

## APPENDICES

### Appendix 1 – Taxonomy of the Immune Evasion Literature

Table A1-1 – Eight Aggregate Category Taxonomy of the Immune Evasion Literature

CL #	CLUSTER THEME	# REC
73	<b>ENHANCE CYTOTOXIC FUNCTION OF NATURAL KILLER CELLS TO REDUCE TUMOR CELL IMMUNE SYSTEM EVASION</b>	799
16	impact of shedding of MICA/B on tumor cell immune evasion, emphasizing treatments that reduce this shedding and augment the cytotoxic function of Natural Killer cells	171
19	Natural Killer Cell therapies against tumors, emphasizing reduction of immune evasion through enhancing activation and surveillance functions of NK cells	628
109	<b>REVERSE DOWNREGULATION OF MHC CLASS 1 TO REDUCE IMMUNE SYSTEM EVASION BY TUMORS</b>	1234
3	role of the immune checkpoint HLA-G in cancer mainly and viral infections secondarily	299
11	HLA Class 1 expression, emphasizing immune system evasion by cancer resulting from HLA Class 1 loss	278
29	1) tumor MHC Class 1 downregulation, 2) the consequent reduction in tumor cell immunosurveillance, and 3) immunotherapies that can reverse downregulation of MHC-I to aid in anti-tumor immunity	360
33	MHC class 1, especially the viral interference with the MHC Class I antigen processing and presentation pathway and the subsequent viral immune system evasion	297
114	<b>IMMUNOTHERAPIES TO COUNTER T-CELL SUPPRESSION BY TUMOR MICROENVIRONMENT AND ENHANCE IMMUNE CHECKPOINT BLOCKADE</b>	8193
6	role of fas and fasL in cancer, emphasizing immune evasion by T lymphocyte apoptosis	233
13	hepatocellular carcinoma, emphasizing the potential role of immunotherapies in countering tumor microenvironment-mediated immune system evasion	222
15	cancer immunotherapy using IDO1 inhibitors that will transform the tumor microenvironment from tolerogenic to normal immunogenic by interfering with the IDO1-kynurenine-aryl hydrocarbon receptor pathway	159
17	pancreatic cancer/pancreatic ductal adenocarcinoma, and the role of immunotherapies in countering the immune system evasion associated with a tolerogenic tumor microenvironment	199
18	1) the role of exosomes in cancer progression (especially their role in tumor microenvironment therapy resistance), and 2) immunotherapies to reduce the tolerance associated with the immunosuppressive tumor microenvironment	297
22	immune responses to human papillomavirus, especially the evasion of host immune defenses by hpv	135
28	myeloid-derived suppressor cells in the tumor microenvironment, especially immunotherapy that targets these cells and reduces their suppression of T-cell activity	471
32	breast cancer, emphasizing mechanisms of immune evasion and immunotherapy resistance	425

36	metabolic impacts on the immune system, especially on the contribution of metabolic dysfunction to the immunosuppressive tumor microenvironment	256
39	nanoparticles (especially cell membrane-coated biomimetic) for tumor drug delivery, emphasizing increased immune escape/evasion and enhanced circulation time	224
40	tumors, especially their escape and evasion from immune system control	402
44	cancer stem cells, and methods for targeting of cancer stem cells within the tumor microenvironment	318
45	head and neck squamous cell carcinomas, especially immunotherapies to counter evasion of the immune system	300
49	tumor microenvironment (especially the role of tumor-associated macrophages in promoting immunosuppression), emphasizing the targeting of these tumor-associated macrophages for cancer immunotherapy	435
56	construction of a cancer-associated long non-coding RNA signature to predict prognosis and immune landscape in myriad cancers	794
59	cancer immunotherapy, emphasizing the targeting of cancer immune evasion	1368
61	tumor microenvironment, especially immune suppression in the tumor microenvironment, and its role in promoting tumor growth	1086
62	cancer, especially the mechanisms of cancer progression and metastasis and their relation to epithelial-mesenchymal plasticity and immune evasion	869
119	<b>IMMUNOTHERAPIES TO COUNTER IMMUNOSUPPRESSIVE TUMOR MICROENVIRONMENT, EMPHASIZING THE PD-1 AND PD-L1 IMMUNE CHECKPOINT</b>	4069
2	CAR T-cell cancer immunotherapy for solid tumors, emphasizing efforts to overcome the tumor cell evasion of the immune system T-cells	233
14	Non-Small-Cell Lung Cancer, emphasizing regulation of PD-L1 expression on tumor cells to minimize tumor immune evasion which occurs by inhibiting functional activity of cytotoxic lymphocytes	350
25	role of immunosuppressive regulatory T cells in cancer immunology (which augment tumor development and progression by inhibiting antitumor immunity), emphasizing strategies to deplete Treg cells and control Treg cell functions to increase antitumor immune responses	472
41	immune checkpoint inhibitors for cancer immunotherapy, including mechanisms of resistance to immune checkpoint blockade	273
46	PD-L1 expression in advanced cancers, especially its contribution to immune escape	842
50	role of the PD-1 and PD-L1 immune checkpoint in immune escape, emphasizing its cancer immunotherapy targeting to inhibit immune escape	421
58	T cells (especially CD8+), emphasizing the role of T cell response suppression (in the tumor microenvironment) in immune system evasion	1478
12	<b>IMMUNE ESCAPE OF THE SARS-COV-2 OMICRON VARIANT</b>	510
12	immune escape of the SARS-CoV-2 Omicron variant	510
21	<b>SARS-COV-2, ITS VARIANTS AND MUTATIONS, AND THE MECHANISMS OF ITS IMMUNE EVASION/IMMUNE ESCAPE</b>	1260
21	SARS-CoV-2, its variants and mutations, and the mechanisms of its immune evasion/immune escape	1260

120	<b>MECHANISMS USED BY BACTERIA, FUNGI, AND PARASITES FOR IMMUNE SYSTEM EVASION</b>	6303
0	role of <i>Helicobacter pylori</i> in gastric cancer, emphasizing immune system evasion	166
5	biofilms (especially bacterial), emphasizing their survival by evasion of the immune system	163
9	immune system evasion by the Lyme Disease Spirochete <i>Borrelia burgdorferi</i>	155
23	pathogenicity and virulence of <i>Staphylococcus aureus</i> , emphasizing the immune system evasion of this gram-positive bacterium	637
30	variant surface glycoprotein coat density in African trypanosomes, emphasizing the mechanisms responsible for this parasite's evasion of its host's immune response	305
31	malaria parasite <i>Plasmodium falciparum</i> , emphasizing pathogenicity and immune evasion	509
37	complement, emphasizing evasion strategies of bacteria	308
42	<i>Porphyromonas gingivalis</i> primarily and <i>Pseudomonas aeruginosa</i> secondarily, emphasizing their immune escape and evasion mechanisms	233
47	fungal pathogens, especially the interactions of <i>Candida albicans</i> with innate host defense and its escape strategies	359
51	parasites (especially <i>Trypanosoma cruzi</i> , <i>Leishmania</i> , and <i>Toxoplasma gondii</i> ), emphasizing their immune system evasion strategies	495
52	<i>Streptococcus pneumoniae</i> , emphasizing pathogenicity, virulence, and immune system evasion	366
53	<i>Salmonella typhimurium</i> and other bacteria, emphasizing their pathogenicity, virulence, and the molecular mechanisms they use to evade the immune system	375
55	<i>Schistosoma mansoni</i> and other helminth parasites, emphasizing evasion of host immunity during infection	537
57	bacteria (especially <i>Neisseria meningitidis</i> ), emphasizing their pathogenesis and immune suppressive mechanisms	394
63	mechanisms associated with evasion of the host immune system by bacterial and viral pathogens	1301
121	<b>MECHANISMS USED BY VIRUSES FOR IMMUNE SYSTEM EVASION</b>	6668
1	Porcine Reproductive and Respiratory Syndrome Virus, emphasizing the role of immune system evasion in promoting viral replication	91
4	hepatitis C virus, especially on the role of immune system evasion in hcv persistence	375
7	immune system evasion by the Epstein-Barr Virus	197
8	escape mutants of the hepatitis B virus	560
10	mechanisms used by African Swine Fever Virus to evade host immune system response and promote viral replication	158
20	herpes simplex virus infections (especially hsv-1), emphasizing immune system evasion by hsv-1, and therapies to reduce this immune evasion	198
24	Kaposi's sarcoma-associated herpesvirus, emphasizing its immune evasion strategies	233
26	human cytomegalovirus disease, with emphasis on countering HCMV immune system evasion and immune escape	309
27	<i>Mycobacterium tuberculosis</i> , especially its mechanisms for evading the host immune response	450
34	B-cell lymphoma, emphasizing mechanisms of immune evasion by lymphoma cells	271

35	HIV-1, emphasizing 1) generation of vaccine-based broadly neutralizing antibodies against HIV-1 and 2) HIV-1 mutations and escape	456
38	Zika virus primarily and flaviviruses and Dengue virus secondarily, emphasizing immune system evasion	201
43	HIV escape mutations, emphasizing the role of the HIV-1 accessory protein Nef in viral pathogenesis, viral replication, and immune escape of HIV-infected cells	553
48	Type 1 interferon, emphasizing viral pathogen-induced mechanisms to antagonize mammalian interferon signaling pathways	729
54	influenza viruses, emphasizing the role of mutations in immune escape variants	643
60	viruses, emphasizing viral strategies for immune system evasion	1244

## Appendix 2 – Selected Titles of Papers from Each Leaf Cluster

(Note: the first row of each table contains the leaf cluster number, the leaf theme, and the number of records in the leaf cluster (in parentheses))

Table A2-1, Cluster 0

Cluster 0 focuses on the role of <i>Helicobacter pylori</i> in gastric cancer, emphasizing immune system evasion (166)
The role of non-coding RNA in the diagnosis and treatment of <i>Helicobacter pylori</i> -related gastric cancer, with a focus on inflammation and immune response.
The impact of autophagic processes on the intracellular fate of <i>Helicobacter pylori</i> : more tricks from an enigmatic pathogen?
Strategies used by <i>Helicobacter pylori</i> to establish persistent infection.
Biosynthesis, structure and biological function of cholesterol glucoside in <i>Helicobacter pylori</i> : A review.
<i>Helicobacter pylori</i> and Its Role in Gastric Cancer.
<i>Helicobacter pylori</i> and the Role of Lipopolysaccharide Variation in Innate Immune Evasion.
Unraveling the factors and mechanism involved in persistence: Host-pathogen interactions in <i>Helicobacter pylori</i> .
<i>Helicobacter pylori</i> evasion strategies of the host innate and adaptive immune responses to survive and develop gastrointestinal diseases.
T cell subsets play an important role in the determination of the clinical outcome of <i>Helicobacter pylori</i> infection.
Cholesterol glucosylation by <i>Helicobacter pylori</i> delays internalization and arrests phagosome maturation in macrophages.
Mucosal inflammation and disease in <i>Helicobacter pylori</i> infection.
<i>Helicobacter pylori</i> vaccination: is there a path to protection?
<i>Helicobacter pylori</i> arginase inhibits T cell proliferation and reduces the expression of the TCR zeta-chain (CD3zeta).
Adhesion and Invasion of Gastric Mucosa Epithelial Cells by <i>Helicobacter pylori</i> .
<i>Helicobacter pylori</i> immune escape is mediated by dendritic cell-induced Treg skewing and Th17 suppression in mice.
Compromised autophagy by MIR30B benefits the intracellular survival of <i>Helicobacter pylori</i> .
Systems-wide analyses of mucosal immune responses to <i>Helicobacter pylori</i> at the interface between pathogenicity and symbiosis.
<i>Helicobacter pylori</i> and T Helper Cells: Mechanisms of Immune Escape and Tolerance.
Oral immunization of mice with a multivalent therapeutic subunit vaccine protects against <i>Helicobacter pylori</i> infection.
<i>Helicobacter pylori</i> downregulates expression of human $\beta$ -defensin 1 in the gastric mucosa in a type IV secretion-dependent fashion.
Induction of maturation and cytokine release of human dendritic cells by <i>Helicobacter pylori</i> .
Effect of <i>Helicobacter pylori</i> on gastric epithelial cells.
Suppression of cell division-associated genes by <i>Helicobacter pylori</i> attenuates proliferation of RAW264.7 monocytic macrophage cells.
Immune evasion strategies used by <i>Helicobacter pylori</i> .
Mechanisms of persistence, innate immune activation and immunomodulation by the gastric pathogen <i>Helicobacter pylori</i> .

Host genetic factors respond to pathogenic step-specific virulence factors of <i>Helicobacter pylori</i> in gastric carcinogenesis.
<i>Helicobacter pylori</i> resists phagocytosis by macrophages: quantitative assessment by confocal microscopy and fluorescence-activated cell sorting.
Serum-derived IgG1-mediated immune exclusion as a mechanism of protection against <i>H. pylori</i> infection.
Immune evasion by <i>Helicobacter pylori</i> : gastric spiral bacteria lack surface immunoglobulin deposition and reactivity with homologous antibodies.
Cholesterol glucosylation-based survival strategy in <i>Helicobacter pylori</i> .
Partial characterization of a cell proliferation-inhibiting protein produced by <i>Helicobacter pylori</i> .
Review: Pathogenesis of <i>Helicobacter pylori</i> infection.
<i>Helicobacter pylori</i> inflammation, immunity, and vaccines.
The Cyclopropane Fatty Acid Synthase Mediates Antibiotic Resistance and Gastric Colonization of <i>Helicobacter pylori</i> .
<i>Helicobacter pylori</i> cholesterol- $\alpha$ -glucosyltransferase manipulates cholesterol for bacterial adherence to gastric epithelial cells.
<i>Helicobacter pylori</i> arginase inhibits nitric oxide production by eukaryotic cells: a strategy for bacterial survival.
Immunity, inflammation, and vaccines for <i>Helicobacter pylori</i> .
[Pathogenesis of <i>Helicobacter pylori</i> infection].
<i>Helicobacter pylori</i> : present status and future prospects in Japan.
<i>Helicobacter pylori</i> attenuates lipopolysaccharide-induced nitric oxide production by murine macrophages.
Immune response modulation in inflammatory bowel diseases by <i>Helicobacter pylori</i> infection.
Interplay of the Gastric Pathogen <i>Helicobacter pylori</i> with Toll-Like Receptors.
<i>Helicobacter pylori</i> inhibits dendritic cell maturation via interleukin-10-mediated activation of the signal transducer and activator of transcription 3 pathway.
<i>Helicobacter pylori</i> Deregulates T and B Cell Signaling to Trigger Immune Evasion.
<i>Helicobacter pylori</i> inhibits phagocytosis by professional phagocytes involving type IV secretion components.
Multi-functional vesicles improve <i>Helicobacter pylori</i> eradication by a comprehensive strategy based on complex pathological microenvironment.
Negative selection of T cells by <i>Helicobacter pylori</i> as a model for bacterial strain selection by immune evasion.
<i>Helicobacter pylori</i> is invasive and it may be a facultative intracellular organism.
<i>Helicobacter pylori</i> cag pathogenicity island's role in B7-H1 induction and immune evasion.
Phagocytosis and persistence of <i>Helicobacter pylori</i> .
<i>Helicobacter pylori</i> virulence factors: subversion of host immune system and development of various clinical outcomes.
<i>Helicobacter pylori</i> targets dendritic cells to induce immune tolerance, promote persistence and confer protection against allergic asthma.
<i>Helicobacter pylori</i> -induced apoptosis in T cells is mediated by the mitochondrial pathway independent of death receptors.
Masking of typical TLR4 and TLR5 ligands modulates inflammation and resolution by <i>Helicobacter pylori</i> .
Prior to Foxp3 <sup>+</sup> regulatory T-cell induction, interleukin-10-producing B cells expand after <i>Helicobacter pylori</i> infection.

Lactobacillus gasseri Suppresses the Production of Proinflammatory Cytokines in Helicobacter pylori-Infected Macrophages by Inhibiting the Expression of ADAM17.
Helicobacter pylori infection: pathogenesis.
Cholesterol glucosylation promotes immune evasion by Helicobacter pylori.
Genome and population dynamics during chronic infection with Helicobacter pylori.
Stable isotope labeling by amino acids in cell culture based proteomics reveals differences in protein abundances between spiral and coccoid forms of the gastric pathogen Helicobacter pylori.
The Role of Helicobacter pylori Outer Membrane Proteins in Adherence and Pathogenesis.
H pylori and host interactions that influence pathogenesis.
[The chemical structure of Helicobacter pylori lipopolysaccharide and innate immune response].
Immune Response in H. pylori-Associated Gastritis and Gastric Cancer.
Helicobacter pylori Infection.
Colonize, evade, flourish: how glyco-conjugates promote virulence of Helicobacter pylori.
Events at the host-microbial interface of the gastrointestinal tract IV. The pathogenesis of Helicobacter pylori persistence.
Helicobacter pylori $\gamma$ -glutamyltranspeptidase impairs T-lymphocyte function by compromising metabolic adaption through inhibition of cMyc and IRF4 expression.
Why can't we make an effective vaccine against Helicobacter pylori?
Pathogenesis of Helicobacter pylori infection.
Statins' Regulation of the Virulence Factors of Helicobacter pylori and the Production of ROS May Inhibit the Development of Gastric Cancer.
Comparative analysis of the interaction of Helicobacter pylori with human dendritic cells, macrophages, and monocytes.
Chimeric flagellin as the self-adjuvanting antigen for the activation of immune response against Helicobacter pylori.
Redox biology and gastric carcinogenesis: the role of Helicobacter pylori.
The Helicobacter pylori autotransporter ImaA (HP0289) modulates the immune response and contributes to host colonization.
Enhanced expression of indoleamine 2,3-dioxygenase in Helicobacter pylori-infected human gastric mucosa modulates Th1/Th2 pathway and interleukin 17 production.
Immune Cell Signaling by Helicobacter pylori: Impact on Gastric Pathology.
Helicobacter pylori VacA Targets Myeloid Cells in the Gastric Lamina Propria To Promote Peripherally Induced Regulatory T-Cell Differentiation and Persistent Infection.
In silico experiment with an-antigen-toll like receptor-5 agonist fusion construct for immunogenic application to Helicobacter pylori.
Helicobacter pylori versus the host: remodeling of the bacterial outer membrane is required for survival in the gastric mucosa.
Arginase II restricts host defense to Helicobacter pylori by attenuating inducible nitric oxide synthase translation in macrophages.
Immune response to H. pylori.
Prediction of Host-Pathogen Interactions for Helicobacter pylori by Interface Mimicry and Implications to Gastric Cancer.
Helicobacter pylori induces apoptosis of human monocytes but not monocyte-derived dendritic cells: role of the cag pathogenicity island.
Update on the immunologic basis of Helicobacter pylori gastritis.
Immune Evasion Strategies and Persistence of Helicobacter pylori.

Helicobacter pylori outer membrane protein, HomC, shows geographic dependent polymorphism that is influenced by the Bab family.
HomA and HomB, outer membrane proteins of Helicobacter pylori down-regulate activation-induced cytidine deaminase (AID) and Ig switch germline transcription and thereby affect class switch recombination (CSR) of Ig genes in human B-cells.
Periplasmic cleavage and modification of the 1-phosphate group of Helicobacter pylori lipid A.
Microbial carcinogenesis: Lactic acid bacteria in gastric cancer.
Helicobacter pylori interferes with leukocyte migration via the outer membrane protein HopQ and via CagA translocation.
Immunology of Helicobacter pylori: insights into the failure of the immune response and perspectives on vaccine studies.
Helicobacter pylori and the innate immune system.
Nod1 Imprints Inflammatory and Carcinogenic Responses toward the Gastric Pathogen Helicobacter pylori.
Helicobacter pylori type IV secretion, host cell signalling and vaccine development.
Activity and Functional Importance of Helicobacter pylori Virulence Factors.
Potential animal models of Helicobacter pylori infection in immunological and vaccine research.
Lipopolysaccharide Lewis Antigens.
Cellular evasion strategies of Helicobacter pylori in regulating its intracellular fate.
The role of genome diversity and immune evasion in persistent infection with Helicobacter pylori.
Vitamin D3 activates the autolysosomal degradation function against Helicobacter pylori through the PDIA3 receptor in gastric epithelial cells.
The leukocyte-associated immunoglobulin (Ig)-like receptor-1 modulating cell apoptosis and inflammatory cytokines secretion in THP-1 cells after Helicobacter pylori infection.
Helicobacter pylori induces apoptosis of macrophages in association with alterations in the mitochondrial pathway.
Helicobacter pylori gene silencing in vivo demonstrates urease is essential for chronic infection.
Inhibition of heat shock protein expression by Helicobacter pylori.
Lewis X biosynthesis in Helicobacter pylori. Molecular cloning of an alpha(1,3)-fucosyltransferase gene.
Why Helicobacter pylori has Lewis antigens.
High Helicobacter pylori Bacterial Load and Low Cytokine Expression Levels Are Associated with Nodular Gastropathy.
Virulence mechanisms and persistence strategies of the human gastric pathogen Helicobacter pylori.
Biochemical and pathophysiological characterization of Helicobacter pylori asparaginase.
Lipopolysaccharide Structure and Biosynthesis in Helicobacter pylori.
Helicobacter pylori flagellins have very low intrinsic activity to stimulate human gastric epithelial cells via TLR5.
Immune evasion by Helicobacter pylori is mediated by induction of macrophage arginase II.
Moonlighting of Helicobacter pylori catalase protects against complement-mediated killing by utilising the host molecule vitronectin.
mRNA expression profiling reveals a role of Helicobacter pylori vacuolating toxin in escaping host defense.
Role of AmiA in the morphological transition of Helicobacter pylori and in immune escape.
The Helicobacter pylori virulence effector CagA abrogates human $\beta$ -defensin 3 expression via inactivation of EGFR signaling.



Lipopolysaccharide in bacterial chronic infection: insights from <i>Helicobacter pylori</i> lipopolysaccharide and lipid A.
Relevance of fucosylation and Lewis antigen expression in the bacterial gastroduodenal pathogen <i>Helicobacter pylori</i> .
Phase-variable restriction/modification systems are required for <i>Helicobacter pylori</i> colonization.
HLA-G 14-bp Ins/Ins Genotype in Patients Harboring <i>Helicobacter pylori</i> Infection: A Potential Risk Factor?
Modulation of lymphocyte proliferation induced by gastric MALT lymphoma-associated <i>Helicobacter pylori</i> strains.
<i>Helicobacter pylori</i> infection interferes with epithelial Stat6-mediated interleukin-4 signal transduction independent of cagA, cagE, or VacA.
In vitro suppression of dendritic cells by <i>Helicobacter pylori</i> OipA.
<i>Helicobacter pylori</i> induces ERK-dependent formation of a phospho-c-Fos c-Jun activator protein-1 complex that causes apoptosis in macrophages.
Phase variation mediated niche adaptation during prolonged experimental murine infection with <i>Helicobacter pylori</i> .
Modification in the ppk gene of <i>Helicobacter pylori</i> during single and multiple experimental murine infections.
Induction of Mincle by <i>Helicobacter pylori</i> and consequent anti-inflammatory signaling denote a bacterial survival strategy.
<i>Helicobacter pylori</i> flagellin: TLR5 evasion and fusion-based conversion into a TLR5 agonist.
Modulation of the CD4+ T-cell response by <i>Helicobacter pylori</i> depends on known virulence factors and bacterial cholesterol and cholesterol $\alpha$ -glucoside content.
How <i>Helicobacter pylori</i> acquired its name, and how it overcomes gastric defence mechanisms.
A comprehensive analysis of <i>Helicobacter pylori</i> plasticity zones reveals that they are integrating conjugative elements with intermediate integration specificity.
Immunosuppressive proteins isolated from spiral and coccoid cytoplasmic solutions of <i>Helicobacter pylori</i> .
Outer membrane biogenesis in <i>Escherichia coli</i> , <i>Neisseria meningitidis</i> , and <i>Helicobacter pylori</i> : paradigm deviations in <i>H. pylori</i> .
Transcriptional phase variation of a type III restriction-modification system in <i>Helicobacter pylori</i> .
An enzymatic ruler modulates Lewis antigen glycosylation of <i>Helicobacter pylori</i> LPS during persistent infection.
sncRNAs packaged by <i>Helicobacter pylori</i> outer membrane vesicles attenuate IL-8 secretion in human cells.
Serologic IgG response to urease in <i>Helicobacter pylori</i> -infected persons from Mexico.
Species-specific immunity to <i>Helicobacter suis</i> .
Molecular modelling of the gastric barrier response, from infection to carcinogenesis.
<i>Helicobacter pylori</i> proteins response to nitric oxide stress.
Crystallization and preliminary crystallographic studies of <i>Helicobacter pylori</i> arginase.
Oxidative stress-induced peptidoglycan deacetylase in <i>Helicobacter pylori</i> .
Variations in <i>Helicobacter pylori</i> lipopolysaccharide to evade the innate immune component surfactant protein D.
Arginase 2 deletion leads to enhanced M1 macrophage activation and upregulated polyamine metabolism in response to <i>Helicobacter pylori</i> infection.
IgA antibodies impair resistance against <i>Helicobacter pylori</i> infection: studies on immune evasion in IL-10-deficient mice.

Regulation of vacuolar pH and its modulation by some microbial species.
Molecular anatomy and pathogenic actions of <i>Helicobacter pylori</i> CagA that underpin gastric carcinogenesis.
Beta2 integrin mediates entry of a bacterial toxin into T lymphocytes.
<i>Helicobacter pylori</i> colonization of the human gastric epithelium: a bug's first step is a novel target for us.
Who ate whom? Adaptive <i>Helicobacter</i> genomic changes that accompanied a host jump from early humans to large felines.
Therapeutic potential of cholesteryl O-acyl $\alpha$ -glucoside found in <i>Helicobacter pylori</i> .
Comparison of enzymatic properties and small molecule inhibition of $\gamma$ -glutamyltranspeptidases from pathogenic and commensal bacteria.
East-Asian <i>Helicobacter pylori</i> strains synthesize heptan-deficient lipopolysaccharide.
Epigenetics and <i>Helicobacter pylori</i> .
Antiparasitic drug nitazoxanide inhibits the pyruvate oxidoreductases of <i>Helicobacter pylori</i> , selected anaerobic bacteria and parasites, and <i>Campylobacter jejuni</i> .
Recombinant herpesvirus glycoprotein G improves the protective immune response to <i>Helicobacter pylori</i> vaccination in a mouse model of disease.
<i>Helicobacter pylori</i> Outer Membrane Vesicles Protect the Pathogen From Reactive Oxygen Species of the Respiratory Burst.
Molecular mimicry of ferret gastric epithelial blood group antigen A by <i>Helicobacter mustelae</i> .
Preclinical studies of amoxicillin, a systemic therapeutic developed for treatment of <i>Clostridium difficile</i> infections that also shows efficacy against <i>Helicobacter pylori</i> .
Characterization of <i>Helicobacter pylori</i> VacA-containing vacuoles (VAVs), VacA intracellular trafficking and interference with calcium signalling in T lymphocytes.
At home with hostility: How do pathogenic bacteria evade mammalian immune surveillance to establish persistent infection?
Characterization of the non-glandular gastric region microbiota in <i>Helicobacter suis</i> -infected versus non-infected pigs identifies a potential role for <i>Fusobacterium gastrois</i> in gastric ulceration.
Potential role of enterohepatic <i>Helicobacter</i> species as a facilitating factor in the development of <i>Chlamydia trachomatis</i> proctitis.
Structural, enzymatic and biochemical studies on <i>Helicobacter pylori</i> arginase.
Role of lipid rafts in persistent <i>Helicobacter pylori</i> infection: a narrative review.

Table A2-2, Cluster 1

Cluster 1 focuses on Porcine Reproductive and Respiratory Syndrome Virus, emphasizing the role of immune system evasion in promoting viral replication (91)
MicroRNA-376b-3p Promotes Porcine Reproductive and Respiratory Syndrome Virus Replication by Targeting Viral Restriction Factor TRIM22.
MicroRNA 373 Facilitates the Replication of Porcine Reproductive and Respiratory Syndrome Virus by Its Negative Regulation of Type I Interferon Induction.
Taming PRRSV: revisiting the control strategies and vaccine design.
Porcine Reproductive and Respiratory Syndrome Virus: Immune Escape and Application of Reverse Genetics in Attenuated Live Vaccine Development.
The Function of the PRRSV-Host Interactions and Their Effects on Viral Replication and Propagation in Antiviral Strategies.

Phylogenetic analysis of porcine reproductive and respiratory syndrome virus in Vietnam, 2021.
Nonstructural protein 11 (nsp11) of porcine reproductive and respiratory syndrome virus (PRRSV) promotes PRRSV infection in MARC-145 cells.
Evasion of Antiviral Innate Immunity by Porcine Reproductive and Respiratory Syndrome Virus.
Role of transcription factors in porcine reproductive and respiratory syndrome virus infection: A review.
Evasion strategies of porcine reproductive and respiratory syndrome virus.
N-Acetyltransferase 9 Inhibits Porcine Reproductive and Respiratory Syndrome Virus Proliferation by N-Terminal Acetylation of the Structural Protein GP5.
Exosomes Mediate Intercellular Transmission of Porcine Reproductive and Respiratory Syndrome Virus.
Porcine Reproductive and Respiratory Syndrome Virus Nonstructural Protein 4 Cleaves Porcine DCP1a To Attenuate Its Antiviral Activity.
Innate and adaptive immunity against Porcine Reproductive and Respiratory Syndrome Virus.
Interleukin-1 receptor antagonist: an early immunomodulatory cytokine induced by porcine reproductive and respiratory syndrome virus.
RBM39 Alters Phosphorylation of c-Jun and Binds to Viral RNA to Promote PRRSV Proliferation.
Both Nsp1 $\beta$ and Nsp11 are responsible for differential TNF- $\alpha$ production induced by porcine reproductive and respiratory syndrome virus strains with different pathogenicity in vitro.
Porcine reproductive and respiratory syndrome virus vaccines: current status and strategies to a universal vaccine.
Induction of HOXA3 by Porcine Reproductive and Respiratory Syndrome Virus Inhibits Type I Interferon Response through Negative Regulation of HO-1 Transcription.
SILAC-based quantitative proteomic analysis of secretome of Marc-145 cells infected with porcine reproductive and respiratory syndrome virus.
Interaction between innate immunity and porcine reproductive and respiratory syndrome virus.
Porcine Reproductive and Respiratory Syndrome Virus Enhances Self-Replication via AP-1-Dependent Induction of SOCS1.
Porcine reproductive and respiratory syndrome virus-infected alveolar macrophages contain no detectable levels of viral proteins in their plasma membrane and are protected against antibody-dependent, complement-mediated cell lysis.
Regulation and evasion of antiviral immune responses by porcine reproductive and respiratory syndrome virus.
miR-382-5p promotes porcine reproductive and respiratory syndrome virus (PRRSV) replication by negatively regulating the induction of type I interferon.
Porcine Reproductive and Respiratory Syndrome Virus Evades Antiviral Innate Immunity via MicroRNAs Regulation.
Phenotypic and functional modulation of bone marrow-derived dendritic cells by porcine reproductive and respiratory syndrome virus.
[Advance in immunology and immune evasion of PRRSV].
Interferon-Induced Transmembrane Protein 3 Is a Virus-Associated Protein Which Suppresses Porcine Reproductive and Respiratory Syndrome Virus Replication by Blocking Viral Membrane Fusion.
Porcine reproductive and respiratory syndrome virus counteracts the porcine intrinsic virus restriction factors-IFITM1 and Tetherin in MARC-145 cells.
Replication-competent recombinant porcine reproductive and respiratory syndrome (PRRS) viruses expressing indicator proteins and antiviral cytokines.

Quantitative interactome reveals that porcine reproductive and respiratory syndrome virus nonstructural protein 2 forms a complex with viral nucleocapsid protein and cellular vimentin.
The N-N non-covalent domain of the nucleocapsid protein of type 2 porcine reproductive and respiratory syndrome virus enhances induction of IL-10 expression.
Porcine reproductive and respiratory syndrome virus increases SOCS3 production via activation of p38/AP-1 signaling pathway to promote viral replication.
Transcriptomic Analysis of Liver Indicates Novel Vaccine to Porcine Reproductive and Respiratory Virus Promotes Homeostasis in T-Cell and Inflammatory Immune Responses Compared to a Commercial Vaccine in Pigs.
Porcine reproductive and respiratory syndrome virus-mediated lactate facilitates virus replication by targeting MAVS.
Evaluation of immune responses to porcine reproductive and respiratory syndrome virus in pigs during early stage of infection under farm conditions.
Key Gaps in the Knowledge of the Porcine Respiratory Reproductive Syndrome Virus (PRRSV).
Targeting Swine Leukocyte Antigen Class I Molecules for Proteasomal Degradation by the nsp1 $\alpha$ Replicase Protein of the Chinese Highly Pathogenic Porcine Reproductive and Respiratory Syndrome Virus Strain JXwn06.
Highly Pathogenic Porcine Reproductive and Respiratory Syndrome Virus Nsp4 Cleaves VISA to Impair Antiviral Responses Mediated by RIG-I-like Receptors.
Strategies to broaden the cross-protective efficacy of vaccines against porcine reproductive and respiratory syndrome virus.
Impact of PRRSV on activation and viability of antigen presenting cells.
Porcine Reproductive and Respiratory Syndrome Virus Nonstructural Protein 1 Beta Interacts with Nucleoporin 62 To Promote Viral Replication and Immune Evasion.
PRRSV Non-Structural Proteins Orchestrate Porcine E3 Ubiquitin Ligase RNF122 to Promote PRRSV Proliferation.
Porcine reproductive and respiratory syndrome virus (PRRSV) could be sensed by professional beta interferon-producing system and had mechanisms to inhibit this action in MARC-145 cells.
Porcine reproductive and respiratory syndrome virus genetic variability a management and diagnostic dilemma.
Downregulation of miR-122 by porcine reproductive and respiratory syndrome virus promotes viral replication by targeting SOCS3.
Nonstructural proteins nsp2TF and nsp2N of porcine reproductive and respiratory syndrome virus (PRRSV) play important roles in suppressing host innate immune responses.
Porcine Reproductive and Respiratory Syndrome Virus nsp11 Antagonizes Type I Interferon Signaling by Targeting IRF9.
Challenges for porcine reproductive and respiratory syndrome virus (PRRSV) vaccinology.
Pathogenesis of porcine reproductive and respiratory syndrome virus.
Porcine Reproductive and Respiratory Syndrome Virus (PRRSV) Inhibits RNA-Mediated Gene Silencing by Targeting Ago-2.
LGP2 Promotes Type I Interferon Production To Inhibit PRRSV Infection via Enhancing MDA5-Mediated Signaling.
Transcriptome sequencing analysis of porcine alveolar macrophages infected with PRRSV strains to elucidate virus pathogenicity and immune evasion strategies.
Identification of Differentially Expressed Non-coding RNA in Porcine Alveolar Macrophages from Tongcheng and Large White Pigs Responded to PRRSV.

ORF5 of porcine reproductive and respiratory syndrome virus (PRRSV) is a target of diversifying selection as infection progresses from acute infection to virus rebound.
Identification of three site mutations in nonstructural protein 1 $\beta$ , glycoprotein 3 and glycoprotein 5 that correlate with increased interferon $\alpha$ resistance of porcine reproductive and respiratory syndrome virus.
Nsp1 $\alpha$ of Porcine Reproductive and Respiratory Syndrome Virus Strain BB0907 Impairs the Function of Monocyte-Derived Dendritic Cells via the Release of Soluble CD83.
Porcine reproductive and respiratory syndrome virus productively infects monocyte-derived dendritic cells and compromises their antigen-presenting ability.
Mucosal vaccines to prevent porcine reproductive and respiratory syndrome: a new perspective.
Post-transcriptional control of type I interferon induction by porcine reproductive and respiratory syndrome virus in its natural host cells.
Porcine reproductive and respiratory syndrome virus nonstructural protein 1 $\beta$ modulates host innate immune response by antagonizing IRF3 activation.
Porcine reproductive and respiratory syndrome virus infection inhibits NF- $\kappa$ B signaling pathway through cleavage of IKK $\beta$ by Nsp4.
Porcine reproductive and respiratory syndrome virus nucleocapsid protein modulates interferon- $\beta$ production by inhibiting IRF3 activation in immortalized porcine alveolar macrophages.
Modulation of host cell responses and evasion strategies for porcine reproductive and respiratory syndrome virus.
A Bayesian phylogeographical analysis of type 1 porcine reproductive and respiratory syndrome virus (PRRSV).
Porcine reproductive and respiratory syndrome virus 3C protease cleaves the mitochondrial antiviral signalling complex to antagonize IFN- $\beta$ expression.
Pathogenic characteristics of three genotype II porcine reproductive and respiratory syndrome viruses isolated from China.
Identification of differentially expressed proteins in porcine alveolar macrophages infected with virulent/attenuated strains of porcine reproductive and respiratory syndrome virus.
Porcine arterivirus activates the NF- $\kappa$ B pathway through I $\kappa$ B degradation.
Acute phase response in porcine reproductive and respiratory syndrome virus infection.
Targeted mutations in a highly conserved motif of the nsp1 $\beta$ protein impair the interferon antagonizing activity of porcine reproductive and respiratory syndrome virus.
Genetic diversity and phylogenetic analysis of glycoprotein 5 of European-type porcine reproductive and respiratory virus strains in Spain.
Control of porcine reproductive and respiratory syndrome (PRRS) through genetic improvements in disease resistance and tolerance.
The role of porcine reproductive and respiratory syndrome virus infection in immune phenotype and Th1/Th2 balance of dendritic cells.
Amino acid residues in the non-structural protein 1 of porcine reproductive and respiratory syndrome virus involved in down-regulation of TNF- $\alpha$ expression in vitro and attenuation in vivo.
Mutations in a Highly Conserved Motif of nsp1 $\beta$ Protein Attenuate the Innate Immune Suppression Function of Porcine Reproductive and Respiratory Syndrome Virus.
Activation of activating Fc gamma receptors down-regulates the levels of interferon $\beta$ , interferon $\gamma$ and interferon $\lambda$ 1 in porcine alveolar macrophages during PRRSV infection.
Whole genome sequencing of clinical specimens reveals the genomic diversity of porcine reproductive and respiratory syndrome viruses emerging in China.

MicroRNA-30c targets the interferon-alpha/beta receptor beta chain to promote type 2 PRRSV infection.
A DNA Prime Immuno-Potentiates a Modified Live Vaccine against the Porcine Reproductive and Respiratory Syndrome Virus but Does Not Improve Heterologous Protection.
Type I interferon suppression-negative and host mRNA nuclear retention-negative mutation in nsp1 $\beta$ confers attenuation of porcine reproductive and respiratory syndrome virus in pigs.
Mapping the Key Residues within the Porcine Reproductive and Respiratory Syndrome Virus nsp1 $\alpha$ Replicase Protein Required for Degradation of Swine Leukocyte Antigen Class I Molecules.
Porcine reproductive and respiratory syndrome virus non-structural protein 1 suppresses tumor necrosis factor-alpha promoter activation by inhibiting NF- $\kappa$ B and Sp1.
A highly pathogenic porcine reproductive and respiratory syndrome virus type 1 (PRRSV-1) strongly modulates cellular innate and adaptive immune subsets upon experimental infection.
Arterivirus molecular biology and pathogenesis.
Molecular characterization of the RNA-protein complex directing -2/-1 programmed ribosomal frameshifting during arterivirus replicase expression.
Arterivirus nsp4 Antagonizes Interferon Beta Production by Proteolytically Cleaving NEMO at Multiple Sites.
Viral evasion of PKR restriction by reprogramming cellular stress granules.
Modulation of Innate Antiviral Immune Response by Porcine Enteric Coronavirus.
Porcine circovirus type 2 exploits cap to inhibit PKR activation through interaction with Hsp40.

Table A2-3, Cluster 2

Cluster 2 focuses on CAR T-cell cancer immunotherapy for solid tumors, emphasizing efforts to overcome the tumor cell evasion of the immune system T-cells (233)
Improving the ability of CAR-T cells to hit solid tumors: Challenges and strategies.
Mechanisms of response and resistance to CAR T cell therapies.
Current Challenges and Strategies for Chimeric Antigen Receptor-T-Cell Therapy for Solid Tumors.
Resistance and recurrence of malignancies after CAR-T cell therapy.
Determinants of response and resistance to CAR T cell therapy.
CAR-T Cell Therapy in Cancer: Tribulations and Road Ahead.
Chimeric Antigen Receptor (CAR) T Cell Therapy for Cancer. Challenges and Opportunities: An Overview.
Bright future or blind alley? CAR-T cell therapy for solid tumors.
CAR-T cell therapy for hematological malignancies: Limitations and optimization strategies.
Overcome tumor relapse in CAR T cell therapy.
Fine-Tuning through Generations: Advances in Structure and Production of CAR-T Therapy.
Genetic Modification of Cytokine Signaling to Enhance Efficacy of CAR T Cell Therapy in Solid Tumors.
CAR-T Cell Therapy: A Door Is Open to Find Innumerable Possibilities of Treatments for Cancer Patients.
Immunotherapy and immunoengineering for breast cancer; a comprehensive insight into CAR-T cell therapy advancements, challenges and prospects.
CAR-T Cells Hit the Tumor Microenvironment: Strategies to Overcome Tumor Escape.
Driving an improved CAR for cancer immunotherapy.
Immunobiology of chimeric antigen receptor T cells and novel designs.
Cytokine IL-36 $\gamma$ improves CAR T-cell functionality and induces endogenous antitumor response.

Gene modification strategies for next-generation CAR T cells against solid cancers.
CAR Talk: How Cancer-Specific CAR T Cells Can Instruct How to Build CAR T Cells to Cure HIV.
[Mechanisms of resistance and escape to CAR-T cells].
CAR-T cell therapy in melanoma: A future success story?
[Not Available].
Compromised antigen binding and signaling interfere with bispecific CD19 and CD79a chimeric antigen receptor function.
Engineering CAR-T Cells for Improved Function Against Solid Tumors.
Chimeric antigen receptor T cells for acute myeloid leukemia.
Chimeric Antigen Receptor T Cells for Glioblastoma: Current Concepts, Challenges, and Future Perspectives.
Nanobody-based CAR T cells that target the tumor microenvironment inhibit the growth of solid tumors in immunocompetent mice.
Engineered CAR-T and novel CAR-based therapies to fight the immune evasion of glioblastoma: gutta cavat lapidem.
Chimeric Antigen Receptor T Cells for B-Cell Acute Lymphoblastic Leukemia.
Current challenges for CAR T-cell therapy of acute myeloid leukemia.
Allogeneic CAR T Cells: An Alternative to Overcome Challenges of CAR T Cell Therapy in Glioblastoma.
Shed antigen-induced blocking effect on CAR-T cells targeting Glypican-3 in Hepatocellular Carcinoma.
New development in CAR-T cell therapy.
Engineering chimeric antigen receptor-T cells for cancer treatment.
The one-two punch (of CAR T cells).
Chimeric Antigen Receptor (CAR) T Cell Therapy for Metastatic Melanoma: Challenges and Road Ahead.
Hypoimmune anti-CD19 chimeric antigen receptor T cells provide lasting tumor control in fully immunocompetent allogeneic humanized mice.
Dual Targeting CAR-T Cells with Optimal Costimulation and Metabolic Fitness enhance Antitumor Activity and Prevent Escape in Solid Tumors.
Advances in chimeric antigen receptor T cells therapy in the treatment of breast cancer.
Central memory phenotype drives success of checkpoint inhibition in combination with CAR T cells.
Trogocytosis and fratricide killing impede MSLN-directed CAR T cell functionality.
Combining selective inhibitors of nuclear export (SINEs) with chimeric antigen receptor (CAR) T cells for CD19-positive malignancies.
The IAP antagonist birinapant enhances chimeric antigen receptor T cell therapy for glioblastoma by overcoming antigen heterogeneity.
Born to survive: how cancer cells resist CAR T cell therapy.
A deep insight into CRISPR/Cas9 application in CAR-T cell-based tumor immunotherapies.
Tumour Escape from CAR-T Cells.
Current progress in chimeric antigen receptor T cell therapy for glioblastoma multiforme.
JAK/STAT-Dependent Chimeric Antigen Receptor (CAR) Expression: A Design Benefiting From a Dual AND/OR Gate Aiming to Increase Specificity, Reduce Tumor Escape and Affect Tumor Microenvironment.
Role of CAR T Cell Metabolism for Therapeutic Efficacy.
CD58 loss in tumor cells confers functional impairment of CAR T cells.
Combination Immunotherapy with CAR T Cells and Checkpoint Blockade for the Treatment of Solid Tumors.

Engineering Novel CD19/CD22 Dual-Target CAR-T Cells for Improved Anti-Tumor Activity.
[Perspectives for the evolution and use of CAR-T cells].
Multi Targeted CAR-T Cell Therapies for B-Cell Malignancies.
Driving CARs on the uneven road of antigen heterogeneity in solid tumors.
The Role of Chimeric Antigen Receptor-T Cell Therapy in the Treatment of Hematological Malignancies: Advantages, Trials, and Tribulations, and the Road Ahead.
CD40 Ligand-Modified Chimeric Antigen Receptor T Cells Enhance Antitumor Function by Eliciting an Endogenous Antitumor Response.
[CAR-T cells immunotherapy in multiple myeloma: Present and future].
Chimeric Antigen Receptor T-Cell Therapies for Aggressive B-Cell Lymphomas: Current and Future State of the Art.
Point mutation in CD19 facilitates immune escape of B cell lymphoma from CAR-T cell therapy.
High-Affinity Chimeric Antigen Receptor With Cross-Reactive scFv to Clinically Relevant EGFR Oncogenic Isoforms.
CAR-T cell therapy for triple-negative breast cancer and other solid tumors: preclinical and clinical progress.
Loop CD20/CD19 CAR-T cells eradicate B-cell malignancies efficiently.
Advances in CAR-T Cell Genetic Engineering Strategies to Overcome Hurdles in Solid Tumors Treatment.
Chimeric antigen receptor (CAR) T-cell therapy for multiple myeloma.
Beyond CD19 CAR-T cells in lymphoma.
IL-15 armoring enhances the antitumor efficacy of claudin 18.2-targeting CAR-T cells in syngeneic mouse tumor models.
Exploring the Dilemma of Allogeneic Hematopoietic Cell Transplantation after Chimeric Antigen Receptor T Cell Therapy: To Transplant or Not?
Tumor-Derived Extracellular Vesicles Impair CD171-Specific CD4(+) CAR T Cell Efficacy.
Application of Chimeric Antigen Receptor T Cells in the Treatment of Hematological Malignancies.
Vaccine-boosted CAR T crosstalk with host immunity to reject tumors with antigen heterogeneity.
Adapter CAR T Cell Therapy for the Treatment of B-Lineage Lymphomas.
Engineered CAR-T cells targeting TAG-72 and CD47 in ovarian cancer.
Chimeric antigen receptor-engineered T-cell therapy for liver cancer.
GPC3-targeted CAR-T cells secreting B7H3-targeted BiTE exhibit potent cytotoxicity activity against hepatocellular carcinoma cell in the in vitro assay.
T cells expressing CD5/CD7 bispecific chimeric antigen receptors with fully human heavy-chain-only domains mitigate tumor antigen escape.
Chimeric antigen receptor-modified T cells strike back.
Chimeric antigen receptor T cell therapy for non-Hodgkin lymphoma.
Tandem chimeric antigen receptor (CAR) T cells targeting EGFRvIII and IL-13R $\alpha$ 2 are effective against heterogeneous glioblastoma.
In Like a Lamb; Out Like a Lion: Marching CAR T Cells Toward Enhanced Efficacy in B-ALL.
CAR T targets and microenvironmental barriers of osteosarcoma.
Immunogenetic Metabolomics Reveals Key Enzymes That Modulate CAR T-cell Metabolism and Function.
CAR T-cell therapy of solid tumors.
Generation and functional characterization of CAR exosomes.



Acquisition of a CD19-negative myeloid phenotype allows immune escape of MLL-rearranged B-ALL from CD19 CAR-T-cell therapy.
CAR T-cells in acute lymphoblastic leukemia: Current results.
Chimeric-Antigen-Receptor (CAR) T Cells and the Factors Influencing their Therapeutic Efficacy.
Cellular Immunotherapy in B-Cell Malignancy.
Mechanisms underlying CD19-positive ALL relapse after anti-CD19 CAR T cell therapy and associated strategies.
Chimeric Antigen Receptor Therapy: How Are We Driving in Solid Tumors?
Chimeric antigen receptor-modified T cells: CD19 and the road beyond.
Tandem CAR T cells targeting HER2 and IL13R $\alpha$ 2 mitigate tumor antigen escape.
Bone Marrow Mesenchymal Stromal Cells Can Render Multiple Myeloma Cells Resistant to Cytotoxic Machinery of CAR T Cells through Inhibition of Apoptosis.
The rational development of CD5-targeting biepitopic CARs with fully human heavy-chain-only antigen recognition domains.
Current status and future development of anti-HIV chimeric antigen receptor T-cell therapy.
Genetically Modified T-Cell-Based Adoptive Immunotherapy in Hematological Malignancies.
Prospects to improve chimeric antigen receptor T-cell therapy for solid tumors.
SJI 2020 special issue: A catalogue of Ovarian Cancer targets for CAR therapy.
Increased antitumor activities of glypican-3-specific chimeric antigen receptor-modified T cells by coexpression of a soluble PD1-CH3 fusion protein.
Driving CAR T cells towards dermatologic oncology.
Nanosensors in clinical development of CAR-T cell immunotherapy.
Kinetics of tumor destruction by chimeric antigen receptor-modified T cells.
Targeting glycosylated antigens on cancer cells using siglec-7/9-based CAR T-cells.
CRISPR/Cas9 mediated deletion of the adenosine A2A receptor enhances CAR T cell efficacy.
CAR T cell trogocytosis and cooperative killing regulate tumour antigen escape.
[Not Available].
Arming Anti-EGFRvIII CAR-T With TGF $\beta$ Trap Improves Antitumor Efficacy in Glioma Mouse Models.
Non-viral TRAC-knocked-in CD19(KI)CAR-T and gp350(KI)CAR-T cells tested against Burkitt lymphomas with type 1 or 2 EBV infection: In vivo cellular dynamics and potency.
Improving Chimeric Antigen Receptor-Modified T Cell Function by Reversing the Immunosuppressive Tumor Microenvironment of Pancreatic Cancer.
Combining BCMA-targeting CAR-T with TCR-engineered T-cell therapy to prevent immune escape of multiple myeloma.
Immunogenetic metabolomics revealed key enzymes that modulate CAR-T metabolism and function.
SEAKER cells coordinate cellular immunotherapy with localized chemotherapy.
CAR T cells targeting tumor endothelial marker CLEC14A inhibit tumor growth.
Cancer immunotherapy with lymphocytes genetically engineered with T cell receptors for solid cancers.
Multi-Specific CAR Targeting to Prevent Antigen Escape.
Chimeric antigen receptors: unleashing a new age of anti-cancer therapy.
CX3CR1 deficiency-induced TIL tumor restriction as a novel addition for CAR-T design in solid malignancies.
Mesothelin-targeted CAR-T therapy combined with irinotecan for the treatment of solid cancer.
APDL1-CART cells exhibit strong PD-L1-specific activity against leukemia cells.
The Generation of CAR-Transfected Natural Killer T Cells for the Immunotherapy of Melanoma.

The molecular perspective on the melanoma and genome engineering of T-cells in targeting therapy.
Anti-ROR1 CAR-T cells: Architecture and performance.
Genetically Engineered T-Cells for Malignant Glioma: Overcoming the Barriers to Effective Immunotherapy.
A tandem CD19/CD20 CAR lentiviral vector drives on-target and off-target antigen modulation in leukemia cell lines.
B7-H3-redirected chimeric antigen receptor T cells target glioblastoma and neurospheres.
Preclinical Development of Bivalent Chimeric Antigen Receptors Targeting Both CD19 and CD22.
T Cells Expressing CD19/CD20 Bispecific Chimeric Antigen Receptors Prevent Antigen Escape by Malignant B Cells.
FCγ Chimeric Receptor-Engineered T Cells: Methodology, Advantages, Limitations, and Clinical Relevance.
Enhanced AC133-specific CAR T cell therapy induces durable remissions in mice with metastatic small cell lung cancer.
CAR-T with License to Kill Solid Tumors in Search of a Winning Strategy.
Targeted glycan degradation potentiates cellular immunotherapy for solid tumors.
Chimeric antigen receptor (CAR)-engineered lymphocytes for cancer therapy.
[CAR-T cells in acute lymphoblastic leukemias: What's new?].
Redirecting T cells with Chimeric Antigen Receptor (CAR) for the treatment of childhood acute lymphoblastic leukemia.
Immunotherapy: enhancing the efficacy of this promising therapeutic in multiple cancers.
CSPG4 Shows Promise for Glioblastoma CAR T Therapy.
Does lineage plasticity enable escape from CAR-T cell therapy? Lessons from MLL-r leukemia.
CAR19/22 T cell cocktail therapy for B-ALL relapsed after allogeneic hematopoietic stem cell transplantation.
Clinical Trials of Cellular Therapies in Solid Tumors.
Three-Dimensional Bioconjugated Liquid-Like Solid (LLS) Enhance Characterization of Solid Tumor - Chimeric Antigen Receptor T cell interactions.
Engineering Hematopoietic Cells for Cancer Immunotherapy: Strategies to Address Safety and Toxicity Concerns.
Tumor interferon signaling and suppressive myeloid cells are associated with CAR T-cell failure in large B-cell lymphoma.
The Interplay between T Cells and Cancer: The Basis of Immunotherapy.
Engineered T-cell Receptor T Cells for Cancer Immunotherapy.
NK-92 cell, another ideal carrier for chimeric antigen receptor.
Realism and pragmatism in developing an effective chimeric antigen receptor T-cell product for solid cancers.
Immune dysregulation in multiple myeloma: the current and future role of cell-based immunotherapy.
CD30-Redirected Chimeric Antigen Receptor T Cells Target CD30(+) and CD30(-) Embryonal Carcinoma via Antigen-Dependent and Fas/FasL Interactions.
Leukemia escape in immune desert: intraocular relapse of pediatric pro-B-ALL during systemic control by CD19-CAR T cells.
Immunomodulators in Lymphoma.
Augmenting engineered T-cell strategies in solid cancers through epigenetic priming.
Fully human antibody V(H) domains to generate mono and bispecific CAR to target solid tumors.

Multispecific Targeting with Synthetic Ankyrin Repeat Motif Chimeric Antigen Receptors.
Novel chimeric antigen receptor T cells based on T-cell receptor-like antibodies.
Adoptive immunotherapies in neuro-oncology: classification, recent advances, and translational challenges.
Overcoming key challenges in cancer immunotherapy with engineered T cells.
CD4(+)CD25(+)CD127(low) regulatory T cells associated with the effect of CD19 CAR-T therapy for relapsed/refractory B-cell acute lymphoblastic leukemia.
Auto T cells expressing chimeric antigen receptor derived from auto antibody might be a new treatment for osteosarcoma.
Overcoming CAR-Mediated CD19 Downmodulation and Leukemia Relapse with T Lymphocytes Secreting Anti-CD19 T-cell Engagers.
Programmed cell death protein 1 activation preferentially inhibits CD28.CAR-T cells.
Prospects for adoptive immunotherapy of pancreatic cancer using chimeric antigen receptor-engineered T-cells.
Antigen-Specific TCR-T Cells for Acute Myeloid Leukemia: State of the Art and Challenges.
Chimeric Antigen Receptor Immunotherapy for Solid Tumors: Choosing the Right Ingredients for the Perfect Recipe.
Fifteen years of gene therapy based on chimeric antigen receptors: "are we nearly there yet?"
Host Interactions with Engineered T-cell Micropharmacies.
Current Headway in Cancer Immunotherapy Emphasizing the Practice of Genetically Engineered T Cells to Target Selected Tumor Antigens.
Targeting Acute Myeloid Leukemia Using the RevCAR Platform: A Programmable, Switchable and Combinatorial Strategy.
Leveraging Endogenous Dendritic Cells to Enhance the Therapeutic Efficacy of Adoptive T-Cell Therapy and Checkpoint Blockade.
Engineered T cells: the promise and challenges of cancer immunotherapy.
Single variable domains from the T cell receptor $\beta$ chain function as mono- and bifunctional CARs and TCRs.
Actors on the Scene: Immune Cells in the Myeloma Niche.
Adoptive cellular therapy with T cells expressing the dendritic cell growth factor Flt3L drives epitope spreading and antitumor immunity.
Immunotherapy of metastatic melanoma using genetically engineered GD2-specific T cells.
Cardiotoxicities of novel cancer immunotherapies.
Genetically engineered T cells bearing chimeric nanoconstructed receptors harboring TAG-72-specific camelid single domain antibodies as targeting agents.
Extracellular Matrix Viscosity Reprogramming by In Situ Au Bioreactor-Boosted Microwavegenetics Disables Tumor Escape in CAR-T Immunotherapy.
Genetically modified T cells in cancer therapy: opportunities and challenges.
Chimeric antigen receptor engineered stem cells: a novel HIV therapy.
Arming T Cells with a gp100-Specific TCR and a CSPG4-Specific CAR Using Combined DNA- and RNA-Based Receptor Transfer.
Chimeric antigen receptor engineered NK cellular immunotherapy overcomes the selection of T-cell escape variant cancer cells.
HLA-independent T cell receptors for targeting tumors with low antigen density.
Engineering T cells for adoptive therapy: outsmarting the tumor.
Immunotherapy for the treatment of multiple myeloma.
Investigational immunotherapy targeting CD19 for the treatment of acute lymphoblastic leukemia.

Efficient elimination of primary B-ALL cells in vitro and in vivo using a novel 4-1BB-based CAR targeting a membrane-distal CD22 epitope.
Treating hematological malignancies with cell therapy: where are we now?
Epitope editing enables targeted immunotherapy of acute myeloid leukaemia.
Homozygous BCMA gene deletion in response to anti-BCMA CAR T cells in a patient with multiple myeloma.
Present and Future Role of Immune Targets in Acute Myeloid Leukemia.
CD24: A Novel Target for Cancer Immunotherapy.
Cbl induced ubiquitination of HER2 mediate immune escape from HER2-targeted CAR-T.
Immunosenescence, inflammaging, and cancer immunotherapy efficacy.
Genome editing in engineered T cells for cancer immunotherapy.
The progress and current status of immunotherapy in acute myeloid leukemia.
Combinational targeting offsets antigen escape and enhances effector functions of adoptively transferred T cells in glioblastoma.
Combining a chimeric antigen receptor and a conventional T-cell receptor to generate T cells expressing two additional receptors (TETARs) for a multi-hit immunotherapy of melanoma.
T cell-engaging therapies - BiTEs and beyond.
Chimeric antigen receptors and bispecific antibodies to retarget T cells in pediatric oncology.
TIGIT: A promising target to overcome the barrier of immunotherapy in hematological malignancies.
Targeting Solid Tumors Using CD3 Bispecific Antibodies.
ATF3 and CH25H regulate effector trogocytosis and anti-tumor activities of endogenous and immunotherapeutic cytotoxic T lymphocytes.
In Vitro and In Vivo Antitumor Effect of Anti-CD33 Chimeric Receptor-Expressing EBV-CTL against CD33 Acute Myeloid Leukemia.
Combining T-cell immunotherapy and anti-androgen therapy for prostate cancer.
Adenosine-A2A Receptor Pathway in Cancer Immunotherapy.
The Role of T Cell Immunity in Monoclonal Gammopathy and Multiple Myeloma: From Immunopathogenesis to Novel Therapeutic Approaches.
Multiple cancer-specific antigens are targeted by a chimeric antigen receptor on a single cancer cell.
TCR-based therapy for multiple myeloma and other B-cell malignancies targeting intracellular transcription factor BOB1.
Targeting EGFRvIII for glioblastoma multiforme.
Immunology in the clinic review series; focus on cancer: double trouble for tumours: bi-functional and redirected T cells as effective cancer immunotherapies.
Nanoparticle T-cell engagers as a modular platform for cancer immunotherapy.
Bone Marrow Microenvironment Interplay and Current Clinical Practice in Multiple Myeloma: A Review of the Balkan Myeloma Study Group.
Novel immunotherapies for hematologic malignancies.
The Leading Role of the Immune Microenvironment in Multiple Myeloma: A New Target with a Great Prognostic and Clinical Value.
From Anti-HER-2 to Anti-HER-2-CAR-T Cells: An Evolutionary Immunotherapy Approach for Gastric Cancer.
T cell therapy for nasopharyngeal carcinoma.
An APRIL-based chimeric antigen receptor for dual targeting of BCMA and TACI in multiple myeloma.
Unleashing the immune response against childhood solid cancers.
Immunotherapy in leukaemia.

Novel approaches to immunotherapy for B-cell malignancies.
Upregulation of CD22 by Chidamide promotes CAR T cells functionality.
T-cell Therapies Targeting Multiple Cancer Antigens: The Power of Many.
Is adoptive T-cell therapy for solid tumors coming of age?
Myelodysplastic syndrome and immunotherapy novel to next in-line treatments.
Advances in immunotherapy for acute myeloid leukemia.
Novel Approaches to Acute Myeloid Leukemia Immunotherapy.
Combining mTor inhibitors with rapamycin-resistant T cells: a two-pronged approach to tumor elimination.
Immunotherapy of multiple myeloma.
CD19 Isoforms Enabling Resistance to CART-19 Immunotherapy Are Expressed in B-ALL Patients at Initial Diagnosis.
Cellular immunotherapy strategies for Ewing sarcoma.
Adoptive cellular immunotherapy for virus-associated cancers: a new paradigm in personalized medicine.
Epstein-Barr virus-induced ectopic CD137 expression helps nasopharyngeal carcinoma to escape immune surveillance and enables targeting by chimeric antigen receptors.
Targeting CD19: the good, the bad, and CD81.

Table A2-4, Cluster 3

Cluster 3 focuses on the role of the immune checkpoint HLA-G in cancer mainly and viral infections secondarily (299)
HLA-G/sHLA-G and HLA-G-Bearing Extracellular Vesicles in Cancers: Potential Role as Biomarkers.
The Molecular and Functional Characteristics of HLA-G and the Interaction with Its Receptors: Where to Intervene for Cancer Immunotherapy?
HLA-G: A New Immune Checkpoint in Cancer?
Role of HLA-G in Viral Infections.
HLA-G in melanoma: can the current controversies be solved?
The dual role of HLA-G in cancer.
Human leukocyte antigen G expression: as a significant prognostic indicator for patients with colorectal cancer.
Heterogeneity of HLA-G Expression in Cancers: Facing the Challenges.
The non-classical antigens of HLA-G and HLA-E as diagnostic and prognostic biomarkers and as therapeutic targets in transplantation and tumors.
HLA-G expression correlates with histological grade but not with prognosis in colorectal carcinoma.
[HLA-G: a tolerance molecule implicated in the escape of tumors from immunosurveillance].
Expression of tolerogenic HLA-G molecules in cancer prevents antitumor responses.
HLA-G*0105N null allele encodes functional HLA-G isoforms.
Identification of novel microRNAs regulating HLA-G expression and investigating their clinical relevance in renal cell carcinoma.
Prognostic Value of HLA-G in Malignant Liver and Pancreas Lesions.
Promoter methylation and mRNA expression of HLA-G in relation to HLA-G protein expression in colorectal cancer.
Differential in vivo and in vitro HLA-G expression in melanoma cells: potential mechanisms.
Role of microRNAs on HLA-G expression in human tumors.

New insights into HLA-G mediated tolerance.
A Critical Assessment of the Association between HLA-G Expression by Carcinomas and Clinical Outcome.
HLA-G mediated immune regulation is impaired by a single amino acid exchange in the alpha 2 domain.
HLA-G expression on blasts and tolerogenic cells in patients affected by acute myeloid leukemia.
HLA-G Genotype/Expression/Disease Association Studies: Success, Hurdles, and Perspectives.
Specific activation of the non-classical class I histocompatibility HLA-G antigen and expression of the ILT2 inhibitory receptor in human breast cancer.
HLA-G in cancer: a way to turn off the immune system.
Clinical relevance of miR-mediated HLA-G regulation and the associated immune cell infiltration in renal cell carcinoma.
Hypoxic Modulation of HLA-G Expression through the Metabolic Sensor HIF-1 in Human Cancer Cells.
Human leukocyte antigen G up-regulation in lung cancer associates with high-grade histology, human leukocyte antigen class I loss and interleukin-10 production.
Human leukocyte antigen-G is frequently expressed in glioblastoma and may be induced in vitro by combined 5-aza-2'-deoxycytidine and interferon-γ treatments: results from a multicentric study.
Analysis of HLA-G expression in breast cancer tissues.
HLA-G and immune evasion in cancer cells.
Unfavourable clinical implications for HLA-G expression in acute myeloid leukaemia.
Structure, expression and function of HLA-G in renal cell carcinoma.
HLA-G expression in gastric carcinoma: clinicopathological correlations and prognostic impact.
Mouse models for studies of HLA-G functions in basic science and pre-clinical research.
HLA-G: Too Much or Too Little? Role in Cancer and Autoimmune Disease.
The role of HLA-G in gastrointestinal inflammatory disease and malignancy.
The Role of HLA-G in Tumor Escape: Manipulating the Phenotype and Function of Immune Cells.
HLA-G expression in gastric cancer.
HLA-G expression in human melanoma cells: protection from NK cytotoxicity.
Research on HLA-G: an update.
Hypoxia inducible factor-1 mediates the expression of the immune checkpoint HLA-G in glioma cells through hypoxia response element located in exon 2.
Human Leukocyte Antigen-G Inhibits the Anti-Tumor Effect of Natural Killer Cells via Immunoglobulin-Like Transcript 2 in Gastric Cancer.
Human leukocyte antigen-G expression and polymorphisms promote cancer development and guide cancer diagnosis/treatment.
HLA-G gene activation in tumor cells involves cis-acting epigenetic changes.
Human Leukocyte Antigen-G (HLA-G) Expression in Cancers: Roles in Immune Evasion, Metastasis and Target for Therapy.
A functional role of HLA-G expression in human gliomas: an alternative strategy of immune escape.
Human Leukocyte Antigen-G (HLA-G) Expression in Precancerous and Cancerous Cervical Lesions: Association with Human Papilloma Virus Infection and Host Immune Response.
HLA-G: An Immune Checkpoint Molecule.
The immunosuppressive molecule HLA-G and its clinical implications.
Human leukocyte antigen-G (HLA-G) polymorphism and expression in breast cancer patients.
HLA-G and its KIR ligands in cancer--another enigma yet to be solved?
HLA-G gene repression is reversed by demethylation.

HLA-G Expression in Tumor Tissues and Soluble HLA-G Plasma Levels in Patients with Gastrointestinal Cancer.
HLA-G in melanoma: A new strategy to escape from immunosurveillance?
HLA-G expression and role in advanced-stage classical Hodgkin lymphoma.
HLA-G and Other Immune Checkpoint Molecules as Targets for Novel Combined Immunotherapies.
The Impact of HLA-G 3'UTR Polymorphisms in Breast Cancer in a Tunisian Population.
HLA-G and lymphoproliferative disorders.
The role of HLA-G for protection of human renal cell-carcinoma cells from immune-mediated lysis: implications for immunotherapies.
First immunotherapeutic CAR-T cells against the immune checkpoint protein HLA-G.
Human leukocyte antigen-G is closely associated with tumor immune escape in gastric cancer by increasing local regulatory T cells.
HLA-G expression in malignant melanoma.
No expression of human leukocyte antigen G (HLA-G) in colorectal cancer cells.
Human leukocyte antigen-G (HLA-G) expression plays an important role in the diagnosis and grading of endometrial cancer.
HLA-G is differentially expressed in thyroid tissues.
HLA-G peptide preferences change in transformed cells: impact on the binding motif.
Increased soluble human leukocyte antigen-G levels in peripheral blood from climbers on Mount Everest.
Recipient HLA-G +3142 CC Genotype and Concentrations of Soluble HLA-G Impact on Occurrence of CMV Infection after Living-Donor Kidney Transplantation.
Role of HLA-G in innate immunity through direct activation of NF-kappaB in natural killer cells.
Human leukocyte antigen-G in cancer: are they clinically relevant?
Association of HLA-G*01:01:02:01/G*01:04:01 polymorphism with gastric adenocarcinoma.
Analysis of HLA-G expression in malignant hematopoietic cells from leukemia patients.
T cell infiltration into Ewing sarcomas is associated with local expression of immune-inhibitory HLA-G.
Clinical relevance and functional implications for human leukocyte antigen-g expression in non-small-cell lung cancer.
The microRNA-152/human leukocyte antigen-G axis affects proliferation and immune escape of non-small cell lung cancer cells.
Role of HLA-G in tumor escape through expansion of myeloid-derived suppressor cells and cytokinic balance in favor of Th2 versus Th1/Th17.
Switch of HLA-G alternative splicing in a melanoma cell line causes loss of HLA-G1 expression and sensitivity to NK lysis.
Analysis of HLA-G gene polymorphism and protein expression in invasive breast ductal carcinoma.
Analysis of HLA antigen expression in benign and malignant melanocytic lesions reveals that upregulation of HLA-G expression correlates with malignant transformation, high inflammatory infiltration and HLA-A1 genotype.
The clinicopathologic significance of the expression of HLA-G in oral squamous cell carcinoma.
Epigenetic modification augments the immunogenicity of human leukocyte antigen G serving as a tumor antigen for T cell-based immunotherapy.
Human leukocyte antigen-G (HLA-G) as a marker for diagnosis, prognosis and tumor immune escape in human malignancies.
Functional role of human leukocyte antigen-G up-regulation in renal cell carcinoma.
HLA-G is a component of the chronic lymphocytic leukemia escape repertoire to generate immune suppression: impact of the HLA-G 14 base pair (rs66554220) polymorphism.

Clinical and biological significance of HLA-G expression in ovarian cancer.
HLA-G expression in acute lymphoblastic leukemia: a significant prognostic tumor biomarker.
HLA-G protein expression in colorectal cancer evaluated by immunohistochemistry and western blot analysis: Its expression characteristics remain enigmatic.
Implication of Soluble HLA-G and HLA-G +3142G/C Polymorphism in Breast Cancer Patients Receiving Adjuvant Therapy in Tanzania.
Targeting human leukocyte antigen G with chimeric antigen receptors of natural killer cells convert immunosuppression to ablate solid tumors.
Human leukocyte antigen G expression in breast cancer: role in immunosuppression.
Altered HLA class I and HLA-G expression is associated with IL-10 expression in patients with cervical cancer.
A comparative study on regulation of HLA-G expression in bad obstetric history and in head and neck squamous cell carcinoma from Northeast India.
HLA-G inhibition of NK-cell cytolytic function is uncoupled from tumor cell lipid raft reorganization.
High Expression of Human Leukocyte Antigen-G is Associated with a Poor Prognosis in Patients with PDAC.
Inhibiting HLA-G restores IFN- $\gamma$ and TNF- $\alpha$ producing T cell in pleural Tuberculosis.
Characterization of the expression and immunological impact of the transcriptional activator CREB in renal cell carcinoma.
Expression of HLA-G in malignant mesothelioma and clinically aggressive breast carcinoma.
HLA-G and IL-10 expression in human cancer--different stories with the same message.
HLA-G molecules and clinical outcome in Chronic Myeloid Leukemia.
Roles of HLA-G/KIR2DL4 in Breast Cancer Immune Microenvironment.
Human leukocyte antigen-G polymorphism in relation to expression, function, and disease.
Inhibition of host immune response in colorectal cancer: human leukocyte antigen-G and beyond.
Human leukocyte antigen-G in gynaecological tumours.
In vivo evidence that secretion of HLA-G by immunogenic tumor cells allows their evasion from immunosurveillance.
HLA-G 3'UTR Polymorphisms Impact the Prognosis of Stage II-III CRC Patients in Fluoropyrimidine-Based Treatment.
HLA-G and classical HLA class I expression in primary colorectal cancer and associated liver metastases.
PD1/PD1L pathway, HLA-G and T regulatory cells as new markers of immunosuppression in cancers.
Immunoregulatory function of HLA-G in gastric cancer.
Certain haplotypes of the 3'-UTR region of the HLA-G gene are linked to breast cancer.
HLA-G Neo-Expression on Tumors.
Hide-and-seek in the brain: a role for HLA-G mediating immune privilege for glioma cells.
Human leukocyte antigen-G protein expression is an unfavorable prognostic predictor of hepatocellular carcinoma following curative resection.
Induction of cell surface human leukocyte antigen-G expression in pandemic H1N1 2009 and seasonal H1N1 influenza virus-infected patients.
HLA-G in the skin--friend or foe?
HLA-G 14 bp In/Del and +3142 C/G genotypes are differentially expressed between patients with grade IV gliomas and controls.
HLA-G 3' untranslated region polymorphic sites associated with increased HLA-G production are more frequent in patients exhibiting differentiated thyroid tumours.



The Influence of Cytomegalovirus on Expression of HLA-G and its Ligand KIR2DL4 by Human Peripheral Blood Leucocyte Subsets.
Expression of the Classical and Nonclassical HLA Molecules in Breast Cancer.
HLA-G: from biology to clinical benefits.
Retinoblastoma: expression of HLA-G.
Association between the HLA-G molecule and lymph node metastasis in papillary thyroid cancer.
Tumor border sharpness correlates with HLA-G expression in low-grade gliomas.
The Human Leukocyte Antigen G as an Immune Escape Mechanism and Novel Therapeutic Target in Urological Tumors.
TGF- $\beta$ induces HLA-G expression through inhibiting miR-152 in gastric cancer cells.
$\beta$ -defensin-3 negatively regulates TLR4-HMGB1 axis mediated HLA-G expression in IL-1 $\beta$ treated glioma cells.
Intercellular transfer of HLA-G: its potential in cancer immunology.
Increase in HLA-G1 proteolytic shedding by tumor cells: a regulatory pathway controlled by NF-kappaB inducers.
Possible roles of HLA-G regulating immune cells in pregnancy and endometrial diseases via KIR2DL4.
A specific interferon (IFN)-stimulated response element of the distal HLA-G promoter binds IFN-regulatory factor 1 and mediates enhancement of this nonclassical class I gene by IFN-beta.
Major histocompatibility complex abnormalities in non-Hodgkin lymphomas.
HLA-G as a prognostic marker in stage II/III colorectal cancer: not quite there yet.
HLA-G in B-chronic lymphocytic leukaemia: clinical relevance and functional implications.
Decreased intraindividual HLA class I expression is due to reduced transcription in advanced melanoma and does not correlate with HLA-G expression.
Human leukocyte antigen G is associated with esophageal squamous cell carcinoma progression and poor prognosis.
Immune-refractory cancers and their little helpers--an extended role for immunetolerogenic MHC molecules HLA-G and HLA-E?
Vesicular-Bound HLA-G as a Predictive Marker for Disease Progression in Epithelial Ovarian Cancer.
HLA-G expression in Merkel cell carcinoma and the correlation with Merkel cell polyomavirus infection.
Expression analysis of immune-regulatory molecules HLA-G, HLA-E and IDO in endometrial cancer.
Total expression of HLA-G and TLR-9 in chronic lymphocytic leukemia patients.
HLA-G up-regulates ILT2, ILT3, ILT4, and KIR2DL4 in antigen presenting cells, NK cells, and T cells.
Soluble HLA-G (sHLA-G) measurement might be useful as an early diagnostic biomarker and screening test for gastric cancer.
High level of soluble human leukocyte antigen (HLA)-G at beginning of pregnancy as predictor of risk of malaria during infancy.
Soluble HLA-G molecules are increased in lymphoproliferative disorders.
HLA-G and HLA-E Immune Checkpoints Are Widely Expressed in Ewing Sarcoma but Have Limited Functional Impact on the Effector Functions of Antigen-Specific CAR T Cells.
The prognostic impact of soluble and vesicular HLA-G and its relationship to circulating tumor cells in neoadjuvant treated breast cancer patients.
Classical and non-classical HLA class I aberrations in primary cervical squamous- and adenocarcinomas and paired lymph node metastases.
Correlation of serum sHLA-G levels with cyst stage in patients with cystic echinococcosis: is it an immune evasion strategy?

Contribution of HLA-G and FOXP3 genes and proteins in the severity of cervical intraepithelial neoplasia during HPV infection.
Non-classical HLA-G antigen and its role in the cancer progression.
Natural Killer Cell Cytotoxicity Against SKOV3 after HLA-G Downregulation by shRNA.
Prognostic value of HLA class I, HLA-E, HLA-G and Tregs in rectal cancer: a retrospective cohort study. [HLA-G: from feto-maternal tolerance to organ acceptance].
Ionizing radiation modulates the surface expression of human leukocyte antigen-G in a human melanoma cell line.
Human leukocyte antigen G polymorphism is associated with an increased risk of invasive cancer of the uterine cervix.
Human neuroblastoma cells trigger an immunosuppressive program in monocytes by stimulating soluble HLA-G release.
Late-onset intrauterine growth restriction and HHV-6 infection: A pilot study.
Case-control study of HLA-G promoter methylation status, HPV infection and cervical neoplasia in Curitiba, Brazil: a pilot analysis.
Immune response and evasion mechanisms in lip carcinogenesis: An immunohistochemical study.
Expression of classical human leukocyte antigen class I antigens, HLA-E and HLA-G, is adversely prognostic in pancreatic cancer patients.
A comparison of cancer stem cell markers and nonclassical major histocompatibility complex antigens in colorectal tumor and noncancerous tissues.
Relevance of HLA-G, HLA-E and IL-10 expression in lip carcinogenesis.
HLA-G 3' untranslated region gene variants are promising prognostic factors for BK polyomavirus replication and acute rejection after living-donor kidney transplant.
HLA-G protein expression as a potential immune escape mechanism in classical Hodgkin's lymphoma.
Altered pattern of major histocompatibility complex expression in renal carcinoma: tumor-specific expression of the nonclassical human leukocyte antigen-G molecule is restricted to clear cell carcinoma while up-regulation of other major histocompatibility complex antigens is primarily distributed in all subtypes of renal carcinoma.
The 14bp-deletion allele in the HLA-G gene confers susceptibility to the development of hepatocellular carcinoma in the Brazilian population.
Inhibition of iNKT Cells by the HLA-G-ILT2 Checkpoint and Poor Stimulation by HLA-G-Expressing Tolerogenic DC.
Immune Modulation in HLA-G Expressing Head and Neck Squamous Cell Carcinoma in Relation to Human Papilloma Virus Positivity: A Study From Northeast India.
Exosomes bearing HLA-G are released by melanoma cells.
Human leukocyte antigen-G donor-recipient matching of the 14-base pair polymorphism protects against cancer after heart transplant.
Immunohistochemical evaluation of HLA-G and FoxP3+ T regulatory cells in oral cavity and lower lip squamous cell carcinomas.
Combined analysis of HLA class I, HLA-E and HLA-G predicts prognosis in colon cancer patients.
Long non-coding RNA HOTAIR promotes HLA-G expression via inhibiting miR-152 in gastric cancer cells.
HLA-E and HLA-F Are Overexpressed in Glioblastoma and HLA-E Increased After Exposure to Ionizing Radiation.
Soluble HLA-G is a differential prognostic marker in sequential colorectal cancer disease stages.
Between Innate and Adaptive Immune Responses: NKG2A, NKG2C, and CD8 <sup>+</sup> T Cell Recognition of HLA-E Restricted Self-Peptides Acquired in the Absence of HLA-Ia.

Immunohistochemical investigations on the expression of programmed cell death ligand 1, human leukocyte antigens G and E, and granzyme B in intraoral mucoepidermoid carcinoma.
Immunosuppressive mediators of oral squamous cell carcinoma in tumour samples and saliva.
The association between genetic variants at 3'-UTR and 5'-UTR of HLA-G gene and the clinical outcomes of patients with leukemia receiving hematopoietic stem cell transplantation.
Prognostic significance of high circulating sHLA-G in ovarian carcinoma.
HLA-E expression and its clinical relevance in human renal cell carcinoma.
Soluble HLA-G molecules increase during acute leukemia, especially in subtypes affecting monocytic and lymphoid lineages.
Soluble human leukocyte antigen -G during pregnancy and infancy in Benin: Mother/child resemblance and association with the risk of malaria infection and low birth weight.
Personalized HLA typing leads to the discovery of novel HLA alleles and tumor-specific HLA variants.
Primary HIV-1 Strains Use Nef To Downmodulate HLA-E Surface Expression.
HLA and tumour immunology: immune escape, immunotherapy and immune-related adverse events.
The diversity of the HLA-E-restricted peptide repertoire explains the immunological impact of the Arg107Gly mismatch.
Is immune escape via human leukocyte antigen expression clinically relevant in chronic lymphocytic leukemia? Focus on the controversies.
HLA-mediated tumor escape mechanisms that may impair immunotherapy clinical outcomes via T-cell activation.
Classic and nonclassic HLA class I expression in penile cancer and relation to HPV status and clinical outcome.
Protein disulfide isomerase A1 regulates breast cancer cell immunorecognition in a manner dependent on redox state.
Human leukocyte antigen-G and cancer immunoediting.
Soluble HLA-G and HLA-G Bearing Extracellular Vesicles Affect ILT-2 Positive and ILT-2 Negative CD8 T Cells Complementary.
Elevated sHLA-G plasma levels post chemotherapy combined with ILT-2 rs10416697C allele status of the sHLA-G-related receptor predict poorest disease outcome in early triple-negative breast cancer patients.
Immune microenvironment and evasion mechanisms in adenoid cystic carcinomas of salivary glands.
HLA-G/LILRBs: A Cancer Immunotherapy Challenge.
Clinical significance of HLA-E genotype and surface/soluble expression levels between healthy individuals and patients with acute leukemia.
Human leukocyte antigens in cancer metastasis: Prognostic approach and therapeutic susceptibility.
Human leukocyte antigen-E alleles and expression in patients with serous ovarian cancer.
Human leukocyte antigen-G 14-bp InDel polymorphism and oral squamous cell carcinoma risk in Chinese Han population: A case-control study.
Clinical Characteristics and Outcome Analysis for HLA Loss Patients Following Partially Mismatched Related Donor Transplantation Using HLA Chimerism for Loss of Heterozygosity Analysis by Next-Generation Sequencing.
Leptin promotes HLA-G expression on placental trophoblasts via the MEK/Erk and PI3K signaling pathways.
Integration of high-risk human papillomavirus DNA correlates with HLA genotype aberration and reduced HLA class I molecule expression in human cervical carcinoma.
Generation of a blockage monoclonal antibody of LILRB1 against HLA-G.
Advances in the study of HLA class Ib in maternal-fetal immune tolerance.

Interleukin-37 gene polymorphism and susceptibility to pulmonary tuberculosis among Iraqi patients.
High levels of soluble MICA are significantly related to increased disease-free and disease-specific survival in patients with cervical adenocarcinoma.
HLA-E allelic genotype correlates with HLA-E plasma levels and predicts early progression in chronic lymphocytic leukemia.
HLA-G in skin cancer: a wolf in sheep's clothing?
Human leukocyte antigen E in human cytomegalovirus infection: friend or foe?
Involvement of IL-10 and TGF- $\beta$ in HLA-E-mediated neuroblastoma migration and invasion.
Co-expression of HLA-I loci improved prognostication in HER2+ breast cancers.
Impact of HLA-E gene polymorphism on HLA-E expression in tumor cells and prognosis in patients with stage III colorectal cancer.
Toward eliminating HLA class I expression to generate universal cells from allogeneic donors.
HLA-E protects glioma cells from NKG2D-mediated immune responses in vitro: implications for immune escape in vivo.
HLA-J, a Non-Pseudogene as a New Prognostic Marker for Therapy Response and Survival in Breast Cancer.
HLA Allele E*01:01 Is Associated with a Reduced Risk of EBV-Related Classical Hodgkin Lymphoma Independently of HLA-A*01/*02.
European Patent in Immunoncology: From Immunological Principles of Implantation to Cancer Treatment.
Genetically engineered blood pharming: generation of HLA-universal platelets derived from CD34+ progenitor cells.
HLA-E/ $\beta$ 2 microglobulin overexpression in colorectal cancer is associated with recruitment of inhibitory immune cells and tumor progression.
Dynamic Alteration in HLA-E Expression and Soluble HLA-E via Interaction with Natural Killer Cells in Gastric Cancer.
Loss of Fas expression and high expression of HLA-E promoting the immune escape of early colorectal cancer cells.
Incidence, risk factors and clinical outcome of leukemia relapses with loss of the mismatched HLA after partially incompatible hematopoietic stem cell transplantation.
Serum sHLA-G levels: a useful indicator in distinguishing colorectal cancer from benign colorectal diseases.
Comprehensive analysis of cancer-associated somatic mutations in class I HLA genes.
Soluble HLA revisited.
Allele-Specific HLA Loss and Immune Escape in Lung Cancer Evolution.
Is there a role played by HLA-E, if any, in HPV immune evasion?
Novel deletion mutation of HLA-B*40:02 gene in acquired aplastic anemia.
Review: HLA loss and detection in the setting of relapse from HLA-mismatched hematopoietic cell transplant.
Galectin-1 influences trophoblast immune evasion and emerges as a predictive factor for the outcome of pregnancy.
Presentation of a Conserved Adenoviral Epitope on HLA-C*0702 Allows Evasion of Natural Killer but Not T Cell Responses.
HLA-universal platelet transfusions prevent platelet refractoriness in a mouse model.
GPER mediated estradiol reduces miR-148a to promote HLA-G expression in breast cancer.
Individual HLA heterogeneity and its implications for cellular immune evasion in cancer and beyond.
Cancer and pregnancy share similar mechanisms of immunological escape.

Long non-coding RNA HOTAIR modulates HLA-G expression by absorbing miR-148a in human cervical cancer.
High-Risk Human Papillomavirus E7 Alters Host DNA Methylome and Represses HLA-E Expression in Human Keratinocytes.
Cancer treatment and the KIR-HLA system: an overview.
HLA mismatching as a strategy to reduce relapse after alternative donor transplantation.
Overexpression of immunomodulatory mediators in oral precancerous lesions.
Cetuximab-mediated cellular cytotoxicity is inhibited by HLA-E membrane expression in colon cancer cells.
Characterization and clinical enrichment of HLA-C*07:02-restricted Cytomegalovirus-specific CD8+ T cells.
Genetic polymorphism of NK receptors and their ligands in melanoma patients: prevalence of inhibitory over activating signals.
The Presence of HLA-E-Restricted, CMV-Specific CD8+ T Cells in the Blood of Lung Transplant Recipients Correlates with Chronic Allograft Rejection.
Immunogenicity and Immune Silence in Human Cancer.
Immunosuppressive microenvironment in neuroblastoma.
HLA-E expression in cervical adenocarcinomas: association with improved long-term survival.
Variations in BK Polyomavirus Immunodominant Large Tumor Antigen-Specific 9mer CD8 T-Cell Epitopes Predict Altered HLA-Presentation and Immune Failure.
Serum soluble HLA-E in melanoma: a new potential immune-related marker in cancer.
Role of HLA-G and extracellular vesicles in renal cancer stem cell-induced inhibition of dendritic cell differentiation.
Assessment of Patient-Specific Human Leukocyte Antigen Genomic Loss at Relapse After Antithymocyte Globulin-Based T-Cell-Replete Haploidentical Hematopoietic Stem Cell Transplant.
Insignificant effects of loss of heterozygosity in HLA in the efficacy of immune checkpoint blockade treatment.
HLA-DR signaling inhibits Fas-mediated apoptosis in A375 melanoma cells.
Loss of Human Leukocyte Antigen and Immune Escape in Head and Neck Cancer.
Immune Editing: Overcoming Immune Barriers in Stem Cell Transplantation.
Integrated investigation of the prognostic role of HLA LOH in advanced lung cancer patients with immunotherapy.
Non-classical major histocompatibility complex proteins as determinants of tumour immunosurveillance.
Clinical Utility of the Detection of the Loss of the Mismatched HLA in Relapsed Hematological Patients After Haploidentical Stem Cell Transplantation With High-Dose Cyclophosphamide.
Immunological properties of extraembryonic human mesenchymal stromal cells derived from gestational tissue.
No cancer in cancers: evolutionary trade-off between successful viviparity and tumor escape from the adaptive immune system.
Senescent cells evade immune clearance via HLA-E-mediated NK and CD8(+) T cell inhibition.
Suppression of HLA expression by lentivirus-mediated gene transfer of siRNA cassettes and in vivo chemoselection to enhance hematopoietic stem cell transplantation.
[Associations of HLA gene with leukemia in 1186 cases].
Enriched HLA-E and CD94/NKG2A Interaction Limits Antitumor CD8(+) Tumor-Infiltrating T Lymphocyte Responses.

Expression of HLA-DR is reduced in tumor infiltrating immune cells (TIICs) and regional lymph nodes of non-small-cell lung carcinomas. A putative mechanism of tumor-induced immunosuppression?
Dual IFN- $\gamma$ /hypoxia priming enhances immunosuppression of mesenchymal stromal cells through regulatory proteins and metabolic mechanisms.
A cancer-array approach elucidates the immune escape mechanism and defects in the DNA repair system in esophageal squamous cell carcinoma.
HLA Loss Facilitates Immune Escape.
Are HLA DQB1 alleles correlated with breast cancer histopronostic parameters in Tunisia?
Predisposition to HPV16/18-related cervical cancer because of proline homozygosity at codon 72 of p53 among Indian women is influenced by HLA-B*07 and homozygosity of HLA-DQB1*03.
Comparing effects of BK virus agnoprotein and herpes simplex-1 ICP47 on MHC-I and MHC-II expression.
Bone marrow-infiltrating human neuroblastoma cells express high levels of calprotectin and HLA-G proteins.
Loss of mismatched HLA in acute myeloid leukemia relapse after haploidentical peripheral blood stem cell transplantation combined with unrelated cord blood: A case report.
Direct identification of an HPV-16 tumor antigen from cervical cancer biopsy specimens.
Recurrent genetic HLA loss in AML relapsed after matched unrelated allogeneic hematopoietic cell transplantation.
Selective deletion of human leukocyte antigens protects stem cell-derived islets from immune rejection.
Open conformers of HLA-F are high-affinity ligands of the activating NK-cell receptor KIR3DS1.
Genomic loss of HLA alleles may affect the clinical outcome in low-risk myelodysplastic syndrome patients.
Generation of HLA Universal Megakaryocytes and Platelets by Genetic Engineering.
Immunomodulatory strategies for relapse after haploidentical hematopoietic stem cell transplantation in hematologic malignancy patients.
Mechanisms of Leukemia Immune Evasion and Their Role in Relapse After Haploidentical Hematopoietic Cell Transplantation.
G Protein-Coupled receptors and heterotrimeric G proteins as cancer drivers.
Genomic loss of mismatched human leukocyte antigen and leukemia immune escape from haploidentical graft-versus-leukemia.
HLA-G1-expressing antigen-presenting cells induce immunosuppressive CD4+ T cells.
Frequent mutations in HLA and related genes in extranodal NK/T cell lymphomas.
Escape of leukemia blasts from HLA-specific CTL pressure in a recipient of HLA one locus-mismatched bone marrow transplantation.
Graft-versus-leukemia effect of HLA-haploidentical central-memory T-cells expanded with leukemic APCs and modified with a suicide gene.
Immunosuppressive activity of CD14+ HLA-DR- cells in squamous cell carcinoma of the head and neck.
Extramedullary leukemia relapse after allogeneic stem cell transplantation: a novel mechanism of immune escape?
Trogocytic intercellular membrane exchanges among hematological tumors.
Lack of detectable neoantigen depletion signals in the untreated cancer genome.
Association of genomic variants at the human leukocyte antigen locus with cervical cancer risk, HPV status and gene expression levels.
Indoleamine 2,3-dioxygenase activity as a potential biomarker of immune suppression during visceral leishmaniasis.

Tracking the Genetic Susceptibility Background of B-Cell Non-Hodgkin's Lymphomas from Genome-Wide Association Studies.
Somatic Mutations in Aplastic Anemia.
A novel clinically relevant graft-versus-leukemia model in humanized mice.
Increased blood CD226(-) inflammatory monocytes with low antigen presenting potential correlate positively with severity of hemorrhagic fever with renal syndrome.
Donor-derived acute myeloid leukemia in solid organ transplantation.

Table A2-5, Cluster 4

Cluster 4 focuses on hepatitis C virus, especially on the role of immune system evasion in hcv persistence (375)
Innate immune responses in hepatitis C virus infection.
Immune modulation by the hepatitis C virus core protein.
A new insight into hepatitis C vaccine development.
Experimental models to study the immunobiology of hepatitis C virus.
Immunogenetic studies of the hepatitis C virus infection in an era of pan-genotype antiviral therapies - Effective treatment is coming.
Activation and evasion of antiviral innate immunity by hepatitis C virus.
Extracellular Interactions between Hepatitis C Virus and Secreted Apolipoprotein E.
Mechanisms and Consequences of Genetic Variation in Hepatitis C Virus (HCV).
Status of hepatitis C virus vaccination: Recent update.
The hepatitis C virus persistence: how to evade the immune system?
Hepatitis C virus evasion of adaptive immune responses: a model for viral persistence.
Hepatitis C virus infection inhibits a Src-kinase regulatory phosphatase and reduces T cell activation in vivo.
Direct effects of hepatitis C virus on the lymphoid cells.
Hepatitis C virus strategies to evade the specific-T cell response: a possible mission favoring its persistence.
Innate immunity against hepatitis C virus.
Inhibition of the interferon antiviral response by hepatitis C virus.
Programmed death-1/programmed death-L1 signaling pathway and its blockade in hepatitis C virus immunotherapy.
Subversion of immune responses by hepatitis C virus: immunomodulatory strategies beyond evasion?
Hepatitis C virus-mediated modulation of cellular immunity.
Immune responses to HCV and other hepatitis viruses.
High-programmed death-1 levels on hepatitis C virus-specific T cells during acute infection are associated with viral persistence and require preservation of cognate antigen during chronic infection.
Toll-like receptors in hepatitis C infection: implications for pathogenesis and treatment.
Natural killer cells in hepatitis C: Current progress.
Multi-step regulation of interferon induction by hepatitis C virus.
Immunopathogenesis of hepatitis C virus infection.
Hepatitis C Virus Genetic Variability, Human Immune Response, and Genome Polymorphisms: Which Is the Interplay?
Hepatitis C Virus: Evading the Intracellular Innate Immunity.
Hepatitis C virus infection: when silence is deception.

Mononuclear phagocyte system in hepatitis C virus infection.
Natural killer cells: primary target for hepatitis C virus immune evasion strategies?
Clinical significance of hepatitis C virus genotypes and quasispecies.
The hepatitis C virus glycan shield and evasion of the humoral immune response.
Hepatitis C virus targets the T cell secretory machinery as a mechanism of immune evasion.
Hepatitis C virus and antiviral innate immunity: who wins at tug-of-war?
Hepatitis C Virus Glycan-Dependent Interactions and the Potential for Novel Preventative Strategies.
The induction of type I interferon production in hepatitis C-infected patients.
Structure-Based and Rational Design of a Hepatitis C Virus Vaccine.
Interaction between the hepatitis C virus and the immune system.
microRNA-125a targets MAVS and TRAF6 to modulate interferon signaling and promote HCV infection.
Neutralizing antibodies and broad, functional T cell immune response following immunization with hepatitis C virus proteins-based vaccine formulation.
Inter and inpatient evolution of hepatitis C virus.
A Hepatitis C Virus Envelope Polymorphism Confers Resistance to Neutralization by Polyclonal Sera and Broadly Neutralizing Monoclonal Antibodies.
The role of human lipoproteins for hepatitis C virus persistence.
Chronic HCV-related autoimmunity: a consequence of viral persistence and lymphotropism.
Hepatitis C virus triggers apoptosis of a newly developed hepatoma cell line through antiviral defense system.
Hepatitis C virus--T-cell responses and viral escape mutations.
Evasion of host immune surveillance by hepatitis C virus: potential roles in viral persistence.
Hepatitis C Virus. Strategies to Evade Antiviral Responses.
Update on hepatitis C virus-specific immunity.
Hepatitis C virus infection: establishment of chronicity and liver disease progression.
Structural perspectives on HCV humoral immune evasion mechanisms.
A Role for B Cells to Transmit Hepatitis C Virus Infection.
Hepatitis C virus infection: virus/host interactions.
Hepatitis C virus evasion mechanisms from neutralizing antibodies.
The genetics of hepatitis C virus underlie its ability to escape humoral immunity.
Hepatitis C Virus (HCV)-Apolipoprotein Interactions and Immune Evasion and Their Impact on HCV Vaccine Design.
Toll-like receptors and hepatitis C virus infection.
Roles of the two distinct proteasome pathways in hepatitis C virus infection.
Enhanced IL-10 production in response to hepatitis C virus proteins by peripheral blood mononuclear cells from human immunodeficiency virus-monoinfected individuals.
Chronic hepatitis C.
Innate and Adaptive Immune Responses in Chronic HCV Infection.
The Role of ApoE in HCV Infection and Comorbidity.
Hepatitis C virus: immunosuppression by complement regulatory pathway.
Factors that determine the antiviral efficacy of HCV-specific CD8(+) T cells ex vivo.
Modeling and analysis of innate immune responses induced by the host cells against hepatitis C virus infection.
CXCL9 chemokine level is associated with spontaneous clearance and sustained virological response in Egyptian Chronic Hepatitis C patients receiving direct acting antivirals.



Adaptive immunity to the hepatitis C virus.
Regulation of hepatic innate immunity by hepatitis C virus.
Tupaia MAVS Is a Dual Target during Hepatitis C Virus Infection for Innate Immune Evasion and Viral Replication via NF- $\kappa$ B.
CD8 epitope escape and reversion in acute HCV infection.
Network based analysis of hepatitis C virus core and NS4B protein interactions.
Specialization of Hepatitis C Virus Envelope Glycoproteins for B Lymphocytes in Chronically Infected Patients.
Hepatitis C virus core activates proteasomal activator 28 $\gamma$ expression via upregulation of p53 levels to control virus propagation.
Immunomodulation by hepatitis C virus-derived proteins: targeting human dendritic cells by multiple mechanisms.
HCV inhibits antigen processing and presentation and induces oxidative stress response in gastric mucosa.
Synthesized peptides 705-734 from hepatitis C virus E2 glycoprotein induce dendritic cell maturation by activating p38 MAPK signaling.
Failure of innate and adaptive immune responses in controlling hepatitis C virus infection.
Therapeutic control of hepatitis C virus: the role of neutralizing monoclonal antibodies.
Three different functional microdomains in the hepatitis C virus hypervariable region 1 (HVR1) mediate entry and immune evasion.
A Novel Approach To Display Structural Proteins of Hepatitis C Virus Quasispecies in Patients Reveals a Key Role of E2 HVR1 in Viral Evolution.
HCV Interplay with Lipoproteins: Inside or Outside the Cells?
Immune system control of hepatitis C virus infection.
Apolipoprotein E Mediates Evasion From Hepatitis C Virus Neutralizing Antibodies.
HCV viremia drives an increment of CD86 expression by myeloid dendritic cells.
An overview about hepatitis C: a devastating virus.
HCV Defective Genomes Promote Persistent Infection by Modulating the Viral Life Cycle.
Hepatitis C virus epitope-specific neutralizing antibodies in Igs prepared from human plasma.
Immunology of hepatitis B virus and hepatitis C virus infection.
Development of a heterologous, multigenotype vaccine against hepatitis C virus infection.
Mutations that alter use of hepatitis C virus cell entry factors mediate escape from neutralizing antibodies.
Hepatitis C virus infection suppresses the interferon response in the liver of the human hepatocyte chimeric mouse.
The HCV life cycle: in vitro tissue culture systems and therapeutic targets.
The dynamics of T-lymphocyte responses during combination therapy for chronic hepatitis C virus infection.
Host-virus interactions in hepatitis B and hepatitis C infection.
Incorporation of hepatitis C virus E1 and E2 glycoproteins: the keystones on a peculiar virion.
Innate detection of hepatitis B and C virus and viral inhibition of the response.
Dendritic Cells in HIV-1 and HCV Infection: Can They Help Win the Battle?
Can Broadly Neutralizing Monoclonal Antibodies Lead to a Hepatitis C Virus Vaccine?
Hepatitis C virus-infected cells downregulate NKp30 and inhibit ex vivo NK cell functions.
Impaired expression and function of toll-like receptor 7 in hepatitis C virus infection in human hepatoma cells.

Immune control and failure in HCV infection--tipping the balance.
ISG56 and IFITM1 proteins inhibit hepatitis C virus replication.
Modulation of the Immune System in Chronic Hepatitis C and During Antiviral Interferon-Free Therapy.
Hepatitis C virus attenuates interferon-induced major histocompatibility complex class I expression and decreases CD8+ T cell effector functions.
Glycan shifting on hepatitis C virus (HCV) E2 glycoprotein is a mechanism for escape from broadly neutralizing antibodies.
Neutralizing antibodies and pathogenesis of hepatitis C virus infection.
Emerging concepts in immunity to hepatitis C virus infection.
Structural basis of hepatitis C virus neutralization by broadly neutralizing antibody HCV1.
Structural Versatility of Hepatitis C Virus Proteins: Implications for the Design of Novel Anti-HCV Intervention Strategies.
Hepatitis C virus impairs natural killer cell activity via viral serine protease NS3.
[Immune evasion mechanisms of hepatitis C virus. Review].
HLA-G 3'UTR haplotype analyses in HCV infection and HCV-derived cirrhosis, hepatocellular carcinoma and fibrosis.
Antigen-driven patterns of TCR bias are shared across diverse outcomes of human hepatitis C virus infection.
Ficolin-2 inhibits hepatitis C virus infection, whereas apolipoprotein E3 mediates viral immune escape.
Hepatitis C virions subvert natural killer cell activation to generate a cytokine environment permissive for infection.
HCV innate immune responses.
Hepatitis C Virus Manipulates Humans as its Favorite Host for a Long-Term Relationship.
The role of humoral innate immunity in hepatitis C virus infection.
Hepatitis C virus vaccine candidates inducing protective neutralizing antibodies.
Positive selection of cytotoxic T lymphocyte escape variants during acute hepatitis C virus infection.
Induction of Selenoprotein P mRNA during Hepatitis C Virus Infection Inhibits RIG-I-Mediated Antiviral Immunity.
Systems biology analyses to define host responses to HCV infection and therapy.
The role of the hepatitis C virus glycoproteins in infection.
Effect of immune pressure on hepatitis C virus evolution: insights from a single-source outbreak.
Characterization of antigenic variants of hepatitis C virus in immune evasion.
A Review of Hepatitis B Virus and Hepatitis C Virus Immunopathogenesis.
The heat shock cognate protein 70 is associated with hepatitis C virus particles and modulates virus infectivity.
Hepatitis C virus entry into hepatocytes: molecular mechanisms and targets for antiviral therapies.
Viral entry and escape from antibody-mediated neutralization influence hepatitis C virus reinfection in liver transplantation.
Tim-3 expression on PD-1+ HCV-specific human CTLs is associated with viral persistence, and its blockade restores hepatocyte-directed in vitro cytotoxicity.
Intra-host analysis of hepaciviral glycoprotein evolution reveals signatures associated with viral persistence and clearance.
Antigenic cooperation among intrahost HCV variants organized into a complex network of cross-immunoreactivity.

Influence of HIV and HCV on T cell antigen presentation and challenges in the development of vaccines.
Set7 facilitates hepatitis C virus replication via enzymatic activity-dependent attenuation of the IFN-related pathway.
A bioinformatics pipeline for the analyses of viral escape dynamics and host immune responses during an infection.
Dendritic cells, regulatory T cells and the pathogenesis of chronic hepatitis C.
The hepatitis C virus enigma.
Hepatitis C virus NS4B blocks the interaction of STING and TBK1 to evade host innate immunity.
Hepatitis C virus: the role of molecular mimicry in response to interferon treatment.
Expression of hepatitis C virus proteins does not interfere with major histocompatibility complex class I processing and presentation in vitro.
Naturally selected hepatitis C virus polymorphisms confer broad neutralizing antibody resistance.
The Role of Cytosolic Lipid Droplets in Hepatitis C Virus Replication, Assembly, and Release.
Structure-Based Design of Hepatitis C Virus Vaccines That Elicit Neutralizing Antibody Responses to a Conserved Epitope.
Computational Modeling of Hepatitis C Virus Envelope Glycoprotein Structure and Recognition.
Loss of immune escape mutations during persistent HCV infection in pregnancy enhances replication of vertically transmitted viruses.
Sequence evolution of putative cytotoxic T cell epitopes in NS3 region of hepatitis C virus.
The neutralizing activity of anti-hepatitis C virus antibodies is modulated by specific glycans on the E2 envelope protein.
Escape mutations alter proteasome processing of major histocompatibility complex class I-restricted epitopes in persistent hepatitis C virus infection.
Interferon-Induced Transmembrane Proteins Mediate Viral Evasion in Acute and Chronic Hepatitis C Virus Infection.
CD81(+) Exosomes Play a Pivotal Role in the Establishment of Hepatitis C Persistent Infection and Contribute Toward the Progression of Hepatocellular Carcinoma.
[Immune dysregulation mediated by hepatitis C virus infection].
Efficient acute and chronic infection of stem cell-derived hepatocytes by hepatitis C virus.
Recombination in hepatitis C virus is not uncommon among people who inject drugs in Kolkata, India.
Upregulation of soluble and membrane-bound human leukocyte antigen G expression is primarily observed in the milder histopathological stages of chronic hepatitis C virus infection.
[The cytokine secretion of peripheral blood mononucleocytes from patients infected with HCV].
Evolutionary dynamics of hepatitis C virus envelope genes during chronic infection.
Inhibition of protease-inhibitor-resistant hepatitis C virus replicons and infectious virus by intracellular intrabodies.
Mapping Determinants of Virus Neutralization and Viral Escape for Rational Design of a Hepatitis C Virus Vaccine.
Extracellular Vesicle Release Promotes Viral Replication during Persistent HCV Infection.
Regulatory T cells in viral hepatitis.
Hepatitis C Virus-Induced Myeloid-Derived Suppressor Cells Suppress NK Cell IFN- $\gamma$ Production by Altering Cellular Metabolism via Arginase-1.
Characterization of chronic HCV infection-induced apoptosis.
Hepatitis C virus transmission bottlenecks analyzed by deep sequencing.
Recombinant retrovirus-derived virus-like particle-based vaccines induce hepatitis C virus-specific cellular and neutralizing immune responses in mice.

Autophagy induction plays time-dependent role in viral load of HCV infected Huh7.5 cell line.
Immune Evasion Strategies during Chronic Hepatitis B and C Virus Infection.
Elucidating novel hepatitis C virus-host interactions using combined mass spectrometry and functional genomics approaches.
Hepatitis C virus resistance to protease inhibitors.
Apolipoprotein E, but Not Apolipoprotein B, Is Essential for Efficient Cell-to-Cell Transmission of Hepatitis C Virus.
Glycan Shielding and Modulation of Hepatitis C Virus Neutralizing Antibodies.
Virological footprint of CD4+ T-cell responses during chronic hepatitis C virus infection.
Upregulation of major histocompatibility complex class I on liver cells by hepatitis C virus core protein via p53 and TAP1 impairs natural killer cell cytotoxicity.
Immunopathogenesis of hepatitis C virus infection.
Hepatitis C virus suppresses C9 complement synthesis and impairs membrane attack complex function.
Hepatitis C virus induces regulatory T cells by naturally occurring viral variants to suppress T cell responses.
Immune evasion versus recovery after acute hepatitis C virus infection from a shared source.
Increased cytotoxic T-lymphocyte epitope variant cross-recognition and functional avidity are associated with hepatitis C virus clearance.
Bivalent vaccine platform based on Japanese encephalitis virus (JEV) elicits neutralizing antibodies against JEV and hepatitis C virus.
Interaction Between the Neglected Tropical Disease Human Schistosomiasis and HCV Infection in Egypt: a Puzzling Relationship.
Dendritic cell maturation in HCV infection: altered regulation of MHC class I antigen processing-presenting machinery.
Humoral immune system targets clonotypic antibody-associated hepatitis C virus.
Viral hepatitis.
Exosome-mediated transmission of hepatitis C virus between human hepatoma Huh7.5 cells.
Understanding human immunodeficiency virus type 1 and hepatitis C virus coinfection.
Up-regulation of A20/ABIN1 contributes to inefficient M1 macrophage polarization during Hepatitis C virus infection.
Induction of Profibrotic Microenvironment via TLR4 MyD88-Dependent and -Independent Inflammatory Signaling in Chronic Hepatitis C Virus Infection.
Subversion of innate host antiviral strategies by the hepatitis C virus.
Sequence variations of the hypervariable region of hepatitis C virus and their clinical significance.
Hepatitis C Virus-Associated Extrahepatic Manifestations in Lung and Heart and Antiviral Therapy-Related Cardiopulmonary Toxicity.
Glycometabolism regulates hepatitis C virus release.
Hepatitis C virus fails to activate NF-κB signaling in plasmacytoid dendritic cells.
Hepatitis C Virus and Hepatocellular Carcinoma: When the Host Loses Its Grip.
[Chronic hepatitis C infection--mechanisms of virus "immune escape"].
Relation between viral fitness and immune escape within the hepatitis C virus protease.
Extra-epitopic hepatitis C virus polymorphisms confer resistance to broadly neutralizing antibodies by modulating binding to scavenger receptor B1.
Hepatitis C virus: Enslavement of host factors.

Hepatitis C virus envelope glycoprotein signatures are associated with treatment failure and modulation of viral entry and neutralization.
Hepatitis C Virus.
Glances in Immunology of HIV and HCV Infection.
Frequency of Interferon-Resistance Conferring Substitutions in Amino Acid Positions 70 and 91 of Core Protein of the Russian HCV 1b Isolates Analyzed in the T-Cell Epitopic Context.
Control of hepatitis C virus replication in mouse liver-derived cells by MAVS-dependent production of type I and type III interferons.
A single amino acid change in the hypervariable region 1 of hepatitis C virus genotype 4a aids humoral immune escape.
Virus-neutralizing antibodies to hepatitis C virus.
A target on the move: innate and adaptive immune escape strategies of hepatitis C virus.
Constraints on viral evolution during chronic hepatitis C virus infection arising from a common-source exposure.
Impaired clearance of virus-infected hepatocytes in transgenic mice expressing the hepatitis C virus polyprotein.
Challenges to the development of vaccines to hepatitis C virus that elicit neutralizing antibodies.
Hepatitis C virus escape from the interferon regulatory factor 3 pathway by a passive and active evasion strategy.
Activation of Vgamma9Vdelta2 T cells by non-peptidic antigens induces the inhibition of subgenomic HCV replication.
Hepatitis C virus infection upregulates CD55 expression on the hepatocyte surface and promotes association with virus particles.
[VIRAL HEPATITIS C: EVOLUTION OF THE EPIDEMIOLOGIC PROCESS, EVOLUTION OF THE VIRUS].
Hepatitis C Virus NS5A Targets Nucleosome Assembly Protein NAP1L1 To Control the Innate Cellular Response.
Hepatitis C virus, cholesterol and lipoproteins--impact for the viral life cycle and pathogenesis of liver disease.
Distinct Toll-like Receptor 3 and 7 Expression in Peripheral Blood Mononuclear Cells From Patients with Chronic Hepatitis C Infection.
Flying under the radar: the immunobiology of hepatitis C.
Hepatitis C virus core protein induces expression of genes regulating immune evasion and anti-apoptosis in hepatocytes.
Organ system view of the hepatic innate immunity in HCV infection.
Toll-like receptor 3 and 7/8 function is impaired in hepatitis C rapid fibrosis progression post-liver transplantation.
Immunization with Recombinant Adenoviral Vectors Expressing HCV Core or F Proteins Leads to T Cells with Reduced Effector Molecules Granzyme B and IFN- $\gamma$ : A Potential New Strategy for Immune Evasion in HCV Infection.
Hepatitis C virus triggers mitochondrial fission and attenuates apoptosis to promote viral persistence.
Hypervariable Region 1 in Envelope Protein 2 of Hepatitis C Virus: A Linchpin in Neutralizing Antibody Evasion and Viral Entry.
Do antiviral CD8+ T cells select hepatitis C virus escape mutants? Analysis in diverse epitopes targeted by human intrahepatic CD8+ T lymphocytes.
Complement Regulation and Immune Evasion by Hepatitis C Virus.
DDB1 is a cellular substrate of NS3/4A protease and required for hepatitis C virus replication.
Hepatitis C virus modulates signal peptide peptidase to alter host protein processing.

Evolution of viral quasispecies in four dominant HIA-A2 restricted T cell epitopes is not a major reason for viral persistence in interferon-treated patients with chronic hepatitis C.
Characteristics of the intrahepatic cytotoxic T lymphocyte response in chronic hepatitis C virus infection.
Highly heterogeneous mutation rates in the hepatitis C virus genome.
Hepatitis C: clinical management and debated issues.
ISG15, a ubiquitin-like interferon-stimulated gene, promotes hepatitis C virus production in vitro: implications for chronic infection and response to treatment.
Pathogen-Associated Molecular Pattern Recognition of Hepatitis C Virus Transmitted/Founder Variants by RIG-I Is Dependent on U-Core Length.
Identification of antigenic escape variants in an immunodominant epitope of hepatitis C virus.
Selection-driven immune escape is not a significant factor in the failure of CD4 T cell responses in persistent hepatitis C virus infection.
Factors Influencing the Prevalence of Resistance-Associated Substitutions in NS5A Protein in Treatment-Naïve Patients with Chronic Hepatitis C.
Divergent adaptation of hepatitis C virus genotypes 1 and 3 to human leukocyte antigen-restricted immune pressure.
Positional effect of phosphorylation sites 266 and 267 in the cytoplasmic domain of the E2 protein of hepatitis C virus 3a genotype: interferon resistance analysis via sequence alignment.
Distinct Escape Pathway by Hepatitis C Virus Genotype 1a from a Dominant CD8+ T Cell Response by Selection of Altered Epitope Processing.
Comparison of HAV and HCV infections in vivo and in vitro reveals distinct patterns of innate immune evasion and activation.
Conventional protein kinase C inhibition prevents alpha interferon-mediated hepatitis C virus replicon clearance by impairing STAT activation.
Evolution of within-host variants of the hepatitis C virus.
Activation of nuclear factor kappaB in hepatitis C virus infection: implications for pathogenesis and hepatocarcinogenesis.
Limited T cell receptor diversity of HCV-specific T cell responses is associated with CTL escape.
Viral escape and T cell exhaustion in hepatitis C virus infection analysed using Class I peptide tetramers.
A laboratory-adapted HCV JFH-1 strain is sensitive to neutralization and can gradually escape under the selection pressure of neutralizing human plasma.
Combination of three adjuvants enhances the immunogenicity of a recombinant protein containing the CTL epitopes of non-structural proteins of hepatitis C virus.
Variability or conservation of hepatitis C virus hypervariable region 1? Implications for immune responses.
The full length hepatitis C virus polyprotein and interactions with the interferon-beta signalling pathways in vitro.
Applying antibody-sensitive hypervariable region 1-deleted hepatitis C virus to the study of escape pathways of neutralizing human monoclonal antibody AR5A.
Modulation of dendritic cell function by persistent viruses.
Long-term follow-up of chimpanzees inoculated with the first infectious clone for hepatitis C virus.
Inhibition of intrahepatic gamma interferon production by hepatitis C virus nonstructural protein 5A in transgenic mice.
A new system to measure and compare hepatitis C virus replication capacity using full-length, replication competent viruses.

Naturally occurring CD4+ T-cell epitope variants act as altered peptide ligands leading to impaired helper T-cell responses in hepatitis C virus infection.
Ultrastructural Localization and Molecular Associations of HCV Capsid Protein in Jurkat T Cells.
Identification of CTL epitopes in hepatitis C virus by a genome-wide computational scanning and a rational design of peptide vaccine.
Hepatitis C virus immune escape via exploitation of a hole in the T cell repertoire.
Human single chain-transbodies that bound to domain-I of non-structural protein 5A (NS5A) of hepatitis C virus.
Hepatitis C virus dynamics and pathology: the role of CTL and antibody responses.
Role of the E2 Hypervariable Region (HVR1) in the Immunogenicity of a Recombinant Hepatitis C Virus Vaccine.
Effective epitope identification employing phylogenetic, mutational variability, sequence entropy, and correlated mutation analysis targeting NS5B protein of hepatitis C virus: from bioinformatics to therapeutics.
Targeting naturally occurring epitope variants of hepatitis C virus with high-affinity T-cell receptors.
Impact of sequence variation in a dominant HLA-A*02-restricted epitope in hepatitis C virus on priming and cross-reactivity of CD8+ T cells.
Occurrence of antibodies reactive with more than one variant of the putative envelope glycoprotein (gp70) hypervariable region 1 in viremic hepatitis C virus-infected patients.
Inflammatory status in human hepatic cirrhosis.
Structural flexibility of a conserved antigenic region in hepatitis C virus glycoprotein E2 recognized by broadly neutralizing antibodies.
Virion-independent transfer of replication-competent hepatitis C virus RNA between permissive cells.
Cytokines and HCV-related disorders.
An Optimized Hepatitis C Virus E2 Glycoprotein Core Adopts a Functional Homodimer That Efficiently Blocks Virus Entry.
ISGylation of Hepatitis C Virus NS5A Protein Promotes Viral RNA Replication via Recruitment of Cyclophilin A.
Hypervariable region 1 and N-linked glycans of hepatitis C regulate virion neutralization by modulating envelope conformations.
Development of an efficient in vivo cell-based assay system for monitoring hepatitis C virus genotype 4a NS3/4A protease activity.
CD8(+) T-Cell Exhaustion Phenotype in Chronic Hepatitis C Virus Infection Is Associated With Epitope Sequence Variation.
Incorporation of apolipoprotein E into HBV-HCV subviral envelope particles to improve the hepatitis vaccine strategy.
Relation of pretreatment sequence diversity in NS5A region of HCV genotype 1 with immune response between pegylated-INF/ribavirin therapy outcomes.
Lack of variant specific CD8+ T-cell response against mutant and pre-existing variants leads to outgrowth of particular clones in acute hepatitis C.
Toll-like receptors 1 and 6 are involved in TLR2-mediated macrophage activation by hepatitis C virus core and NS3 proteins.
[A HLA-A2-restricted-CTL epitope variation at the N-terminal of HCV helicase and the immune response of CTLs].
Sequence evolution and cross-reactive antibody responses to hypervariable region 1 in acute hepatitis C virus infection.

Incomplete humoral immunity against hepatitis C virus is linked with distinct recognition of putative multiple receptors by E2 envelope glycoprotein.
Heterogeneity and coexistence of oncogenic mechanisms involved in HCV-associated B-cell lymphomas.
Hepatitis C virus quasispecies and pseudotype analysis from acute infection to chronicity in HIV-1 co-infected individuals.
Structural basis for penetration of the glycan shield of hepatitis C virus E2 glycoprotein by a broadly neutralizing human antibody.
The ISG15/USP18 ubiquitin-like pathway (ISGylation system) in hepatitis C virus infection and resistance to interferon therapy.
Role of N-linked glycans in the functions of hepatitis C virus envelope proteins incorporated into infectious virions.
HCV Pit Stop at the Lipid Droplet: Refuel Lipids and Put on a Lipoprotein Coat before Exit.
The hepatitis C virus (HCV) protein, p7, suppresses inflammatory responses to tumor necrosis factor (TNF)- $\alpha$ via signal transducer and activator of transcription (STAT)3 and extracellular signal-regulated kinase (ERK)-mediated induction of suppressor of cytokine signaling (SOCS)3.
Translocation of outer mitochondrial membrane 70 induces interferon response and is impaired by hepatitis C virus NS3.
Immunoglobulin mimicry by Hepatitis C Virus envelope protein E2.
HCV persistence and immune evasion in the absence of memory T cell help.
HCV Glycoprotein Structure and Implications for B-Cell Vaccine Development.
Differential escape of HCV from CD8(+) T cell selection pressure between China and Germany depends on the presenting HLA class I molecule.
Stable cytotoxic T cell escape mutation in hepatitis C virus is linked to maintenance of viral fitness.
Hepatitis C Virus Escape Studies of Human Antibody AR3A Reveal a High Barrier to Resistance and Novel Insights on Viral Antibody Evasion Mechanisms.
Hepatitis C virus present in the sera of infected patients interferes with the autophagic process of monocytes impairing their in-vitro differentiation into dendritic cells.
An interplay between hypervariable region 1 of the hepatitis C virus E2 glycoprotein, the scavenger receptor BI, and high-density lipoprotein promotes both enhancement of infection and protection against neutralizing antibodies.
Systematic identification of anti-interferon function on hepatitis C virus genome reveals p7 as an immune evasion protein.
Toll-like Receptor Response to Human Immunodeficiency Virus Type 1 or Co-Infection with Hepatitis B or C Virus: An Overview.
Transmitted/Founder Viruses Rapidly Escape from CD8+ T Cell Responses in Acute Hepatitis C Virus Infection.
Discovery of danoprevir (ITMN-191/R7227), a highly selective and potent inhibitor of hepatitis C virus (HCV) NS3/4A protease.
Mutational escape of CD8+ T cell epitopes: implications for prevention and therapy of persistent hepatitis virus infections.
Human dendritic cells expressing hepatitis C virus core protein display transcriptional and functional changes consistent with maturation.
Previous failure of interferon-based therapy does not alter the frequency of HCV NS3 protease or NS5B polymerase inhibitor resistance-associated variants: longitudinal analysis in HCV/HIV co-infected patients.



Next-Generation Sequencing Analysis of CpG Methylation of a Tumor Suppressor Gene SHP-1 Promoter in Stable Cell Lines and HCV-Positive Patients.
Affinity maturation of a broadly neutralizing human monoclonal antibody that prevents acute hepatitis C virus infection in mice.
Development of a high-throughput pyrosequencing assay for monitoring temporal evolution and resistance associated variant emergence in the Hepatitis C virus protease coding-region.
Peripheral B cells may serve as a reservoir for persistent hepatitis C virus infection.
Effect of Direct Acting Antivirals (DAAs) on Myeloid-Derived Suppressor Cells Population in Egyptian Chronic Hepatitis C Virus Patients: A Potential Immunomodulatory Role of DAAs.
Elevation of plasma soluble human leukocyte antigen-G in patients with chronic hepatitis C virus infection.
Recruitment of an interferon molecular signaling complex to the mitochondrial membrane: disruption by hepatitis C virus NS3-4A protease.
Toward a vaccine against hepatitis C virus.
Virus-related liver cirrhosis: molecular basis and therapeutic options.
CD4+ immune escape and subsequent T-cell failure following chimpanzee immunization against hepatitis C virus.
Induction of potent cellular immune response in mice by hepatitis C virus NS3 protein with double-stranded RNA.
Tumor necrosis family receptor superfamily member 9/tumor necrosis factor receptor-associated factor 1 pathway on hepatitis C viral persistence and natural history.
Temporal and tight hepatitis C virus gene activation in cultured human hepatoma cells mediated by a cell-permeable Cre recombinase.
Hepatitis G virus/GBV-C persistence: absence of hypervariable E2 region and genetic analysis of viral quasispecies in serum and lymphocytes.
Role of HVR1 sequence similarity in the cross-genotypic neutralization of HCV.
Identification of a residue in hepatitis C virus E2 glycoprotein that determines scavenger receptor BI and CD81 receptor dependency and sensitivity to neutralizing antibodies.
Identification of human progenitors of exhausted CD8(+) T cells associated with elevated IFN- $\gamma$ response in early phase of viral infection.
A novel T cell evasion mechanism in persistent RNA virus infection.
Specific targeting of hepatitis C virus core protein by an intracellular single-chain antibody of human origin.
Hepatitis C virus-specific Th17 cells are suppressed by virus-induced TGF- $\beta$ .
Hepatitis C Recurrence after Orthotopic Liver Transplantation: Mechanisms and Management.
Activation of the Ca(2+)/NFAT Pathway by Assembly of Hepatitis C Virus Core Protein into Nucleocapsid-like Particles.
[N-glycans as contributors in the biological properties of the hepatitis C virus envelope proteins].
In vivo adaptation of hepatitis C virus in chimpanzees for efficient virus production and evasion of apoptosis.
Coordinated evolution of the hepatitis C virus.
Immunopathogenesis of hepatitis C viral infection: Th1/Th2 responses and the role of cytokines.
Eradication of HCV by direct antiviral agents restores mitochondrial function and energy homeostasis in peripheral blood mononuclear cells.
Combined therapy of interferon plus ribavirin promotes multiple adaptive solutions in hepatitis C virus.
Influence of host resistance on viral adaptation: hepatitis C virus as a case study.

Genetic characterization of hypervariable region 1 in patients chronically infected with hepatitis C virus genotype 2.
HVR1-mediated antibody evasion of highly infectious in vivo adapted HCV in humanised mice.
Next-generation sequencing reveals large connected networks of intra-host HCV variants.
Monoclonal Antibodies Directed toward the Hepatitis C Virus Glycoprotein E2 Detect Antigenic Differences Modulated by the N-Terminal Hypervariable Region 1 (HVR1), HVR2, and Intergenotypic Variable Region.
Determining the cellular diversity of hepatitis C virus quasispecies by single-cell viral sequencing.
In vitro adaptation and characterization of attenuated hypervariable region 1 swap chimeras of hepatitis C virus.
[Quasispecies sequence analyses of envelope protein E1/E2 coding genes from four Chinese HCV patients and identification of a novel insertion mutation of HCV].
Genetic variability of hepatitis C virus before and after combined therapy of interferon plus ribavirin.
Modulation of RANTES expression by HCV core protein in liver derived cell lines.
Hepatitis C virus hypervariable region 1 modulates receptor interactions, conceals the CD81 binding site, and protects conserved neutralizing epitopes.
Effect of Glycan Shift on Antibodies against Hepatitis C Virus E2 412-425 Epitope Elicited by Chimeric sHBsAg-Based Virus-Like Particles.
Design of a native-like secreted form of the hepatitis C virus E1E2 heterodimer.
Molecular Determinants of Mouse Adaptation of Rat Hepacivirus.
Enhancement of Immune Responses by Co-delivery of CCL19/MIP-3beta Chemokine Plasmid With HCV Core DNA/Protein Immunization.
Extended interaction networks with HCV protease NS3-4A substrates explain the lack of adaptive capability against protease inhibitors.
Replicative homeostasis: a mechanism of viral persistence.
Synthesis of 1-boraadamantaneamine derivatives with selective astrocyte vs C6 glioma antiproliferative activity. A novel class of anti-hepatitis C agents with potential to bind CD81.
Hypervariable region 1 variant acting as TCR antagonist affects hepatitis C virus-specific CD4+ T cell repertoire by favoring CD95-mediated apoptosis.
Comparison of methods and characterization of small RNAs from plasma extracellular vesicles of HIV/HCV coinfecting patients.
Deep sequencing of hepatitis C virus hypervariable region 1 reveals no correlation between genetic heterogeneity and antiviral treatment outcome.
Complete genome analysis identifies recombinant events and positive selection sites of hepatitis C virus from mainland China during 2010-2019.
The PD-1/PD-L1 Axis and Virus Infections: A Delicate Balance.
Structural and molecular basis of interaction of HCV non-structural protein 5A with human casein kinase 1 $\alpha$ and PKR.
Replicative homeostasis: a fundamental mechanism mediating selective viral replication and escape mutation.
Role of complement component C4 in treatment response and disease progression in chronic hepatitis C patients.
Liver Cancer Emergence Associated with Antiviral Treatment: An Immune Surveillance Failure?
The hepatitis C virus and immune evasion: non-structural 3/4A transgenic mice are resistant to lethal tumour necrosis factor alpha mediated liver disease.
New insights into hepatitis C.
Vaccination to prevent T cell subversion can protect against persistent hepacivirus infection.

The hepatitis C envelope 2 protein inhibits LFA-1-transduced protein kinase C signaling for T-lymphocyte migration.
Hepatitis Viruses Control Host Immune Responses by Modifying the Exosomal Biogenesis Pathway and Cargo.
Long-Term Persistent GBV-B Infection and Development of a Chronic and Progressive Hepatitis C-Like Disease in Marmosets.
Establishment of a novel triple-transgenic mouse: conditionally and liver-specifically expressing hepatitis C virus NS3/4A protease.
Humoral immune response to the E2 protein of hepatitis G virus is associated with long-term recovery from infection and reveals a high frequency of hepatitis G virus exposure among healthy blood donors.
The interplay between N6-methyladenosine and precancerous liver disease: molecular functions and mechanisms.
MHC I-dependent antigen presentation is inhibited by poliovirus protein 3A.
Correlating Flavivirus virulence and levels of intrinsic disorder in shell proteins: protective roles vs. immune evasion.
Epistatic interactions promote persistence of NS3-Q80K in HCV infection by compensating for protein folding instability.
G-Quadruplexes: More Than Just a Kink in Microbial Genomes.
Immunity against the GBV-B hepatitis virus in tamarins can prevent productive infection following rechallenge and is long-lived.
Acute GB virus B infection of marmosets is accompanied by mutations in the NS5A protein.
Generation of the regulatory protein rtTA transgenic mice.
Mutation rate of GB virus C/hepatitis G virus over the entire genome and in subgenomic regions.
Cellular responses during liver fluke infection in sheep and its evasion by the parasite.

Table A2-6, Cluster 5

Cluster 5 focuses on biofilms (especially bacterial), emphasizing their survival by evasion of the immune system (163)
Staphylococcal Biofilm Development: Structure, Regulation, and Treatment Strategies.
A Framework for Understanding the Evasion of Host Immunity by Candida Biofilms.
Staphylococcus aureus biofilms prevent macrophage phagocytosis and attenuate inflammation in vivo.
Image-based 384-well high-throughput screening method for the discovery of skylamycins A to C as biofilm inhibitors and inducers of biofilm detachment in Pseudomonas aeruginosa.
The immune system vs. Pseudomonas aeruginosa biofilms.
Biofilm formation in Streptococcus pneumoniae.
Hijacking of immune defences by biofilms: a multifront strategy.
Current and future approaches to the prevention and treatment of staphylococcal medical device-related infections.
SarZ is a key regulator of biofilm formation and virulence in Staphylococcus epidermidis.
[Neutrophils and bacterial biofilms: dialectics of relationship].
A histone deacetylase complex mediates biofilm dispersal and drug resistance in Candida albicans.
[The research progress of biofilm and biofilm in eye disease].
Advances in Bacterial Biofilm Management for Maintaining Microbiome Homeostasis.

Biofilm formation in the lung contributes to virulence and drug tolerance of <i>Mycobacterium tuberculosis</i> .
<i>Streptococcus pneumoniae</i> biofilm formation and dispersion during colonization and disease.
Deciphering mechanisms of staphylococcal biofilm evasion of host immunity.
Plasma is the main regulator of <i>Staphylococcus epidermidis</i> biofilms virulence genes transcription in human blood.
Biofilm formation avoids complement immunity and phagocytosis of <i>Streptococcus pneumoniae</i> .
Thermonucleases Contribute to <i>Staphylococcus aureus</i> Biofilm Formation in Implant-Associated Infections-A Redundant and Complementary Story.
In vitro efficacy of ciprofloxacin and gentamicin against a biofilm of <i>Pseudomonas aeruginosa</i> and its free-living forms.
Potential Role of Biofilm Formation in the Development of Digestive Tract Cancer With Special Reference to <i>Helicobacter pylori</i> Infection.
The Role of Exopolysaccharides in Oral Biofilms.
<i>Pseudomonas aeruginosa</i> Biofilm Infections: Community Structure, Antimicrobial Tolerance and Immune Response.
Communication is the key: biofilms, quorum sensing, formation and prevention.
Biofilm formation by ST17 and ST19 strains of <i>Streptococcus agalactiae</i> .
Role of Bcr1-activated genes Hwp1 and Hyr1 in <i>Candida albicans</i> oral mucosal biofilms and neutrophil evasion.
Interactions between macrophages and biofilm during <i>Staphylococcus aureus</i> -associated implant infection: difficulties and solutions.
<i>Staphylococcus aureus</i> ATP Synthase Promotes Biofilm Persistence by Influencing Innate Immunity.
Battle royale: Immune response on biofilms - host-pathogen interactions.
The role of extracellular DNA in the establishment, maintenance and perpetuation of bacterial biofilms.
The Extracellular Polysaccharide Matrix of <i>Pseudomonas aeruginosa</i> Biofilms Is a Determinant of Polymorphonuclear Leukocyte Responses.
Biofilm-Leukocyte Cross-Talk: Impact on Immune Polarization and Immunometabolism.
Biofilm Homeostasis Interference Therapy via (1) O(2) -Sensitized Hyperthermia and Immune Microenvironment Re-Rousing for Biofilm-Associated Infections Elimination.
Mycobacterial Biofilms.
Impact of <i>Actinobacillus pleuropneumoniae</i> biofilm mode of growth on the lipid A structures and stimulation of immune cells.
A Formidable Foe Is Sabotaging Your Results: What You Should Know about Biofilms and Wound Healing.
<i>Staphylococcus epidermidis</i> uses distinct mechanisms of biofilm formation to interfere with phagocytosis and activation of mouse macrophage-like cells 774A.1.
GlpC gene is responsible for biofilm formation and defense against phagocytes and imparts tolerance to pH and organic solvents in <i>Proteus vulgaris</i> .
Enhanced Clearing of Wound-Related Pathogenic Bacterial Biofilms Using Protease-Functionalized Antibiotic Nanocarriers.
A global transcriptomic analysis of <i>Staphylococcus aureus</i> biofilm formation across diverse clonal lineages.
How Biofilms Evade Host Defenses.
Biofilms and the Nail Unit.

The immunomodulatory activity of <i>Staphylococcus aureus</i> products derived from biofilm and planktonic cultures.
Targeting macrophage activation for the prevention and treatment of <i>Staphylococcus aureus</i> biofilm infections.
Dispersal and inhibition of biofilms associated with infections.
[Application of bacterial tracking techniques in biofilms].
Staphylococcal Biofilms and Immune Polarization During Prosthetic Joint Infection.
Recent insights into <i>Candida albicans</i> biofilm resistance mechanisms.
Genetic and Biochemical Analysis of CodY-Mediated Cell Aggregation in <i>Staphylococcus aureus</i>
Reveals an Interaction between Extracellular DNA and Polysaccharide in the Extracellular Matrix.
Biofilms and their role on diseases.
<i>Staphylococcus aureus</i> Biofilm: Morphology, Genetics, Pathogenesis and Treatment Strategies.
Proteomic profile of dormancy within <i>Staphylococcus epidermidis</i> biofilms using iTRAQ and label-free strategies.
A crucial role for exopolysaccharide modification in bacterial biofilm formation, immune evasion, and virulence.
Sub-inhibitory tigecycline concentrations induce extracellular matrix binding protein Embp dependent <i>Staphylococcus epidermidis</i> biofilm formation and immune evasion.
2,5-Dimethyl-4-hydroxy-3(2H)-furanone as an Anti-biofilm Agent Against Non- <i>Candida albicans</i> <i>Candida</i> Species.
<i>Staphylococcus aureus</i> biofilms release leukocidins to elicit extracellular trap formation and evade neutrophil-mediated killing.
Immune Responses to <i>Pseudomonas aeruginosa</i> Biofilm Infections.
Antibiotic Tolerance of <i>Staphylococcus aureus</i> Biofilm in Periprosthetic Joint Infections and Antibiofilm Strategies.
<i>Staphylococcus epidermidis</i> biofilms with higher proportions of dormant bacteria induce a lower activation of murine macrophages.
Novel Diagnostic Technologies and Therapeutic Approaches Targeting Chronic Wound Biofilms and Microbiota.
Role of biofilm in children with recurrent upper respiratory tract infections.
The correlation study on antimicrobial resistance and biofilm related genes in clinical isolates of <i>Acinetobacter baumannii</i> .
Protoporphyrin IX derived from dual-species anaerobic biofilms of <i>Fusobacterium necrophorum</i> and <i>Porphyromonas levii</i> attenuates bovine neutrophil function.
DL-2-hydroxyisocaproic acid attenuates inflammatory responses in a murine <i>Candida albicans</i> biofilm model.
Biofilms in wounds: a review of present knowledge.
Busting biofilms: free-living amoebae disrupt preformed methicillin-resistant <i>Staphylococcus aureus</i> (MRSA) and <i>Mycobacterium bovis</i> biofilms.
<i>Candida albicans</i> biofilms: antifungal resistance, immune evasion, and emerging therapeutic strategies.
Proteomic dataset of <i>Candida albicans</i> (ATCC 10231) Biofilm.
Integration of Posttranscriptional Gene Networks into Metabolic Adaptation and Biofilm Maturation in <i>Candida albicans</i> .
Transcriptomic study of <i>Salmonella enterica</i> subspecies <i>enterica</i> serovar Typhi biofilm.
Contributions of the Biofilm Matrix to <i>Candida</i> Pathogenesis.

Exopolysaccharide-Repressing Small Molecules with Antibiofilm and Antivirulence Activity against <i>Pseudomonas aeruginosa</i> .
A Novel Silver Bioactive Glass Elicits Antimicrobial Efficacy Against <i>Pseudomonas aeruginosa</i> and <i>Staphylococcus aureus</i> in an ex Vivo Skin Wound Biofilm Model.
Immune Evasion Mechanisms of <i>Staphylococcus epidermidis</i> Biofilm Infection.
Growth Mode and Physiological State of Cells Prior to Biofilm Formation Affect Immune Evasion and Persistence of <i>Staphylococcus aureus</i> .
From molecules to multispecies ecosystems: the roles of structure in bacterial biofilms.
OmpA-mediated biofilm formation is essential for the commensal bacterium <i>Sodalis glossinidius</i> to colonize the tsetse fly gut.
<i>Candida albicans</i> biofilms do not trigger reactive oxygen species and evade neutrophil killing.
<i>Staphylococcus aureus</i> Biofilm-Conditioned Medium Impairs Macrophage-Mediated Antibiofilm Immune Response by Upregulating KLF2 Expression.
A quantitative proteomic analysis of biofilm adaptation by the periodontal pathogen <i>Tannerella forsythia</i> .
Staphylococcal Vaccine Antigens related to biofilm formation.
Regulation of Biofilm Exopolysaccharide Production by Cyclic Di-Guanosine Monophosphate.
Ultrasound-responsive catalytic microbubbles enhance biofilm elimination and immune activation to treat chronic lung infections.
Role of biofilm formation in <i>Ureaplasma</i> antibiotic susceptibility and development of bronchopulmonary dysplasia in preterm neonates.
Complete Genome Sequences of Two Prolific Biofilm-Forming <i>Staphylococcus aureus</i> Isolates Belonging to USA300 and EMRSA-15 Clonal Lineages.
MyD88-dependent signaling influences fibrosis and alternative macrophage activation during <i>Staphylococcus aureus</i> biofilm infection.
Use of fluorescence foldscope as an effective tool for detection of biofilm formation in <i>Pseudomonas aeruginosa</i> .
Z-form extracellular DNA is a structural component of the bacterial biofilm matrix.
Biofilms of clinical strains of <i>Staphylococcus</i> that do not contain polysaccharide intercellular adhesin.
Proteomic profile of <i>Candida albicans</i> biofilm.
<i>Liberibacter crescens</i> biofilm formation in vitro: establishment of a model system for pathogenic 'Candidatus <i>Liberibacter</i> spp.'.
<i>Staphylococcus epidermidis</i> polysaccharide intercellular adhesin activates complement.
Enzymatic bionanocatalysts for combating peri-implant biofilm infections by specific heat-amplified chemodynamic therapy and innate immunomodulation.
<i>Staphylococcus aureus</i> Evasion of Host Immunity in the Setting of Prosthetic Joint Infection: Biofilm and Beyond.
Comparison of biofilm formation between major clonal lineages of methicillin resistant <i>Staphylococcus aureus</i> .
Clearance of <i>Pseudomonas aeruginosa</i> foreign-body biofilm infections through reduction of the cyclic Di-GMP level in the bacteria.
Therapeutic implications of <i>C. albicans</i> - <i>S. aureus</i> mixed biofilm in a murine subcutaneous catheter model of polymicrobial infection.
Saliva-Derived Commensal and Pathogenic Biofilms in a Human Gingiva Model.
Biofilms on Hospital Shower Hoses: Characterization and Implications for Nosocomial Infections.
Does <i>Streptococcus mitis</i> , a neonatal oropharyngeal bacterium, influence the pathogenicity of <i>Pseudomonas aeruginosa</i> ?

Specific Antibodies to Staphylococcus aureus Biofilm Are Present in Serum from Pigs with Osteomyelitis.
Clearance of mixed biofilms of Streptococcus pneumoniae and methicillin-susceptible/resistant Staphylococcus aureus by antioxidants N-acetyl-L-cysteine and cysteamine.
An Introduction to Bacterial Biofilms and Their Proteases, and Their Roles in Host Infection and Immune Evasion.
Colonization and immune modulation properties of Klebsiella pneumoniae biofilm-dispersed cells.
mazEF Homologue Has a Minor Role in Staphylococcus epidermidis 1457 Virulence Potential.
Targeting and arginine-driven synergizing photodynamic therapy with nutritional immunotherapy nanosystems for combating MRSA biofilms.
Neutrophils From Patients With Invasive Candidiasis Are Inhibited by Candida albicans Biofilms.
Contact-dependent regulation of a Tannerella forsythia virulence factor, BspA, in biofilms.
Lipid and polymer nanoparticles for drug delivery to bacterial biofilms.
Resistance to leukocytes ties benefits of quorum sensing dysfunctionality to biofilm infection.
Biofilms of Pathogenic Nontuberculous Mycobacteria Targeted by New Therapeutic Approaches.
Staphylococcus epidermidis pathogenesis.
Delayed neutrophil recruitment allows nascent Staphylococcus aureus biofilm formation and immune evasion.
Vaginal microbiota dysmicrobism and role of biofilm-forming bacteria.
High levels of genetic recombination during nasopharyngeal carriage and biofilm formation in Streptococcus pneumoniae.
Staphylococcus epidermidis as a cause of bacteremia.
The staphylococcal exopolysaccharide PIA - Biosynthesis and role in biofilm formation, colonization, and infection.
Molecular basis of Staphylococcus epidermidis infections.
Direct Microscopic Observation of Human Neutrophil-Staphylococcus aureus Interaction In Vitro Suggests a Potential Mechanism for Initiation of Biofilm Infection on an Implanted Medical Device.
The extracellular polysaccharide Pel makes the attachment of P. aeruginosa to surfaces symmetric and short-ranged.
[Research progress of c-di-GMP in the regulation of Escherichia coli biofilm].
Micrococcal Nuclease stimulates Staphylococcus aureus Biofilm Formation in a Murine Implant Infection Model.
Reduced phagocytosis and killing of Cryptococcus neoformans biofilm-derived cells by J774.16 macrophages is associated with fungal capsular production and surface modification.
Murine model for the evaluation of candiduria caused by Candida tropicalis from biofilm.
Function analysis of choline binding domains (CBDs) of LytA, LytC and CbpD in biofilm formation of Streptococcus pneumoniae.
Structural and functional insights into the periplasmic detector domain of the GacS histidine kinase controlling biofilm formation in Pseudomonas aeruginosa.
The Complex Mechanism of the Salmonella typhi Biofilm Formation That Facilitates Pathogenicity: A Review.
Pseudomonas aeruginosa chronic colonization in cystic fibrosis patients.
Analysis of Pseudomonas aeruginosa biofilm membrane vesicles supports multiple mechanisms of biogenesis.
Effect of growth conditions on poly-N-acetylglucosamine expression and biofilm formation in Escherichia coli.

The giant staphylococcal protein Embp facilitates colonization of surfaces through Velcro-like attachment to fibrillated fibronectin.
Comment on "Effect of biofilm formation by clinical isolates of <i>Helicobacter pylori</i> on the efflux-mediated resistance to commonly used antibiotics".
[Advances in study on traditional Chinese medicine against biofilms].
Extracellular DNA-induced antimicrobial peptide resistance in <i>Salmonella enterica</i> serovar Typhimurium.
<i>Candida auris</i> Phenotypic Heterogeneity Determines Pathogenicity In Vitro.
Extracellular DNA-induced antimicrobial peptide resistance mechanisms in <i>Pseudomonas aeruginosa</i> .
<i>Staphylococcus epidermidis</i> Contributes to Healthy Maturation of the Nasal Microbiome by Stimulating Antimicrobial Peptide Production.
The Pga59 cell wall protein is an amyloid forming protein involved in adhesion and biofilm establishment in the pathogenic yeast <i>Candida albicans</i> .
Functionalized Chitosan Nanomaterials: A Jammer for Quorum Sensing.
Revealing the Metabolic Alterations during Biofilm Development of <i>Burkholderia cenocepacia</i> Based on Genome-Scale Metabolic Modeling.
<i>Candida albicans</i> and <i>Staphylococcus</i> Species: A Threatening Twosome.
Polysaccharide intercellular adhesin (PIA) protects <i>Staphylococcus epidermidis</i> against major components of the human innate immune system.
Evaluation of biofilm production and characterization of genes encoding type III secretion system among <i>Pseudomonas aeruginosa</i> isolated from burn patients.
Biofilm-derived <i>Legionella pneumophila</i> evades the innate immune response in macrophages.
Catalase Activity is Critical for <i>Proteus mirabilis</i> Biofilm Development, Extracellular Polymeric Substance Composition, and Dissemination during Catheter-Associated Urinary Tract Infection.
The 5'-nucleotidase S5nA is dispensable for evasion of phagocytosis and biofilm formation in <i>Streptococcus pyogenes</i> .
Psl Produced by Mucoid <i>Pseudomonas aeruginosa</i> Contributes to the Establishment of Biofilms and Immune Evasion.
Key role of poly- $\gamma$ -DL-glutamic acid in immune evasion and virulence of <i>Staphylococcus epidermidis</i> .
Implant infections: adhesion, biofilm formation and immune evasion.
Skin-to-blood pH shift triggers metabolome and proteome global remodelling in <i>Staphylococcus epidermidis</i> .
Abrogation of efflux pump activity, biofilm formation, and immune escape by candidacidal geraniol in emerging superbug, <i>Candida auris</i> .
The Human Immune System against <i>Staphylococcus epidermidis</i> .
Chitosan nanoparticles enhance the antibacterial activity of the native polymer against bovine mastitis pathogens.
Phenol-soluble modulins: novel virulence-associated peptides of staphylococci.
Genomic Analysis of Antibiotic-Resistant <i>Staphylococcus epidermidis</i> Isolates From Clinical Sources in the KwaZulu-Natal Province, South Africa.
Comparative proteome profiling of bovine and human <i>Staphylococcus epidermidis</i> strains for screening specifically expressed virulence and adaptation proteins.
The <i>Staphylococcus epidermidis</i> Transcriptional Profile During Carriage.
<i>Staphylococcus epidermidis</i> from prosthetic joint infections induces lower IL-1 $\beta$ release from human neutrophils than isolates from normal flora.
Probiotics for human lactational mastitis.



Mechanisms of resistance to antimicrobial peptides in staphylococci.
Properties and antimicrobial susceptibility of <i>Trueperella pyogenes</i> isolated from bovine mastitis in China.
Multidrug-Resistant and Extensively Drug-Resistant <i>Acinetobacter baumannii</i> Causing Nosocomial Meningitis in the Neurological Intensive Care Unit.
Role of luxS in immune evasion and pathogenicity of piscine <i>Streptococcus agalactiae</i> is not dependent on autoinducer-2.

Table A2-7, Cluster 6

Cluster 6 focuses on the role of fas and fasL in cancer, emphasizing immune evasion by T lymphocyte apoptosis (233)
Irradiation-induced up-regulation of Fas in esophageal squamous cell carcinoma is not accompanied by Fas ligand-mediated apoptosis.
Co-expression of Fas and Fas ligand in malignant glial tumors and cell lines.
Fas (APO-1/CD95) ligand and Fas expression in renal cell carcinomas: correlation with the prognostic factors.
[The significance of Fas/FasL molecular system in breast cancer].
Positive and negative consequences of Fas/Fas ligand interactions in the antitumor response.
Fas ligand is expressed on human squamous cell carcinomas of the head and neck, and it promotes apoptosis of T lymphocytes.
[Fas, fas ligand, immune tolerance, and cancer: implications in cancer of the colon].
Fas and Fas ligand expression in tumor cells and in vascular smooth-muscle cells of colonic and renal carcinomas.
The distribution and intracellular location of Fas and Fas Ligand following gastric carcinogenesis: Fas Ligand expressing gastric carcinoma cells can inhibit local immune response.
Prognostic value of the Fas/Fas ligand system in breast cancer.
[The role of Fas/Fas ligand in tumorigenesis, immune escape, and counterattack in colonic cancer].
Mechanism of counterattack of colorectal cancer cell by Fas/Fas ligand system.
Fas ligand expression in primary colon adenocarcinomas: evidence that the Fas counterattack is a prevalent mechanism of immune evasion in human colon cancer.
The Fas/Fas ligand system and cancer: immune privilege and apoptosis.
Transitional cell carcinoma expresses high levels of Fas ligand in vivo.
Serial in vivo loss and in vitro gain of Fas expression and function in human cancerous pancreatic duct cells.
Fas ligand expression is correlated with metastasis in colorectal carcinoma.
Expression of Fas ligand in retinoblastoma.
Fas-related apoptosis in gastric adenocarcinoma.
Addressing the "Fas counterattack" controversy: blocking fas ligand expression suppresses tumor immune evasion of colon cancer in vivo.
Blockade of the Fas-triggered intracellular signaling pathway in human melanomas is circumvented by cytotoxic lymphocytes.
Resistance to Fas (APO-1/CD95)-mediated apoptosis and expression of Fas ligand in esophageal cancer: the Fas counterattack.
[The evaluation of Fas/Fas ligand system in renal cell carcinoma--the effect of preoperative interferon-alpha therapy].

Expression of Fas and Fas ligand in human gastric adenomas and intestinal-type carcinomas: correlation with proliferation and apoptosis.
Fas/Fas ligand system and apoptosis induction in testicular carcinoma.
Up-regulation of Fas ligand and down-regulation of Fas expression in oral carcinogenesis.
Fas function and tumor progression: use it and lose it.
Tumor-derived Fas ligand induces toxicity in lymphoid organs and plays an important role in successful chemotherapy.
Pre-existing Fas ligand (FasL) in cancer cells elicits tumor-specific protective immunity, but delayed induction of FasL expression after inoculation facilitates tumor formation.
FASL -844C polymorphism is associated with increased activation-induced T cell death and risk of cervical cancer.
Functional polymorphisms in FAS/FASL system increase the risk of neuroblastoma in Chinese population.
Frequent loss of Fas expression and function in human lung tumours with overexpression of FasL in small cell lung carcinoma.
Immunohistochemical analysis of Fas ligand expression in sarcomas. Sarcomas express high level of FasL in vivo.
Expression of Fas ligand in liver metastases of human colonic adenocarcinomas.
Fas and Fas ligand are highly expressed in lymphocytes from cervical intraepithelial neoplasia and cervical cancer patients: A possible role for immune escaping.
Co-expression of Fas (APO-1, CD95)/Fas ligand by BeWo and NJG choriocarcinoma cell lines.
Melanoma cell expression of Fas(Apo-1/CD95) ligand: implications for tumor immune escape.
The effect of Fas/FasL pathway blocking on apoptosis and stemness within breast cancer tumor microenvironment (preclinical study).
The role of fas/fas ligand system in the pathogenesis of liver cirrhosis and hepatocellular carcinoma.
Fas (Apo-1/CD95) and Fas ligand interaction between gastric cancer cells and immune cells.
Functional polymorphisms in cell death pathway genes FAS and FASL contribute to risk of lung cancer.
Significance of Fas and Fas ligand in tuberculous lymphadenitis.
Human lung carcinomas express Fas ligand.
[Mechanism of immune escape in renal cell carcinoma].
Apoptosis and Fas system are significantly involved in the process of liver cirrhosis converting into hepatocellular carcinoma.
Human urinary bladder transitional cell carcinomas acquire the functional Fas ligand during tumor progression.
Fas (CD95/APO-1) and Fas ligand expression in normal pancreas and pancreatic tumors. Implications for immune privilege and immune escape.
Fas ligand is expressed in normal skin and in some cutaneous malignancies.
Not so Fas: Re-evaluating the mechanisms of immune privilege and tumor escape.
Influence of cytokines on the expression of fas ligand and CD44 splice variants in colon carcinoma cells.
Relationship of Fas, FasL, p53 and bcl-2 expression in human non-small cell lung carcinomas.
Expression of Fas ligand by human gastric adenocarcinomas: a potential mechanism of immune escape in stomach cancer.
Autocrine secretion of Fas ligand shields tumor cells from Fas-mediated killing by cytotoxic lymphocytes.
Intracellular Fas ligand in normal and malignant breast epithelium does not induce apoptosis in Fas-sensitive cells.

Aberrant expression of Fas and FasL pro-apoptotic proteins in basal cell and squamous cell carcinomas.
Increased Fas ligand expression by T cells and tumour cells in the progression of actinic keratosis to squamous cell carcinoma.
Fas ligand promotes tumor immune evasion of colon cancer in vivo.
p53 promotes selection for Fas-mediated apoptotic resistance.
Tumor-induced death of immune cells: its mechanisms and consequences.
Constitutive expression of Fas (Apo-1/CD95) ligand on multiple myeloma cells: a potential mechanism of tumor-induced suppression of immune surveillance.
The Fas counterattack: Fas-mediated T cell killing by colon cancer cells expressing Fas ligand.
Tissue microarray analysis of Fas and FasL expressions in human non-small cell lung carcinomas; with reference to the p53 and bcl-2 overexpressions.
Activation-induced cell death: the controversial role of Fas and Fas ligand in immune privilege and tumour counterattack.
T cell dependent and independent antitumor immunity generated by the expression of Fas ligand on mouse lung carcinoma cells.
Frequency of apoptosis of tumor-infiltrating lymphocytes induced by fas counterattack in human colorectal carcinoma and its correlation with prognosis.
Suppression of FasL expression in tumor cells and preventing tumor necrosis factor-induced apoptosis by adenovirus 14.7K is an effective escape mechanism for immune cells.
Fas ligand on tumor cells mediates inactivation of neutrophils.
Involvement of doxorubicin-induced Fas expression in the antitumor effect of doxorubicin on Lewis lung carcinoma in vivo.
The effects of FasL on inflammation and tumor survival are dependent on its expression levels.
Direct in vivo transfection of antisense Fas-ligand reduces tumor growth and invasion.
Human pancreatic cancer cells express non-functional Fas receptors and counterattack lymphocytes by expressing Fas ligand; a potential mechanism for immune escape.
H3K9 Trimethylation Silences Fas Expression To Confer Colon Carcinoma Immune Escape and 5-Fluorouracil Chemoresistance.
A polymorphic -844T/C in FasL promoter predicts survival and relapse in non-small cell lung cancer.
Fas ligand expression in thyroid carcinomas: a potential mechanism of immune evasion.
Posttranslational regulation of Fas ligand function.
Predominant expression of Fas (CD95) ligand in metastatic melanoma revealed by longitudinal analysis.
Suppression of FasL expression in tumor cells and preventing TNF-induced apoptosis was better for immune cells survival.
[The role of Fas/FasL system in the regulation of tumor-immune system interactions in papillary thyroid carcinoma in children and adolescents].
Fas ligand downregulation with antisense oligonucleotides in cells and in cultured tissues of normal skin epidermis and basal cell carcinoma.
Tumor escape mechanism involving Fas and Fas-L molecules in human colorectal tumors.
The Btk-dependent PIP5K1γ lipid kinase activation by Fas counteracts FasL-induced cell death.
Fas ligand expression in esophageal carcinomas and their lymph node metastases.
The Fas/Fas-ligand system: a mechanism for immune evasion in human breast carcinomas.
Relationship between Fas-ligand expression on carcinoma cell and cytotoxic T-lymphocyte response in lymphoepithelioma-like cancer of the stomach.

Expression of Fas and Fas-ligand and analysis of argyrophilic nucleolar organizer regions in squamous cell carcinoma: relationships with tumor stage and grade, and apoptosis.
Decreased Fas expression in advanced-stage bladder cancer is not related to p53 status.
Expression of FasL in squamous cell carcinomas of the cervix and cervical intraepithelial neoplasia and its role in tumor escape mechanism.
Expression of Fas and Fas ligand in uveal melanoma: biological implication and prognostic value.
Bladder cancer cells acquire competent mechanisms to escape Fas-mediated apoptosis and immune surveillance in the course of malignant transformation.
Apoptosis induced in T cells by human neuroblastoma cells: role of Fas ligand.
Induction of Fas ligand expression in a human hepatoblastoma cell line by HCV core protein.
Omega-6 fatty acids can inhibit Fas-mediated apoptosis in a human colorectal carcinoma cell line: a potential mechanism for escape from immune surveillance.
Association between up-regulation of Fas ligand expression and apoptosis of tumor-infiltrating lymphocytes in human breast cancer.
Decreased expression of Fas (CD95/APO1) associated with goblet cell metaplasia in Barrett's esophagus.
Resistance to Fas-mediated cell death in BeWo and NJG choriocarcinoma cell lines: implications in immune privilege.
Activated cytotoxic lymphocytes promote tumor progression by increasing the ability of 3LL tumor cells to mediate MDSC chemoattraction via Fas signaling.
Expression of Fas ligand in gastric carcinoma relates to lymph node metastasis.
Fas ligand is frequently expressed in human pancreatic duct cell carcinoma.
Expression of Fas ligand by microglia: possible role in glioma immune evasion.
[Expression of Kai1 and FasL in non-small cell lung cancer and its clinicopathological significance].
Protein expression of Fas, Fas ligand, Bcl-2 and TGFbeta2 and correlation with survival in initial and recurrent human gliomas.
Overexpression of Fas ligand (FasL) during malignant transformation in the large bowel and in Barrett's metaplasia of the esophagus.
Inverse relation of Fas-ligand and tumor-infiltrating lymphocytes in angiosarcoma: indications of apoptotic tumor counterattack.
Increased expression of fas ligand in human tuberculosis and leprosy lesions: a potential novel mechanism of immune evasion in mycobacterial infection.
FasL expression in colorectal carcinoma and its significance in immune escape of cancer.
Effects of alpha fetoprotein on escape of Bel 7402 cells from attack of lymphocytes.
Up-regulation of fas ligand at early stages and down-regulation of Fas at progressed stages of intrahepatic cholangiocarcinoma reflect evasion from immune surveillance.
Short Communication: Expression of Apoptosis Regulating Factors on T Lymphocytes in Multiple Myeloma Patients.
Influence of interferon gamma on modulation of Fas expression by human colon carcinoma cells and their subsequent sensitivity to antigen-specific CD8+ cytotoxic T lymphocyte attack.
A Critical Role for Fas-Mediated Off-Target Tumor Killing in T-cell Immunotherapy.
Fas ligand expression by neoplastic T lymphocytes mediates elimination of CD8+ cytotoxic T lymphocytes in mycosis fungoides: a potential mechanism of tumor immune escape?
Epigenetic changes in tumor Fas levels determine immune escape and response to therapy.
Human melanoma cells do not express Fas (Apo-1/CD95) ligand.
Pancreatic cancer cells can evade immune surveillance via nonfunctional Fas (APO-1/CD95) receptors and aberrant expression of functional Fas ligand.

[The relationship of Fas and Fas L protein expression in oral carcinogenesis].
Fas-ligand gene silencing in basal cell carcinoma tissue with small interfering RNA.
Expression of fas ligand in human hepatoma cell lines: role of hepatitis-B virus X (HBX) in induction of Fas ligand.
Membrane-bound Fas (Apo-1/CD95) ligand on leukemic cells: A mechanism of tumor immune escape in leukemia patients.
Antigen-specific CD8+ T lymphocytes generated from a DNA vaccine control tumors through the Fas-FasL pathway.
Translocation of Fas by LPA prevents ovarian cancer cells from anti-Fas-induced apoptosis.
Decitabine and vorinostat cooperate to sensitize colon carcinoma cells to Fas ligand-induced apoptosis in vitro and tumor suppression in vivo.
[Apoptosis of active T lymphocytes induced by human laryngocarcinoma cell line].
Fas ligand expression in cervical adenocarcinoma: relevance to lymph node metastasis and tumor progression.
Caspase-1 mediates Fas-induced apoptosis and is up-regulated by interferon-gamma in human astrocytoma cells.
Synergistic induction of the Fas (CD95) ligand promoter by Max and NFkappaB in human non-small lung cancer cells.
Apoptosis of T lymphocytes invading glioblastomas multiforme: a possible tumor defense mechanism.
Quantitative characterization and potential function of membrane Fas/APO-1 (CD95) receptors on leukaemic cells from chronic B and T lymphoid leukaemias.
The biological role of the Fas/FasL system during tumor formation and progression.
[Study on expression and function of fas ligand in human myeloid leukemia cells].
A modification of the JAM test is necessary for a correct determination of apoptosis induced by FasL+ adherent tumor cells.
Influence of antitumor drugs on the expression of Fas system in SW480 colon cancer cells.
[Immune evasion of human lung carcinoma cell A549 suppressed by human lymphoma cell Jurkat via Fas/FasL pathway].
Loss of functional Fas ligand enhances intestinal tumorigenesis in the Min mouse model.
Normal breast epithelial cells induce apoptosis of breast cancer cells via Fas signaling.
The evaluation of soluble Fas and soluble Fas ligand levels of bronchoalveolar lavage fluid in lung cancer patients.
Elevated Fas expression is related to increased apoptosis of circulating CD8+ T cell in patients with gastric cancer.
The Inhibitor of Apoptosis Protein Livin Confers Resistance to Fas-Mediated Immune Cytotoxicity in Refractory Lymphoma.
Fas ligand gene transfer to the vessel wall inhibits neointima formation and overrides the adenovirus-mediated T cell response.
Regulative effect of IFN-gamma on the Fas/Fas L system of cholangiocarcinoma cells.
FasL gene knock-down therapy enhances the antiglioma immune response.
Expression level of c-FLIP versus Fas determines susceptibility to Fas ligand-induced cell death in murine thymoma EL-4 cells.
Alteration of apoptotic regulatory molecules expression during carcinogenesis and tumor progression of renal cell carcinoma.
Action and mechanism of Fas and Fas ligand in immune escape of gallbladder carcinoma.
Cross-talk between tumor cells and neutrophils through the Fas (APO-1, CD95)/FasL system: human glioma cells enhance cell viability and stimulate cytokine production in neutrophils.

Role of Fas ligand (CD95L) in immune escape: the tumor cell strikes back.
Loss of Fas Expression and Function Is Coupled with Colon Cancer Resistance to Immune Checkpoint Inhibitor Immunotherapy.
Correlation of Fas/FasL expression to cell apoptosis in Epstein-Barr virus-associated gastric carcinoma.
Depressing the immune escape of acute myelomonocytic leukemia via an anti-Fas ribozyme.
Soluble Fas and Fas-ligand in bladder cancer in vitro and in vivo.
Immunological escape mechanisms in pancreatic carcinoma.
Expression and distribution of S-100, CD83 and apoptosis-related proteins (Fas, FasL and Bcl-2) in tissues of thyroid carcinoma.
Prevalence of HLA-DRB1 genotype and altered Fas/Fas ligand expression in adrenocortical carcinoma.
CD44 stimulation down-regulates Fas expression and Fas-mediated apoptosis of lung cancer cells.
Corticotropin-releasing hormone (CRH) is expressed in the human cervical carcinoma cells (HeLa) and upregulates the expression of Fas ligand.
[Expression of Fas, FasL and IFN-gamma in gastric cancer].
The human melanoma cell line MeJuSo secretes bioactive FasL and APO2L/TRAIL on the surface of microvesicles. Possible contribution to tumor counterattack.
Ceramide mediates FasL-induced caspase 8 activation in colon carcinoma cells to enhance FasL-induced cytotoxicity by tumor-specific cytotoxic T lymphocytes.
Functional characterization of Fas ligand on tumor cells escaping active specific immunotherapy.
Depletion of the natural killer cell population in the peritoneum by AK-5 tumor cells overexpressing fas-ligand: a mechanism of immune evasion.
Influence of FasL gene expression on hepatic metastasis of colorectal carcinoma.
Fas receptor counterattack against tumor-infiltrating lymphocytes in vivo as a mechanism of immune escape in gastric carcinoma.
Soluble Fas-ligand (sFasL) in human astrocytoma cyst fluid is cytotoxic to T-cells: another potential means of immune evasion.
Role of IFN-gamma produced after intraperitoneal transplantation of AK-5 cells in the induction of Fas ligand expression by tumor cells leading to immune evasion.
Prognostic significance of Fas expression in retinoblastoma.
Pancreatic islets under attack: cellular and molecular effectors.
Yin-Yang-1 decreases Fas-induced apoptosis in acute lymphoblastic leukemia under hypoxic conditions: its implications in immune evasion.
EphB receptors trigger Akt activation and suppress Fas receptor-induced apoptosis in malignant T lymphocytes.
CTL adoptive immunotherapy concurrently mediates tumor regression and tumor escape.
[The influence of FasL gene expression upon hepatic metastasis of colorectal carcinoma].
Expression of Fas and Fas ligand and apoptosis in tumor-associated lymphocytes and in tumor cells from malignant pleural effusions.
Expression of Fas(APO-1/CD95) in tumor-infiltrating and peripheral blood lymphocytes in patients with renal cell carcinoma.
Apoptosis in oncology.
[Apoptosis and expression of Fas/FasL in tumor infiltrating dendritic cells in human endometrioid adenocarcinoma].
The role of MAPK-ERK pathway in 67-kDa laminin receptor-induced FasL expression in human cholangiocarcinoma cells.

Overexpression of thymosin beta-4 renders SW480 colon carcinoma cells more resistant to apoptosis triggered by FasL and two topoisomerase II inhibitors via downregulating Fas and upregulating Survivin expression, respectively.
Fas counterattack in cholangiocarcinoma: a mechanism for immune evasion in human hilar cholangiocarcinomas.
Antiretroviral cytolytic T-lymphocyte nonresponsiveness: FasL/Fas-mediated inhibition of CD4(+) and CD8(+) antiviral T cells by viral antigen-positive veto cells.
[The "Fas counterattack": a mechanism for immune evasion in human hilar cholangiocarcinomas].
Fas ligand expression in lynch syndrome-associated colorectal tumours.
CD8+ tumor-infiltrating lymphocytes are primed for Fas-mediated activation-induced cell death but are not apoptotic in situ.
FASL polymorphism is associated with response to bacillus Calmette-Guérin immunotherapy in bladder cancer.
Crystal Structure of the Complex of Human FasL and Its Decoy Receptor DcR3.
Increased angiogenesis and Fas-ligand expression are independent processes in acute myeloid leukemia.
The role of Fas-mediated apoptosis in thyroid disease.
Defective death receptor signaling as a cause of tumor immune escape.
New naphthoquinones, f13102A and B, from a fungus strain F-13102.
[CD95 resistance and Fas ligand synthesis as mechanism of defense by immunocompetent cells in pancreatic tumors].
Oral cancer cell lines can use multiple ligands, including Fas-L, TRAIL and TNF-alpha, to induce apoptosis in Jurkat T cells: possible mechanisms for immune escape by head and neck cancers.
Expression of RCAS1 and FasL in human trophoblasts and uterine glands during pregnancy: the possible role in immune privilege.
Cyclooxygenase 2 modulates killing of cytotoxic T lymphocytes by colon cancer cells.
Multiple mechanisms of immune evasion can coexist in melanoma tumor cell lines derived from the same patient.
The "Fas counterattack" is not an active mode of tumor immune evasion in colorectal cancer with high-level microsatellite instability.
Evasion of myofibroblasts from immune surveillance: a mechanism for tissue fibrosis.
CD95 (Fas) ligand expression of Epstein-Barr virus (EBV)-infected lymphocytes: a possible mechanism of immune evasion in chronic active EBV infection.
[Study on blocking the tumor immune escape by Fas ligand pathway].
Decoy receptor 3 overexpression and immunologic tolerance in hepatocellular carcinoma (HCC) development.
Mouse interleukin-12/FasTI: A novel bi-functional fusion protein for cancer immuno/gene therapy.
Significant association among the Fas -670 A/G (rs1800682) polymorphism and esophageal cancer, hepatocellular carcinoma, and prostate cancer susceptibility: a meta-analysis.
Cytotoxic T lymphocytes carrying a pattern recognition protein Tag7 can detect evasive, HLA-negative but Hsp70-exposing tumor cells, thereby ensuring FasL/Fas-mediated contact killing.
Granzyme H induces cell death primarily via a Bcl-2-sensitive mitochondrial cell death pathway that does not require direct Bid activation.
Survivin-3B promotes chemoresistance and immune escape by inhibiting caspase-8 and -6 in cancer cells.
Requirement of inositol 1,4,5-trisphosphate receptors for tumor-mediated lymphocyte apoptosis.

Soluble Fas ligand released by colon adenocarcinoma cells induces host lymphocyte apoptosis: an active mode of immune evasion in colon cancer.
Serum levels of soluble molecules associated with evasion of immune surveillance: a study in biliary disease.
Role of cytokines in promoting immune escape of FasL-expressing human colon cancer cells.
Malignant counterpart of myeloid dendritic cell (DC) belonging to acute myelogenous leukemia (AML) exhibits a dichotomous immunoregulatory potential.
Fas (APO-1, CD95) receptor expression and new options for immunotherapy in childhood medulloblastomas.
Interleukin-2 enhances susceptibility of colon cancer cells to FasR mediated apoptosis by up-regulating Fas receptor level and down-regulating FAP-1 expression.
[Blocking the escape of leukemic cells from killing of T cell by combining anti-Fas ribozyme and CD80-IgG fusion protein].
Cytokine associated sensitivity of colon carcinoma cells to FasR-mediated cytotoxicity of CTL.
Immunohistochemical analysis of cytokines and apoptosis in tuberculous lymphadenitis.
Role of Fas ligand expression in promoting escape from immune rejection in a spontaneous tumor model.
Expression of human tumor-associated antigen RCAS1 in Reed-Sternberg cells in association with Epstein-Barr virus infection: a potential mechanism of immune evasion.
ATM kinase regulates tumor immunoreactions in lymphocyte-predominant breast cancer through modulation of NKG2D ligand and TNF cytokines on tumor cells.
Fas (Apo-1, CD95) receptor expression in childhood astrocytomas. Is it a marker of the major apoptotic pathway or a signaling receptor for immune escape of neoplastic cells?
Selective expression of FLIP in malignant melanocytic skin lesions.
Host immunosurveillance controls tumor growth via IFN regulatory factor-8 dependent mechanisms.
Valproic acid cooperates with hydralazine to augment the susceptibility of human osteosarcoma cells to Fas- and NK cell-mediated cell death.
Escaping Death: How Cancer Cells and Infected Cells Resist Cell-Mediated Cytotoxicity.
Opposite roles of metastasin (S100A4) in two potentially tumoricidal mechanisms involving human lymphocyte protein Tag7 and Hsp70.
Endogenous interleukin-18 modulates immune escape of murine melanoma cells by regulating the expression of Fas ligand and reactive oxygen intermediates.
Elevated serum levels of soluble Fas/APO-1 (CD95) in patients with hepatocellular carcinoma.
Resistance of infant leukemia with MLL rearrangement to tumor necrosis factor-related apoptosis-inducing ligand: a possible mechanism for poor sensitivity to antitumor immunity.
Study on blocking the leukemia immune escape after BMT by Fas-Fas ligand pathway.
Immunocytochemical detection of leukocyte-associated and apoptosis-related antigen expression in childhood brain tumors.
Amplification and expression of a decoy receptor for fas ligand (DcR3) in virus (EBV or HTLV-I) associated lymphomas.
Induction of apoptosis by IFNgamma in human neuroblastoma cell lines through the CD95/CD95L autocrine circuit.
Azidothymidine Sensitizes Primary Effusion Lymphoma Cells to Kaposi Sarcoma-Associated Herpesvirus-Specific CD4+ T Cell Control and Inhibits vIRF3 Function.
[Experimental study on blocking immune escape of leukemia cells in the recipient after bone marrow transplantation].



Immunophenotypic characterization of sphere-forming cells derived from the human renal cell carcinoma cell line 786-O.
SPI-Cl and SPI-6 cooperate in the protection from effector cell-mediated cytotoxicity.
The granzyme B inhibitor, PI-9, is differentially expressed during placental development and up-regulated in hydatidiform moles.
Amyloid precursor-like protein 2 suppresses irradiation-induced apoptosis in Ewing sarcoma cells and is elevated in immune-evasive Ewing sarcoma cells.

Table A2-8, Cluster 7

Cluster 7 focuses on immune system evasion by the Epstein-Barr Virus (197)
Immunity and Immune Evasion Mechanisms of Epstein-Barr Virus.
MicroRNAs of Epstein-Barr Virus Attenuate T-Cell-Mediated Immune Control In Vivo.
Epstein-Barr virus immunosuppression of innate immunity mediated by phagocytes.
EBV, the human host, and the 7TM receptors: defense or offense?
Epstein-Barr virus: Biology and clinical disease.
[Regulation and evasion of host immune responses by Epstein-Barr virus].
CD4+ T cell responses in the immune control against latent infection by Epstein-Barr virus.
Immunological aspects of Epstein-Barr virus infection.
An update: Epstein-Barr virus and immune evasion via microRNA regulation.
The interplay between Epstein-Barr virus and B lymphocytes: implications for infection, immunity, and disease.
Immune Evasion by Epstein-Barr Virus.
A spectrum of clinical manifestations caused by host immune responses against Epstein-Barr virus infections.
The oncogenic role of Epstein-Barr virus-encoded microRNAs in Epstein-Barr virus-associated gastric carcinoma.
Differential methylation of Epstein-Barr virus latency promoters facilitates viral persistence in healthy seropositive individuals.
Epstein-Barr virus microRNAs in the pathogenesis of human cancers.
Manipulation of immune responses by Epstein-Barr virus.
Immune escape by Epstein-Barr virus associated malignancies.
Epstein-Barr Virus and Cancer.
Epstein-Barr Virus and the Origin of Myalgic Encephalomyelitis or Chronic Fatigue Syndrome.
Cutaneous lymphoproliferative disorders associated with Epstein-Barr virus infection: a clinical overview.
Inhibition of macrophage inflammatory protein-1 alpha production by Epstein-Barr virus.
Dynamic Epstein-Barr virus gene expression on the path to B-cell transformation.
Interplay among viral antigens, cellular pathways and tumor microenvironment in the pathogenesis of EBV-driven lymphomas.
New strategies and patent therapeutics in EBV-associated diseases.
A Temporal Proteomic Map of Epstein-Barr Virus Lytic Replication in B Cells.
Latency and lytic replication in Epstein-Barr virus-associated oncogenesis.
Innate Immune Recognition of EBV.
Epstein-Barr Virus miR-BART6-3p Inhibits the RIG-I Pathway.

The EBV immunoevasins vIL-10 and BNLF2a protect newly infected B cells from immune recognition and elimination.
The role of Epstein-Barr virus in epithelial malignancies.
The roles of EBV-encoded microRNAs in EBV-associated tumors.
A cluster of virus-encoded microRNAs accelerates acute systemic Epstein-Barr virus infection but does not significantly enhance virus-induced oncogenesis in vivo.
Multiple Viral microRNAs Regulate Interferon Release and Signaling Early during Infection with Epstein-Barr Virus.
Understanding the interplay between host immunity and Epstein-Barr virus in NPC patients.
Dithiocarbamates and viral IL-10 collaborate in the immortalization and evasion of immune response in EBV-infected human B lymphocytes.
The hide-and-seek game of the oncogenic Epstein-Barr virus-encoded EBNA1 protein with the immune system: An RNA G-quadruplex tale.
Epstein-Barr virus in systemic autoimmune diseases.
Immune Escape by Non-coding RNAs of the Epstein Barr Virus.
MicroRNAs of Epstein-Barr Virus Control Innate and Adaptive Antiviral Immunity.
Knockout of Epstein-Barr virus BPLF1 retards B-cell transformation and lymphoma formation in humanized mice.
Identification of multiple sclerosis-related genes regulated by EBV-encoded microRNAs in B cells.
Strategies of Epstein-Barr virus to evade innate antiviral immunity of its human host.
Epstein-Barr virus, rapamycin, and host immune responses.
Molecular Properties and Therapeutic Targeting of the EBV-Encoded Receptor BILF1.
Modes of infection and oncogenesis by the Epstein-Barr virus.
Epstein-Barr viral miRNAs inhibit antiviral CD4+ T cell responses targeting IL-12 and peptide processing.
Epstein-Barr virus, interleukin-10 and multiple sclerosis: A ménage à trois.
Human B cells on their route to latent infection--early but transient expression of lytic genes of Epstein-Barr virus.
Diversification and expansion of the EBV-reactive cytotoxic T lymphocyte repertoire following autologous haematopoietic stem cell transplant for multiple sclerosis.
Epstein barr virus encodes miRNAs to assist host immune escape.
Epigenetics of Epstein Barr virus - A review.
Epstein-barr virus vaccines.
Molecular biology of EBV in relationship to AIDS-associated oncogenesis.
Methylation of the Epstein-Barr virus genome in normal lymphocytes.
EBV-infected B cells in infectious mononucleosis: viral strategies for spreading in the B cell compartment and establishing latency.
[Epstein-Barr virus infection and lymphoproliferative disorders].
Stage-specific inhibition of MHC class I presentation by the Epstein-Barr virus BNLF2a protein during virus lytic cycle.
Transcribed B lymphocyte genes and multiple sclerosis risk genes are underrepresented in Epstein-Barr Virus hypomethylated regions.
Epstein-Barr virus infection alters cellular signal cascades in human nasopharyngeal epithelial cells.
MicroRNA and Other Non-Coding RNAs in Epstein-Barr Virus-Associated Cancers.
Methionine metabolism controls the B cell EBV epigenome and viral latency.

Induction of the Lytic Cycle Sensitizes Epstein-Barr Virus-Infected B Cells to NK Cell Killing That Is Counteracted by Virus-Mediated NK Cell Evasion Mechanisms in the Late Lytic Cycle.
Critical review of Epstein-Barr virus microRNAs relation with EBV-associated gastric cancer.
Epstein-Barr Virus BZLF1-Mediated Downregulation of Proinflammatory Factors Is Essential for Optimal Lytic Viral Replication.
Natural killer T cell strategies to combat Epstein-Barr virus infection.
Epstein-Barr Virus Early Protein BFRF1 Suppresses IFN- $\beta$ Activity by Inhibiting the Activation of IRF3.
Multiple roles of LMP1 in Epstein-Barr virus induced immune escape.
MYC Controls the Epstein-Barr Virus Lytic Switch.
Latent Expression of the Epstein-Barr Virus (EBV)-Encoded Major Histocompatibility Complex Class I TAP Inhibitor, BNLF2a, in EBV-Positive Gastric Carcinomas.
Effects of Epstein-Barr virus on the development of dendritic cells derived from cord blood monocytes: an essential role for apoptosis.
Contribution of viral recombinants to the study of the immune response against the Epstein-Barr virus.
Epstein-Barr Virus: How Its Lytic Phase Contributes to Oncogenesis.
Epstein-Barr virus isolates retain their capacity to evade T cell immunity through BNLF2a despite extensive sequence variation.
BZLF1 Attenuates Transmission of Inflammatory Paracrine Senescence in Epstein-Barr Virus-Infected Cells by Downregulating Tumor Necrosis Factor Alpha.
A yeast-based assay identifies drugs that interfere with immune evasion of the Epstein-Barr virus.
Role of Viral and Host microRNAs in Immune Regulation of Epstein-Barr Virus-Associated Diseases.
Epstein-Barr Virus and Multiple Sclerosis.
Epstein-Barr Virus Envelope Glycoprotein gp110 Inhibits IKKi-Mediated Activation of NF- $\kappa$ B and Promotes the Degradation of $\beta$ -Catenin.
Peptide transporter (TAP-1 and TAP-2)-independent endogenous processing of Epstein-Barr virus (EBV) latent membrane protein 2A: implications for cytotoxic T-lymphocyte control of EBV-associated malignancies.
Epstein Barr virus inhibits the stimulatory effect of TLR7/8 and TLR9 agonists but not CD40 ligand in human B lymphocytes.
Epstein-Barr virus in Hodgkin's disease: more than just an innocent bystander.
Impaired transporter associated with antigen processing-dependent peptide transport during productive EBV infection.
Characterization of Epstein-Barr virus strains and LMP1-deletion variants in Portugal.
Role of BamHI-A Rightward Frame 1 in Epstein-Barr Virus-Associated Epithelial Malignancies.
Modulation of LMP2A expression by a newly identified Epstein-Barr virus-encoded microRNA miR-BART22.
An Epstein-Barr virus encoded inhibitor of Colony Stimulating Factor-1 signaling is an important determinant for acute and persistent EBV infection.
Infection of leukaemic B lymphocytes by Epstein Barr virus.
Epstein-Barr virus induces the differentiation of semi-mature dendritic cells from cord blood monocytes.
Selective induction of Th2-attracting chemokines CCL17 and CCL22 in human B cells by latent membrane protein 1 of Epstein-Barr virus.
RNAs in Epstein-Barr virions control early steps of infection.
Epstein-barr virus blocks the autophagic flux and appropriates the autophagic machinery to enhance viral replication.

Epigenomic landscape study reveals molecular subtypes and EBV-associated regulatory epigenome reprogramming in nasopharyngeal carcinoma.
Epstein-Barr virus large tegument protein BPLF1 contributes to innate immune evasion through interference with toll-like receptor signaling.
The many ways Epstein-Barr virus takes advantage of the RNA tool kit.
NIH conference. Epstein-Barr virus infections: biology, pathogenesis, and management.
Epstein-Barr virus interferes with the amplification of IFN $\alpha$ secretion by activating suppressor of cytokine signaling 3 in primary human monocytes.
Epstein-Barr virus evades restrictive host chromatin closure by subverting B cell activation and germinal center regulatory loci.
EBV dUTPase: A Novel Modulator of Inflammation and the Tumor Microenvironment in EBV-Associated Malignancies.
Isolation and Characterization of Exosomes Released by EBV-Immortalized Cells.
The Missing Link in Epstein-Barr Virus Immune Evasion: the BDLF3 Gene Induces Ubiquitination and Downregulation of Major Histocompatibility Complex Class I (MHC-I) and MHC-II.
T cell therapy of human CMV and EBV infection in immunocompromised hosts.
Silencing the shutoff protein of Epstein-Barr virus in productively infected B cells points to (innate) targets for immune evasion.
Host shutoff during productive Epstein-Barr virus infection is mediated by BGLF5 and may contribute to immune evasion.
Epstein-Barr virus-encoded EBNA1 and ZEBRA: targets for therapeutic strategies against EBV-carrying cancers.
Etiological factors of nasopharyngeal carcinoma.
Epstein-Barr virus LF2: an antagonist to type I interferon.
EBV stimulates TLR- and autophagy-dependent pathways and impairs maturation in plasmacytoid dendritic cells: implications for viral immune escape.
A CD8 $^{+}$ T cell immune evasion protein specific to Epstein-Barr virus and its close relatives in Old World primates.
DNA methylation enzymes and PRC1 restrict B-cell Epstein-Barr virus oncoprotein expression.
Mechanism of activation of the BNLF2a immune evasion gene of Epstein-Barr virus by Zta.
Cooperation between Epstein-Barr virus immune evasion proteins spreads protection from CD8 $^{+}$ T cell recognition across all three phases of the lytic cycle.
EBV and the Pathogenesis of NK/T Cell Lymphoma.
A yeast model for the mechanism of the Epstein-Barr virus immune evasion identifies a new therapeutic target to interfere with the virus stealthiness.
The long-lasting love affair between the budding yeast <i>Saccharomyces cerevisiae</i> and the Epstein-Barr virus.
An HLA-A2-restricted T-cell epitope mapped to the BNLF2a immune evasion protein of Epstein-Barr virus that inhibits TAP.
EBV suppresses prostaglandin E2 biosynthesis in human monocytes.
Constitutive interferon-inducible protein 16-inflammasome activation during Epstein-Barr virus latency I, II, and III in B and epithelial cells.
Persistent infection drives the development of CD8 $^{+}$ T cells specific for late lytic infection antigens in lymphocryptovirus-infected macaques and Epstein-Barr virus-infected humans.
Human cytomegalovirus and Epstein-Barr virus infection impact on (18)F-FDG PET/CT SUVmax, CT volumetric and KRAS-based parameters of patients with locally advanced rectal cancer treated with neoadjuvant therapy.

Integrated copy number and gene expression profiling analysis of Epstein-Barr virus-positive diffuse large B-cell lymphoma.
The ABC of Epstein-Barr virus infections.
EBV latent membrane protein 1 is a negative regulator of TLR9.
The Interaction of Human and Epstein-Barr Virus miRNAs with Multiple Sclerosis Risk Loci.
The innate and T-cell mediated immune response during acute and chronic gammaherpesvirus infection.
Cellular immune responses in transplantation-associated chronic viral infections.
Terminal differentiation into plasma cells initiates the replicative cycle of Epstein-Barr virus in vivo.
The lytic cycle of Epstein-Barr virus is associated with decreased expression of cell surface major histocompatibility complex class I and class II molecules.
The Epstein-Barr virus-encoded LMP-1 oncoprotein negatively affects Tyk2 phosphorylation and interferon signaling in human B cells.
Inhibition of antigen presentation by the glycine/alanine repeat domain is not conserved in simian homologues of Epstein-Barr virus nuclear antigen 1.
BART miRNAs: an unimaginable force in the development of nasopharyngeal carcinoma.
Identification and functional comparison of seven-transmembrane G-protein-coupled BILF1 receptors in recently discovered nonhuman primate lymphocryptoviruses.
Dynamic chromatin environment of key lytic cycle regulatory regions of the Epstein-Barr virus genome.
The Epstein-Barr Virus Glycoprotein gp150 Forms an Immune-Evasive Glycan Shield at the Surface of Infected Cells.
The IL-1 and IL-1 receptor antagonist (IL-1Ra) response of human neutrophils to EBV stimulation. Preponderance of IL-1Ra detection.
BamHI-A rightward frame 1, an Epstein-Barr virus-encoded oncogene and immune modulator.
EBV glycoproteins: where are we now?
Oncogenic Properties of the EBV ZEBRA Protein.
Epstein-Barr virus evades CD4+ T cell responses in lytic cycle through BZLF1-mediated downregulation of CD74 and the cooperation of vBcl-2.
The Epstein-Barr virus G-protein-coupled receptor contributes to immune evasion by targeting MHC class I molecules for degradation.
Epstein-Barr virus encoded interleukin-10 inhibits HLA-class I, ICAM-1, and B7 expression on human monocytes: implications for immune evasion by EBV.
[Factors involved in host-pathogen interaction for the risk of Hodgkin lymphoma induced by Epstein Barr virus].
Herpesvirus vaccines. Development, controversies, and applications.
Epstein-Barr Virus in Multiple Sclerosis: Theory and Emerging Immunotherapies.
EBNA1: Oncogenic Activity, Immune Evasion and Biochemical Functions Provide Targets for Novel Therapeutic Strategies against Epstein-Barr Virus- Associated Cancers.
Nucleolin directly mediates Epstein-Barr virus immune evasion through binding to G-quadruplexes of EBNA1 mRNA.
Type I arginine methyltransferases are intervention points to unveil the oncogenic Epstein-Barr virus to the immune system.
Interference with T cell receptor-HLA-DR interactions by Epstein-Barr virus gp42 results in reduced T helper cell recognition.
Epstein-Barr Virus BART Long Non-coding RNAs Function as Epigenetic Modulators in Nasopharyngeal Carcinoma.

The CD8+ T-cell response to an Epstein-Barr virus-related gammaherpesvirus infecting rhesus macaques provides evidence for immune evasion by the EBNA-1 homologue.
Human gamma-herpesviruses: a review of 2 divergent paths to oncogenesis.
Multistep evolution of B-cell-derived tumors in humans and rodents.
Pathogenesis of chronic active Epstein-Barr virus infection: is this an infectious disease, lymphoproliferative disorder, or immunodeficiency?
Bystander inhibition of humoral immune responses by Epstein-Barr virus LMP1.
Human gammaherpesvirus cytokines and chemokine receptors.
Targeting of EBNA1 for rapid intracellular degradation overrides the inhibitory effects of the Gly-Ala repeat domain and restores CD8+ T cell recognition.
Sequence analysis of EBV immune evasion gene BNLF2a in EBV associated tumors and healthy individuals from nasopharyngeal carcinoma endemic and non-endemic regions of China.
The Epstein-Barr virus alkaline exonuclease BGLF5 serves pleiotropic functions in virus replication.
Human herpesvirus diversity is altered in HLA class I binding peptides.
Alterations in the structure of the EBV nuclear antigen, EBNA1, in epithelial cell tumours.
Epstein-Barr virus-encoded EBNA1 inhibits the canonical NF-kappaB pathway in carcinoma cells by inhibiting IKK phosphorylation.
Novel cationic bis(acylhydrazones) as modulators of Epstein-Barr virus immune evasion acting through disruption of interaction between nucleolin and G-quadruplexes of EBNA1 mRNA.
EBV peptide epitope sensitization restores human cytotoxic T cell recognition of Burkitt's lymphoma cells. Evidence for a critical role for ICAM-2.
Whole-genome profiling of nasopharyngeal carcinoma reveals viral-host co-operation in inflammatory NF-kB activation and immune escape.
Epstein-Barr virus latent gene sequences as geographical markers of viral origin: unique EBNA3 gene signatures identify Japanese viruses as distinct members of the Asian virus family.
Epstein-Barr virus gp42 is posttranslationally modified to produce soluble gp42 that mediates HLA class II immune evasion.
Comprehensive Epstein-Barr Virus Transcriptome by RNA-Sequencing in Angioimmunoblastic T Cell Lymphoma (AITL) and Other Lymphomas.
Roles and regulation of microRNAs in cytomegalovirus infection.
Cloning of the Epstein-Barr virus-related rhesus lymphocryptovirus as a bacterial artificial chromosome: a loss-of-function mutation of the rhBARF1 immune evasion gene.
Structural basis for the constitutive activity and immunomodulatory properties of the Epstein-Barr virus-encoded G protein-coupled receptor BILF1.
Corrigendum: Role of Viral and Host microRNAs in Immune Regulation of Epstein-Barr Virus-Associated Diseases.
EBV BILF1 evolved to downregulate cell surface display of a wide range of HLA class I molecules through their cytoplasmic tail.
EBV protein BNLF2a exploits host tail-anchored protein integration machinery to inhibit TAP.
Repression of CIITA by the Epstein-Barr virus transcription factor Zta is independent of its dimerization and DNA binding.
Viral Infections and Systemic Lupus Erythematosus: New Players in an Old Story.
Epstein-Barr virus-encoded BARF1 protein is a decoy receptor for macrophage colony stimulating factor and interferes with macrophage differentiation and activation.
EBV BART MicroRNAs Target Multiple Pro-apoptotic Cellular Genes to Promote Epithelial Cell Survival.
Conserved Herpesvirus Protein Kinases Target SAMHD1 to Facilitate Virus Replication.

In Cellulo Protein-mRNA Interaction Assay to Determine the Action of G-Quadruplex-Binding Molecules.
The genomic sequence of lymphocryptovirus from cynomolgus macaque.
Viral miRNAs exploiting the endosomal-exosomal pathway for intercellular cross-talk and immune evasion.
G-quadruplexes regulate Epstein-Barr virus-encoded nuclear antigen 1 mRNA translation.
DNAM-1 Activating Receptor and Its Ligands: How Do Viruses Affect the NK Cell-Mediated Immune Surveillance during the Various Phases of Infection?
G-quadruplexes in pathogens: a common route to virulence control?
Genetic diversity: frameshift mechanisms alter coding of a gene (Epstein-Barr virus LF3 gene) that contains multiple 102-base-pair direct sequence repeats.
The nested open reading frame in the Epstein-Barr virus nuclear antigen-1 mRNA encodes a protein capable of inhibiting antigen presentation in cis.
c-MYC impairs immunogenicity of human B cells.
Messenger RNA sequence rather than protein sequence determines the level of self-synthesis and antigen presentation of the EBV-encoded antigen, EBNA1.
Epstein-Barr virus, infectious mononucleosis, and posttransplant lymphoproliferative disorders.
Pathogen interactions with cytokines and host defence: an overview.
Cryo-EM structure of the EBV ribonucleotide reductase BORF2 and mechanism of APOBEC3B inhibition.
The Epstein-Barr Virus Immuno-evasins BCRF1 and BPLF1 Are Expressed by a Mechanism Independent of the Canonical Late Pre-initiation Complex.
The Epstein-Barr Virus BART microRNAs target the pro-apoptotic protein Bim.
Induction of CD137 expression by viral genes reduces T cell costimulation.
A potentially immunologically inert derivative of the reverse tetracycline-controlled transactivator.
Developing a potentially immunologically inert tetracycline-regulatable viral vector for gene therapy in the peripheral nerve.
A homolog of interleukin-10 is encoded by the poxvirus orf virus.
Characterization of an immune-evading doxycycline-inducible lentiviral vector for gene therapy in the spinal cord.

Table A2-9, Cluster 8

Cluster 8 focuses on escape mutants of the hepatitis B virus (560)
Molecular epidemiology and genotyping of hepatitis B virus of HBsAg-positive patients in Oman.
Genetic variation of hepatitis B virus and its significance for pathogenesis.
Symptomatic hepatitis B virus (HBV) reactivation despite reduced viral fitness is associated with HBV test and immune escape mutations in an HIV-coinfected patient.
[Present state and the future direction of HBV vaccine].
Interplay between hepatitis B virus and the innate immune responses: implications for new therapeutic strategies.
[Escape mutants of hepatitis B virus].
Molecular characterization of a variant virus that caused de novo hepatitis B without elevation of hepatitis B surface antigen after chemotherapy with rituximab.
Late HBsAg seroreversion of mutated hepatitis B virus after bone marrow transplantation.
Hepatitis B virus mutation in children.

Occult hepatitis B virus infection with positive hepatitis B e antigen.
Hepatitis B virus genotypes and lamivudine resistance mutations in HIV/hepatitis B virus-coinfected patients.
Selection of hepatitis B virus (HBV) vaccine escape mutants in HBV-infected and HBV/HIV-coinfected patients failing antiretroviral drugs with anti-HBV activity.
A case of hepatitis B reactivation due to the hepatitis B virus escape mutant in a patient undergoing chemotherapy.
Natural history of acute and chronic hepatitis B: The role of HBV genotypes and mutants.
Predominance of HBV Genotype B and HDV Genotype 1 in Vietnamese Patients with Chronic Hepatitis.
Prevalence and molecular characterization of occult hepatitis B virus in pregnant women from Gondar, Ethiopia.
Spontaneous reactivation of hepatitis B virus with S gene mutations in an elderly patient with diabetic nephropathy.
Impact of hepatitis B vaccination on hepatitis B disease and nucleic acid testing in high-prevalence populations.
Crosstalk between innate and adaptive immunity in hepatitis B virus infection.
Virology and clinical sequelae of long-term antiviral therapy in a North American cohort of hepatitis B virus (HBV)/human immunodeficiency virus type 1 (HIV-1) co-infected patients.
Characteristics of escape mutations from occult hepatitis B virus infected patients with hematological malignancies in South Egypt.
Genotype, phylogenetic analysis, and transmission pattern of occult hepatitis B virus (HBV) infection in families of asymptomatic HBsAg carriers.
Innate and adaptive immune responses in chronic hepatitis B virus infections: towards restoration of immune control of viral infection.
Innate and adaptive immune escape mechanisms of hepatitis B virus.
Prevalence of vaccine-induced escape mutants of hepatitis B virus in the adult population in China: a prospective study in 176 restaurant employees.
Molecular detection of hepatitis B virus genotype E with immune escape mutations in chronic hepatitis B patients on long-term antiviral therapy in Jos, Nigeria.
Hepatitis B surface gene variants isolated from blood donors with overt and occult HBV infection in north eastern Egypt.
Hepatitis B virus genetic variants: biological properties and clinical implications.
Hepatitis B caused by a hepatitis B surface antigen escape mutant.
Mutation spectra of the surface-protein-coding region of the HBV genome in HBV-vaccinated and non-vaccinated individuals in Hungary.
A novel hepatitis B virus mutant with A-to-G at nt551 in the surface antigen gene.
[IMMUNOPATHOGENESIS OF OCCULT INFECTION CAUSED BY HEPATITIS B VIRUS].
Occult hepatitis B virus infection and S gene escape mutants in HIV-infected patients after hepatitis B virus vaccination.
A Case of S-Variant Hepatitis B Virus: An Immune System Escape Artist.
Innate immune responses in hepatitis B virus (HBV) infection.
Hepatitis B in HIV-infected patients.
Immune-Escape Hepatitis B Virus Mutations Associated with Viral Reactivation upon Immunosuppression.
High Burden of HBV-Infection and Atypical HBV Strains among HIV-infected Cameroonians.



Stable seroepidemiology of hepatitis B after universal immunization in Taiwan: A 3-year study of national surveillance of primary school students.
Hepatitis B virus infection in post-vaccination South Africa: occult HBV infection and circulating surface gene variants.
Genomic Diversity of Hepatitis B Virus Infection Associated With Fulminant Hepatitis B Development.
Vaccine- and hepatitis B immune globulin-induced escape mutations of hepatitis B virus surface antigen.
Clinical relevance and public health significance of hepatitis B virus genomic variations.
Management and treatment of chronic hepatitis B in HIV-positive patients.
A low rate of hepatitis B virus vaccine breakthrough infections in Mongolia.
Optimization of the algorithm diagnosis chronic hepatitis B markers in patients with newly diagnosed HIV infection.
Serological profile of HBV infection among apparently healthy hospital attendees.
Hepatitis B virus surface antigen impairs myeloid dendritic cell function: a possible immune escape mechanism of hepatitis B virus.
Hepatitis B virus: significance of naturally occurring mutants.
Reactivation of resolved hepatitis B virus infection with immune escape mutations after long-term corticosteroid therapy.
[Clinical significance of hepatitis B virus mutants].
HBV vaccine efficacy and detection and genotyping of vaccinee asymptomatic breakthrough HBV infection in Egypt.
Hepatitis B Virus Adaptation to the CD8+ T Cell Response: Consequences for Host and Pathogen.
Hepatitis B virus-persistent infection and innate immunity defect: Cell-related or virus-related?
Molecular characterization of hepatitis B virus in blood donors in Botswana.
Advances in Immunotherapy for Hepatitis B.
Interferon- $\beta$ response is impaired by hepatitis B virus infection in <i>Tupaia belangeri</i> .
Hepatitis B surface antigen genetic elements critical for immune escape correlate with hepatitis B virus reactivation upon immunosuppression.
Hepatitis B virus vaccine breakthrough infection: surveillance of S gene mutants of HBV.
A Hyper-Glycosylation of HBV Surface Antigen Correlates with HBsAg-Negativity at Immunosuppression-Driven HBV Reactivation in Vivo and Hinders HBsAg Recognition in Vitro.
Profiles of mutations in hepatitis B virus surface and polymerase genes isolated from treatment-naïve Nigerians infected with genotype E.
Detection and molecular characterisation of a diagnosis escape variant associated with occult hepatitis B virus in Brazil.
Hepatitis B virus genome replication triggers toll-like receptor 3-dependent interferon responses in the absence of hepatitis B surface antigen.
Post-Vaccination and Post-Infection Immunity to the Hepatitis B Virus and Circulation of Immune-Escape Variants in the Russian Federation 20 Years after the Start of Mass Vaccination.
3D microfluidic liver cultures as a physiological preclinical tool for hepatitis B virus infection.
Hepatitis B virus drug resistance mutations in HIV/HBV co-infected children in Windhoek, Namibia.
Efficacy of hepatitis B vaccine against antiviral drug-resistant hepatitis B virus mutants in the chimpanzee model.
Unique surface gene variants of hepatitis B virus isolated from patients in the Philippines.
New insight in the pathobiology of hepatitis B virus infection.
Multiple Hepatitis B Virus (HBV) Quasispecies and Immune-Escape Mutations Are Present in HBV Surface Antigen and Reverse Transcriptase of Patients With Acute Hepatitis B.

Detection of Hepatitis B Virus (HBV) Genomes and HBV Drug Resistant Variants by Deep Sequencing Analysis of HBV Genomes in Immune Cell Subsets of HBV Mono-Infected and/or Human Immunodeficiency Virus Type-1 (HIV-1) and HBV Co-Infected Individuals.
Substantial variation in the hepatitis B surface antigen (HBsAg) in hepatitis B virus (HBV)-positive patients from South Africa: Reliable detection of HBV by the Elecsys HBsAg II assay.
Disparate distribution of hepatitis B virus genotypes in four sub-Saharan African countries.
Occult and active hepatitis B virus detection in donated blood in São Paulo, Brazil.
A novel hepatitis B virus variant in the sera of immunized children.
Molecular virology of hepatitis B virus and targets for antiviral intervention.
Molecular epidemiology of chronic hepatitis B virus infection in Greece.
Clinical relevance of hepatitis B virus variants.
Hepatitis B Virus Does Not Interfere With Innate Immune Responses in the Human Liver.
Global Occurrence of Clinically Relevant Hepatitis B Virus Variants as Found by Analysis of Publicly Available Sequencing Data.
Reactivation of Hepatitis B Virus With Immune-Escape Mutations After Ocrelizumab Treatment for Multiple Sclerosis.
Clinical features and viral quasispecies characteristics associated with infection by the hepatitis B virus G145R immune escape mutant.
Hepatitis B virus subgenotype F3 reactivation with vaccine escape mutations: A case report and review of the literature.
Cancer prevention by vaccination against hepatitis B.
Genotyping of acute HBV isolates from England, 1997-2001.
Structure and selection of hepatitis B virus variants during the natural course of infection and interferon therapy.
Overview of hepatitis B viral replication and genetic variability.
Molecular characterization of hepatitis B virus (HBV) isolates, including identification of a novel recombinant, in patients with acute HBV infection attending an Irish hospital.
Characterization of occult hepatitis B virus infection among HIV positive patients in Cameroon.
rt269I Type of Hepatitis B Virus (HBV) Polymerase versus rt269L Is More Prone to Mutations within HBV Genome in Chronic Patients Infected with Genotype C2: Evidence from Analysis of Full HBV Genotype C2 Genome.
Discovery of a Novel Mutation (X8Del) Resulting in an 8-bp Deletion in the Hepatitis B Virus X Gene Associated with Occult Infection in Korean Vaccinated Individuals.
Characterization of Antigen Escape Mutations in Chronic HBV-Infected Patients in Upper Egypt.
A novel orally available small molecule that inhibits hepatitis B virus expression.
Subtypes, genotypes and molecular epidemiology of the hepatitis B virus as reflected by sequence variability of the S-gene.
Hepatitis B virus drug resistance tools: one sequence, two predictions.
Drug-resistant and immune-escape hepatitis B virus mutants, occult hepatitis B infection and coinfections in public hospital patients from Argentina.
[Hepatitis B virus mutants--clinical significance].
Clearance of HBV DNA in immunized children born to HBsAg-positive mothers, years after being diagnosed with occult HBV infection.
Acute hepatitis B due to immune-escape mutations in a naturally immune patient.
Natural history of chronic hepatitis B virus infection from infancy to adult life - the mechanism of inflammation triggering and long-term impacts.

Reduced antigenicity of naturally occurring hepatitis B surface antigen variants with substitutions at the amino acid residue 126.
The glucose metabolic reprogramming in hepatitis B virus infection and hepatitis B virus associated diseases.
New paradigms for treating hepatitis B in HIV/hepatitis B virus co-infected patients.
Establishment of monoclonal antibodies broadly neutralize infection of hepatitis B virus.
Unusual naturally occurring humoral and cellular mutated epitopes of hepatitis B virus in a chronically infected argentine patient with anti-HBs antibodies.
High frequency of Lamivudine and Telbivudine resistance mutations in hepatitis B virus isolates from human immunodeficiency virus co-infected patients on highly active antiretroviral therapy in Bucaramanga, Colombia.
The Association Between Hepatitis B Virus Mutations and the Risk of Liver Disease and Hepatocellular Carcinoma.
The virological and clinical significance of mutations in the overlapping envelope and polymerase genes of hepatitis B virus.
Reactivation of a Vaccine Escape Hepatitis B Virus Mutant in a Cambodian Patient During Anti-Hepatitis C Virus Therapy.
Molecular epidemiology of hepatitis B virus mutants associated with vaccine escape, drug resistance and diagnosis failure.
Characteristics of drug resistant HBV in an international collaborative study of HIV-HBV-infected individuals on extended lamivudine therapy.
Functional analysis of 'a' determinant mutations associated with occult HBV in HIV-positive South Africans.
Variability of the Hepatitis B Surface Protein in HBV-Infected Liver Transplant Recipients.
Appearance of HbeAg in an occult persistent hepatitis B virus infection.
Clinical implications of hepatitis B virus mutations: recent advances.
Prevalence and characterization of lamivudine-resistant hepatitis B virus mutations in HIV-HBV co-infected individuals.
[Hepatitis B virus genetic diversity and mutant].
Hepatitis B virus infection assessed 3 to 18 years after vaccination in Cuban children and adolescents born to HBsAg-positive mothers.
Emergence of HBV resistance to lamivudine (3TC) in HIV/HBV co-infected patients in The Gambia, West Africa.
Serologic and genotypic characterization of hepatitis B virus in HIV-1 infected patients from South West and Littoral Regions of Cameroon.
A description of the hepatitis B virus genomic background in a high-prevalence area in China.
Acute hepatitis B virus infection despite vaccination in a patient treated by infliximab: a case report.
Detection of circulating hepatitis B virus immune escape and polymerase mutants among HBV-positive patients attending Institut Pasteur de Bangui, Central African Republic.
[Analysis of hepatitis B virus (HBV) preS1, preS2 and S gene regions from patient groups infected with HBV genotype D].
Molecular and Genetic Characterization of Hepatitis B Virus (HBV) among Saudi Chronically HBV-Infected Individuals.
Investigation of occult hepatitis B virus infection in anti-hbc positive patients from a liver clinic.
Drug-resistant and immune-escape HBV mutants in HIV-infected hosts.
Molecular and serological characterization of occult hepatitis B among blood donors in Maputo, Mozambique.

Clinical significance of hepatitis B surface antigen mutants.
[Atypical serological profiles in hepatitis B infections: investigation of S gene mutations in cases with concurrently positive for HBsAg and anti-HBs].
Envelope protein variability among HBV-Infected asymptomatic carriers and immunized children with breakthrough infections.
Clinical characteristics and molecular analysis of hepatitis B virus reactivation in hepatitis B surface antigen-negative patients during or after immunosuppressive or cytotoxic chemotherapy.
Hepatitis B escape mutants in Scottish blood donors.
Successful interferon treatment in a patient chronically infected with hepatitis B virus carrying unusual S- (and P-) mutants in the presence of anti-HBs antibodies.
Roles of Hepatitis B Virus Mutations in the Viral Reactivation after Immunosuppression Therapies.
Restoration of replication phenotype of lamivudine-resistant hepatitis B virus mutants by compensatory changes in the "fingers" subdomain of the viral polymerase selected as a consequence of mutations in the overlapping S gene.
Variants within the "a" determinant of HBs gene in children and adolescents with and without hepatitis B vaccination as part of Thailand's Expanded Program on Immunization (EPI).
Patterns of hepatitis b virus immune escape and pol/rt mutations across clinical cohorts of patients with genotypes a, e and occult hepatitis b infection in Nigeria: A multi-centre study.
Failed postnatal immunoprophylaxis for hepatitis B: characteristics of maternal hepatitis B virus as risk factors.
A molecular epidemiological study of the hepatitis B virus in Thailand after 22 years of universal immunization.
How viral genetic variants and genotypes influence disease and treatment outcome of chronic hepatitis B. Time for an individualised approach?
Hepatitis B genotypes and surface antigen mutants present in Pakistani blood donors.
Hepatitis B virus reactivation sustained by a hepatitis B virus surface antigen immune-escape mutant isolate in a patient who was hepatitis B core antibody positive during treatment with sofosbuvir and velpatasvir for hepatitis C virus infection: a case report.
Mutations associated with drug resistance and prevalence of vaccine escape mutations in patients with chronic hepatitis B infection.
Hepatitis B Virus-Associated Hepatocellular Carcinoma and Hepatic Cancer Stem Cells.
Impact of deletions and mutations in Hepatitis B virus envelope proteins on serological profile and clinical evolution.
Mutations in the S gene of hepatitis B virus in three generations of patients with chronic hepatitis B. Hepatitis B virus and HIV infection.
A case of acute hepatitis B in a chronic hepatitis C patient after daclatasvir and asunaprevir combination therapy: hepatitis B virus reactivation or acute self-limited hepatitis?
Hepatitis B Virus Reactivation upon Immunosuppression: Is There a Role for Hepatitis B Core-Related Antigen in Patients with Immune-Escape Mutants? A Case Report.
[Infection with hepatitis B virus enhances basal autophagy].
A quasi-monoclonal anti-HBs response can lead to immune escape of 'wild-type' hepatitis B virus.
Occult HBV infection among anti-HBc positive HIV-infected patients in apex referral centre, Eastern India.
Infection with hepatitis B virus carrying novel pre-S/S gene mutations in female siblings vaccinated at birth: two case reports.
Hepatitis B vaccine: prophylactic, therapeutic, and diagnostic dilemma.

Toll-like receptor (TLR)-mediated innate immune responses in the control of hepatitis B virus (HBV) infection.
Occult hepatitis B virus infection: influence of S protein variants.
Comprehensive Analysis of Clinically Significant Hepatitis B Virus Mutations in Relation to Genotype, Subgenotype and Geographic Region.
High rates of chronic HBV genotype E infection in a group of migrants in Italy from West Africa: Virological characteristics associated with poor immune clearance.
Hepatitis B surface antigen variants in voluntary blood donors in Nanjing, China.
Genetic Diversity of the Hepatitis B Virus Strains in Cuba: Absence of West-African Genotypes despite the Transatlantic Slave Trade.
Molecular and serological evaluation of surface antigen negative hepatitis B virus infection in blood donors from Venezuela.
Epidemiology and Genetic Diversity of Hepatitis B Virus and Hepatitis Delta Virus Infection in Indigenous Communities in Colombia.
Targeting hepatitis B vaccine escape using immunogenetics in Bangladeshi infants.
Genetic variability in coding regions of the surface antigen and reverse transcriptase domain of hepatitis B virus polymerase, Colombia, 2002-2014.
The Variability of Amino Acid Sequences in Hepatitis B Virus.
Evolution of hepatitis B virus sequence from a liver transplant recipient with rapid breakthrough despite hepatitis B immune globulin prophylaxis and lamivudine therapy.
High incidence of lamivudine-resistance-associated vaccine-escape HBV mutants among HIV-coinfected patients on prolonged antiretroviral therapy.
$\alpha$ -Mannosidase I Protein Expression in Peripheral Blood Mononuclear Cells Is Upregulated During Hepatitis B Virus Infection.
Absence of chronicity in infants born to immunized mothers with occult HBV infection in Taiwan.
High frequency of complex mutational patterns in lamivudine resistant hepatitis B virus isolates.
Probable corticosteroid-induced reactivation of latent hepatitis B virus infection in an HIV-positive patient involving immune escape.
A molecular analysis of viral persistence in surface antigen-negative chronic hepatitis B.
Current hepatitis B virus infection situation in Indonesia and its genetic diversity.
Fibrosing cholestatic hepatitis secondary to precore/core promoter hepatitis B variant with lamivudine resistance: successful retransplantation with combination adefovir dipivoxil and hepatitis B immunoglobulin.
[Clinical evaluation of a novel HBsAg quantitative assay].
[Hepatitis B Virus (Hepadnaviridae: Orthohepadnavirus: Hepatitis B virus) among Hospitalized Mentally Disabled Patients is not transmitted by their nurses or family members].
Advances in treating drug-resistant hepatitis B virus in HIV-infected patients.
Genomic Variability of Hepatitis B Virus Circulating in Brazilian Western Amazon.
[Identification of vertical transmission of hepatitis B virus from mother to children by direct sequencing a segment of surface gene of hepatitis B virus].
Understanding Amino Acid Mutations in Hepatitis B Virus Proteins for Rational Design of Vaccines and Drugs.
Prevalence and molecular analysis of occult hepatitis B virus infection isolated in a sample of cryptogenic cirrhosis patients in Iran.
Impact of hepatitis B virus surface protein mutations on the diagnosis of occult hepatitis B virus infection.

Renal transplantation from hepatitis B surface antigen (HBsAg)-positive donors to HBsAg-negative recipients: a case of post-transplant fulminant hepatitis associated with an extensively mutated hepatitis B virus strain and review of the current literature.
The variability of hepatitis B envelope is associated with HBs antigen persistence in either chronic or acute HBV genotype A infection.
When does hepatitis B virus meet long-stranded noncoding RNAs?
A Novel Insertion in the Hepatitis B Virus Surface Protein Leading to Hyperglycosylation Causes Diagnostic and Immune Escape.
Endocytosis of hepatitis B immune globulin into hepatocytes inhibits the secretion of hepatitis B virus surface antigen and virions.
A multicentre molecular analysis of hepatitis B and blood-borne virus coinfections in Viet Nam. [HBV vaccine escape mutations in a chronic hepatitis B patient treated with nucleos(t)ide analogues].
Impact of HBV genotypes A and D genetic variability on infection evolution.
Altered antigenicity of 'a' determinant variants of hepatitis B virus.
Transmission of "a" determinant variants of hepatitis B virus in immunized babies born to HBsAg carrier mothers.
A new hepatitis B virus vaccine escape mutation in a renal transplant recipient.
Occult hepatitis B virus infection in hepatitis B vaccinated children in Taiwan.
The effectiveness of hepatitis B vaccine in toddlers based on the five-year period national basic health research (Riskesdas 2007, 2013 and 2018) in Indonesia.
Immune escape by hepatitis B viruses.
Naturally occurring hepatitis B virus reverse transcriptase mutations related to potential antiviral drug resistance and liver disease progression.
The high prevalence of occult hepatitis B infections among the partners of chronically infected HBV blood donors emphasizes the potential residual risk to blood safety.
Circulating and Hepatic BDCA1+, BDCA2+, and BDCA3+ Dendritic Cells Are Differentially Subverted in Patients With Chronic HBV Infection.
Alpha-enolase regulates hepatitis B virus replication through suppression of the interferon signalling pathway.
Sequence analysis of the HBV S protein in Chinese patients with coexisting HBsAg and anti-HBs antibodies.
HIV/hepatitis B virus co-infection: current challenges and new strategies.
Associations between HLA class I alleles and escape mutations in the hepatitis B virus core gene in New Zealand-resident Tongans.
Evaluation of the Abbott HBV RUO sequencing assay combined with laboratory-modified interpretive software.
Hepatitis B Virus Inhibits Neutrophil Extracellular Trap Release by Modulating Reactive Oxygen Species Production and Autophagy.
A new unconventional HLA-A2-restricted epitope from HBV core protein elicits antiviral cytotoxic T lymphocytes.
Whole genome analysis of hepatitis B virus before and during long-term therapy in chronic infected patients: Molecular characterization, impact on treatment and liver disease progression.
HIV therapy with unknown HBV status is responsible for higher rate of HBV genome variability in Ethiopia.
Comparative biology and pathogenesis of AIDS and hepatitis B viruses: related but different.
Prevalence of HBsAg mutants and impact of hepatitis B infant immunisation in four Pacific Island countries.

Animal Models for the Study of Hepatitis B Virus Pathobiology and Immunity: Past, Present, and Future.
Hepatitis B viremia in completely immunized individuals negative for anti-hepatitis B core antibody.
Validation of cross-genotype neutralization by hepatitis B virus-specific monoclonal antibodies by in vitro and in vivo infection.
Low detection rate and maternal provenance of hepatitis B virus S gene mutants in cases of failed postnatal immunoprophylaxis in England and Wales.
Phylogenetic and drug- and vaccine-resistance profiles of Hepatitis B Virus among children with HIV co-infection in Pakistan.
[The prevalence clinically significant virus mutations among patients with chronic viral hepatitis B.].
Epidemiological, virological and clinical characteristics of HBV infection in 223 HIV co-infected patients: a French multi-centre collaborative study.
High Frequency of Antiviral Resistance Mutations in HBV Genotypes A2 and H: Multidrug Resistance Strains in Mexico.
Phenotyping hepatitis B virus variants: from transfection towards a small animal in vivo infection model.
Hepatitis B virus (HBV) variants fluctuate in paired plasma and peripheral blood mononuclear cells among patient cohorts during different chronic hepatitis B (CHB) disease phases.
[Study of the quasispecies dynamics of serum hepatitis B virus in a patient with acute exacerbations of chronic hepatitis B].
Hepatitis B virus rigs the cellular metabolome to avoid innate immune recognition.
Low prevalence of hepatitis B vaccine escape mutants among individuals born after the initiation of a nationwide vaccination program in Iran.
HBV/D1: a major HBV subgenotype circulating in Uyghur patients with chronic HBV infection in Xinjiang, China.
Molecular evolution of hepatitis B vaccine escape variants in China, during 2000-2016.
Deacetylation of Notch1 by SIRT1 contributes to HBsAg- and HBeAg-mediated M2 macrophage polarization.
Induction of humoral and cell-mediated immune responses by hepatitis B virus epitope displayed on the virus-like particles of prawn nodavirus.
HIV-HBV vaccine escape mutant infection with loss of HBV surface antibody and persistent HBV viremia on tenofovir/emtricitabine without antiviral resistance.
Molecular analysis of an HBsAg-negative hepatitis B virus mutant selected in a tenofovir-treated HIV-hepatitis B virus co-infected patient.
Recent and occult hepatitis B virus infections among blood donors in the United States.
A new immune escape mutant of hepatitis B virus with an Asp to Ala substitution in aa144 of the envelope major protein.
Replication efficiency and sequence analysis of full-length hepatitis B virus isolates from hepatocellular carcinoma tissues.
Hepatitis B virus polymerase disrupts K63-linked ubiquitination of STING to block innate cytosolic DNA-sensing pathways.
Adaptation of the hepatitis B virus core protein to CD8(+) T-cell selection pressure.
Intra-host diversity of hepatitis B virus during mother-to-child transmission: the X gene may play a key role in virus survival in children after transmission.
Stable Human Hepatoma Cell Lines for Efficient Regulated Expression of Nucleoside/Nucleotide Analog Resistant and Vaccine Escape Hepatitis B Virus Variants and Woolly Monkey Hepatitis B Virus.

In Silico Prediction of Human Leukocytes Antigen (HLA) Class II Binding Hepatitis B Virus (HBV) Peptides in Botswana.
Association of Pre-S/S and Polymerase Mutations with Acute and Chronic Hepatitis B Virus Infections in Patients from Rio de Janeiro, Brazil.
Detection of lamivudine-resistant variants and mutations related to reduced antigenicity of HBsAg in individuals from the cities of Santos and São Paulo, Brazil.
Hepatitis B in HIV patients: what is the current treatment and what are the challenges?
Effect of HIV co-infection on mutation patterns of HBV in patients with lamivudine-resistant chronic hepatitis B.
Potential threat of drug-resistant and vaccine-escape HBV mutants to public health.
A systematic review of hepatitis B virus (HBV) drug and vaccine escape mutations in Africa: A call for urgent action.
Genetic characterization of hepatitis B virus in peripheral blood leukocytes: evidence for selection and compartmentalization of viral variants with the immune escape G145R mutation.
Hepatitis B eradication: vaccine as a key player.
Diversity of hepatitis B virus infecting Malaysian candidate blood donors is driven by viral and host factors.
Phage display creates innovative applications to combat hepatitis B virus.
Reduced antigenicity of the hepatitis B virus HBsAg protein arising as a consequence of sequence changes in the overlapping polymerase gene that are selected by lamivudine therapy.
Molecular epidemiology of hepatitis B and C virus infections in Asia.
Efficient cellular and humoral immune response and production of virus-neutralizing antibodies by the Hepatitis B Virus S/preS1(16-42) antigen.
Massive APOBEC3 editing of hepatitis B viral DNA in cirrhosis.
Genotyping and Mutation Pattern in the Overlapping MHR Region of HBV Isolates in Southern Khorasan, Eastern Iran.
Compartmental HBV evolution and replication in liver and extrahepatic sites after nucleos(tide) analogue therapy in chronic hepatitis B carriers.
Reactivation of resolved infection with the hepatitis B virus immune escape mutant G145R during dasatinib treatment for chronic myeloid leukemia.
Overlapping structure of hepatitis B virus (HBV) genome and immune selection pressure are critical forces modulating HBV evolution.
Circulating immune complexes and mutations of HBsAg are associated with the undetectable HBsAg in anti-HBs and HBeAg positive occult hepatitis B virus infection.
Hepatitis B Virus Blocks the CRE/CREB Complex and Prevents TLR9 Transcription and Function in Human B Cells.
Hepatitis B Virus (HBV) Variants in Untreated and Tenofovir Treated Chronic Hepatitis B (CHB) Patients during Pregnancy and Post-Partum Follow-Up.
Late hepatitis B virus reactivation after lamivudine prophylaxis interruption in an anti-HBs-positive and anti-HBc-negative patient treated with rituximab-containing therapy.
Identification of a new hepatitis B virus recombinant D2/D3 in the city of São Paulo, Brazil.
Comprehensive analysis of the prevalence of hepatitis B virus escape mutations in the major hydrophilic region of surface antigen.
Prevalence of S gene mutations within the major hydrophilic region of hepatitis B virus in patients in Dongguan, southern China.
Hepatitis B virus genome variability and disease progression: the impact of pre-core mutants and HBV genotypes.



Hepatitis B virus genotypes and resistance mutations in patients under long term lamivudine therapy: characterization of genotype G in Brazil.
Hepatitis B vaccination.
A family cluster of an immune escape variant of hepatitis B virus infecting a mother and her two fully immunized children.
Hepatitis B virus hijacks CTHRC1 to evade host immunity and maintain replication.
Mechanisms of HBV immune evasion.
[Functional analysis of hepatitis B virus immune escape mutants with insertion mutations in the surface antigen].
Variability in hepatitis B virus DNA: phylogenetic, epidemiological and clinical implications.
Alternative splicing of hepatitis B virus: A novel virus/host interaction altering liver immunity.
Detection of hepatitis B virus isolates with mutations associated with immune escape mutants among pregnant women in Ibadan, southwestern Nigeria.
Global prevalence and phylogeny of hepatitis B virus (HBV) drug and vaccine resistance mutations.
Role of Dendritic Cell-Specific ICAM-3-Grabbing Nonintegrin on Dendritic Cells in the Recognition of Hepatitis B Virus.
[Detection of mutants of the "a" determinant region of hepatitis B surface antigen S gene among Wuhan childhood patients].
The role of innate immunity in the immunopathology and treatment of HBV infection.
Biological significance of amino acid substitutions in hepatitis B surface antigen (HBsAg) for glycosylation, secretion, antigenicity and immunogenicity of HBsAg and hepatitis B virus replication.
Establishment of a Seronegative Occult Infection With an Active Hepatitis B Virus Reservoir Enriched of Vaccine Escape Mutations in a Vaccinated Infant After Liver Transplantation.
A recombinant human immunoglobulin with coherent avidity to hepatitis B virus surface antigens of various viral genotypes and clinical mutants.
A Combination of Human Broadly Neutralizing Antibodies against Hepatitis B Virus HBsAg with Distinct Epitopes Suppresses Escape Mutations.
Evolution of hepatitis B virus during primary infection in humans: transient generation of cytotoxic T-cell mutants.
Acute hepatitis B caused by a vaccine-escape HBV strain in vaccinated subject: sequence analysis and therapeutic strategy.
The molecular virology of hepatitis B virus.
Hepatitis B immune globulin and HBV-related liver transplantation.
The effect of introduction of universal childhood hepatitis B immunization in South Africa on the prevalence of serologically negative hepatitis B virus infection and the selection of immune escape variants.
Molecular Epidemiology of Hepatitis B Virus in Turkish Cypriot.
Molecular evolution of hepatitis B virus over 25 years.
Molecular characterization of hepatitis B virus reveals circulation of multiple subgenotypes of genotype D with clinically important mutations in central India.
Hepatitis B virus DNA in serum of healthy black African adults positive for hepatitis B surface antibody alone: possible association with recombination between genotypes A and D.
Occult hepatitis B virus and surface antigen mutant infection in healthy vaccinated cohorts and children with various forms of hepatitis and multiple transfusions.
Naturally occurring polymerase and surface gene variants of hepatitis B virus in Turkish hemodialysis patients with chronic hepatitis B.

Mutational analysis of HBs Ag-positive mothers and their infected children despite immunoprophylaxis.
Immune Complex Mediated Glomerulonephritis with Acute Thrombotic Microangiopathy following Newly Detected Hepatitis B Virus Infection in a Kidney Transplant Recipient.
Molecular characterization of hepatitis B virus in liver disease patients and asymptomatic carriers of the virus in Sudan.
Lack of Association Between Transforming Growth Factor Beta 1 -509C/T and +915G/C Polymorphisms and Chronic Hepatitis B in Iranian Patients.
Molecular epidemiology of hepatitis B in the Indigenous people of northern Australia.
Genetic variation of hepatitis B surface antigen among acute and chronic hepatitis B virus infections in The Netherlands.
Role of hepatitis B virus genetic barrier in drug-resistance and immune-escape development.
Comparative analysis of hepatitis B virus infections in blood donors born before and after the implementation of universal HBV vaccination in southern China.
Impact of Universal Hepatitis B Vaccination on Prevalence, Infection-Associated Morbidity and Mortality, and Circulation of Immune Escape Variants in Russia.
[HBs antigen mutants: prevalence, clinical and diagnostic implications].
Genetic variations of hepatitis B virus.
Evolution and mutations of hepatitis B virus quasispecies in genotype B and C during vertical transmission.
Does rapid oligomerization of hepatitis B envelope proteins play a role in resistance to proteasome degradation and enhance chronicity?
Clade homogeneity and low rate of delta virus despite hyperendemicity of hepatitis B virus in Ethiopia.
Down-regulation of suppressor of cytokine signaling 3 by miR-122 enhances interferon-mediated suppression of hepatitis B virus.
Decreased antigenicity profiles of immune-escaped and drug-resistant hepatitis B surface antigen (HBsAg) double mutants.
Coordinated evolution of the hepatitis B virus polymerase.
Novel concepts on mechanisms underlying Hepatitis Delta virus persistence and related pathogenesis.
Prevalence of occult hepatitis B virus infection and Torque teno virus infection and their association with hepatocellular carcinoma in chronic hepatitis C patients.
A computational approach to identify point mutations associated with occult hepatitis B: significant mutations affect coding regions but not regulative elements of HBV.
Hepatitis B virus polymerase suppresses NF- $\kappa$ B signaling by inhibiting the activity of IKKs via interaction with Hsp90 $\beta$ .
Clade analysis and surface antigen polymorphism of hepatitis B virus American genotypes.
Human hepatitis B virus surface and e antigens inhibit major vault protein signaling in interferon induction pathways.
Hepatitis B Surface Antigen Suppresses the Activation of Nuclear Factor Kappa B Pathway via Interaction With the TAK1-TAB2 Complex.
The sK122R mutation of hepatitis B virus (HBV) is associated with occult HBV infection: Analysis of a large cohort of Chinese patients.
Increased intrahepatic quasispecies heterogeneity correlates with off-treatment sustained response to nucleos(t)ide analogues in e antigen-positive chronic hepatitis B patients.
[Effects of hepatitis B virus X protein on chronic hepatitis B pathophysiology].

Modification of the hepatitis B virus envelope protein glycosylation pattern interferes with secretion of viral particles, infectivity, and susceptibility to neutralizing antibodies.
Antigenic switching of hepatitis B virus by alternative dimerization of the capsid protein.
Naturally occurring core immune-escape and carboxy-terminal mutations\truncations in patients with e antigen negative chronic hepatitis B.
Phylogenetic analysis of new hepatitis B virus isolates from Nigeria supports endemicity of genotype E in West Africa.
Hepatitis viruses: genetic variants and clinical significance.
Cross-Protection of Hepatitis B Vaccination among Different Genotypes.
Coexistence of hepatitis B surface antigen (HBs Ag) and anti-HBs antibodies in chronic hepatitis B virus carriers: influence of "a" determinant variants.
HBV inhibits LPS-induced NLRP3 inflammasome activation and IL-1 $\beta$ production via suppressing the NF- $\kappa$ B pathway and ROS production.
Impairment of hepatitis B virus virion secretion by single-amino-acid substitutions in the small envelope protein and rescue by a novel glycosylation site.
The Computational Analysis of Single Nucleotide Associated with MicroRNA Affecting Hepatitis B Infection.
Plasma soluble human leukocyte antigen-G expression is a potential clinical biomarker in patients with hepatitis B virus infection.
Maternal A90V mutation in the PreS1 gene of sub-genotype C2 hepatitis B virus is associated with intrauterine transmission.
Methyltransferase PRMT1 is a binding partner of HBx and a negative regulator of hepatitis B virus transcription.
Association Between HBx Variations and Development of Severe Liver Disease Among Indonesian Patients.
A new hepatitis B virus e antigen-negative strain gene used as a reference sequence in an animal model.
Expansion of viral variants associated with immune escape and impaired virion secretion in patients with HBV reactivation after resolved infection.
Hot-spot mutations in hepatitis B virus core gene: eliciting or evading immune clearance?
Investigation of immune escape-associated mutations of hepatitis B virus in patients harboring hepatitis B virus drug-resistance mutations.
Chemical-induced degradation of PreS2 mutant surface antigen via the induction of microautophagy.
Preparation, identification, and clinical application of anti-HBs monoclonal antibody that binds both wild-type and immune escape mutant HBsAgs.
Amino acid similarities and divergences in the small surface proteins of genotype C hepatitis B viruses between nucleos(t)ide analogue-naïve and lamivudine-treated patients with chronic hepatitis B.
Peripartum cytokine flares in a multiethnic cohort of chronic hepatitis B carriers does not correlate with hepatitis B virus suppression or increased risk of liver disease.
Hepatitis B virus genotype D strains from Estonia share sequence similarity with strains from Siberia and may specify ayw4.
Impaired antigen processing and presentation machinery is associated with immunotolerant state in chronic hepatitis B virus infection.
Characterization and assessment of HBV chronically infected patients: Identification of those eligible for treatment in the South West region of Cameroon.
High Frequency Occult Hepatitis B Virus Infection Detected in Non-Resolved Donations Suggests the Requirement of Anti-HBc Test in Blood Donors in Southern China.

Hepatitis B virus X protein impairs $\alpha$ -interferon signaling via up-regulation of suppressor of cytokine signaling 3 and protein phosphatase 2A.
Are International Units of Anti-HBs Antibodies Always Indicative of Hepatitis B Virus Neutralizing Activity?
HBV-Induced Increased N6 Methyladenosine Modification of PTEN RNA Affects Innate Immunity and Contributes to HCC.
Immune response induced by immunization with Hepatitis B virus core DNA isolated from chronic active hepatitis patients.
Hepatitis B virus-triggered autophagy targets TNFRSF10B/death receptor 5 for degradation to limit TNFSF10/TRAIL response.
Antiviral drug resistance and hepatitis B: a continuing public health problem.
[Correlation of hepatocyte expression of hepatitis B viral core antigen and the clinicopathological characters in chronic hepatitis B patients].
Cloning and growth-regulated expression of the gene encoding the hepatitis B virus middle surface antigen in <i>Yarrowia lipolytica</i> .
[Impact of amino acid sequence variation of a determinant(s) on the antigenic properties of hepatitis B virus HBsAg].
[Preparation of anti HBs monoclonal antibody and its combination characterization to both wild-type HBsAg and immune escape mutant HBsAg].
Hepatitis B virus X protein induced expression of the Nur77 gene.
Best practice in the treatment of chronic hepatitis B: a summary of the European Viral Hepatitis Educational Initiative (EVHEI).
A hepatitis B virus-derived human hepatic cell-specific heparin-binding peptide: identification and application to a drug delivery system.
Deficiency in virion secretion and decreased stability of the hepatitis B virus immune escape mutant G145R.
Role of RNA secondary structure in emergence of compartment specific hepatitis B virus immune escape variants.
Short duration of lamivudine for the prevention of hepatitis B virus transmission in pregnancy: lack of potency and selection of resistance mutations.
Hypermodification and immune escape of an internally deleted middle-envelope (M) protein of frequent and predominant hepatitis B virus variants.
Suppression of mRNA accumulation by the duck hepatitis B virus reverse transcriptase.
Molecular epidemiology and clinical characteristics of hepatitis B identified through the French mandatory notification system.
HBcAg induces interleukin-10 production, inhibiting HBcAg-specific Th17 responses in chronic hepatitis B patients.
Impaired virion secretion by hepatitis B virus immune escape mutants and its rescue by wild-type envelope proteins or a second-site mutation.
Hepatitis B virus core protein mutations are concentrated in B cell epitopes in progressive disease and in T helper cell epitopes during clinical remission.
Development of a virus-mimicking nanocarrier for drug delivery systems: The bio-nanocapsule.
Viral apoptotic mimicry: an immune evasion strategy developed by the hepatitis B virus?
Aberrant Splicing Events and Epigenetics in Viral Oncogenomics: Current Therapeutic Strategies.
Diverse immune responses to HBV surface epitope variants after vaccine booster in adolescents immunized in infancy.
Nucleic Acid Polymers Are Active against Hepatitis Delta Virus Infection In Vitro.

A 'first loop' linear epitope accessible on native hepatitis B surface antigen that persists in the face of 'second loop' immune escape.
Deep sequencing of hepatitis B surface antigen gene in the preserved umbilical cords in immunoprophylaxis failure against mother-to-child HBV transmission.
Impact of immune escape mutations and N-linked glycosylation on the secretion of hepatitis B virus virions and subviral particles: Role of the small envelope protein.
Hepatitis B virus PreS2-mutant large surface antigen activates store-operated calcium entry and promotes chromosome instability.
Surveillance of the genetic variation in incident HIV, HCV, and HBV infections in blood and plasma donors: implications for blood safety, diagnostics, treatment, and molecular epidemiology.
Effect of Chinese medicine therapy for strengthening-Pi and nourishing-Shen in preventing lamivudine induced YMDD mutation and its immunologic mechanism.
Impact of HBeAg on the maturation and function of dendritic cells.
Evolution and phenotypic characterization of whole HBV genome in compliant patients experiencing unexplained entecavir treatment failure.
Mechanisms behind TB, HBV, and HIV chronic infections.
The -3279C>A and -924A>G polymorphisms in the FOXP3 Gene Are Associated With Viral Load and Liver Enzyme Levels in Patients With Chronic Viral Liver Diseases.
Genotype-specific acquisition, evolution and adaptation of characteristic mutations in hepatitis E virus.
Modulation of the immunogenicity of virus-like particles composed of mutant hepatitis B virus envelope subunits.
Molecular mechanisms underlying the development of hepatocellular carcinoma.
Molecular epidemiology and genetic diversity of hepatitis B virus in Mar del Plata city, Argentina.
[Hepatitis B e antigen perturbs the LPS-stimulated production of inflammatory cytokines by mononuclear-derived dendritic cells].
Hepatitis A virus vaccine escape variants and potential new serotype emergence.
Chronic inflammation, immune escape, and oncogenesis in the liver: a unique neighborhood for novel intersections.
Hepatitis A virus genotype distribution during a decade of universal vaccination of preadolescents.
Advances in Hepatitis E Virus Biology and Pathogenesis.
Optimized AAV rh.10 Vectors That Partially Evade Neutralizing Antibodies during Hepatic Gene Transfer.
Genotype distribution of genital Chlamydia trachomatis in Chiang Mai, Thailand.
An ImmunoSignature test distinguishes Trypanosoma cruzi, hepatitis B, hepatitis C and West Nile virus seropositivity among asymptomatic blood donors.
On the use of immune checkpoint inhibitors in patients with viral infections including COVID-19.
Severe COVID-19 May Impact Hepatic Fibrosis /Hepatic Stellate Cells Activation as Indicated by a Pathway and Population Genetic Study.
Molecular biology and inhibitors of hepatitis A virus.
ON-1 and BA-IX Are the Dominant Sub-Genotypes of Human Orthopneumovirus A&B in Riyadh, Saudi Arabia.
Genotyping of polyomavirus BK by Real Time PCR for VP1 gene.
Investigation on torquetenovirus (TTV) microRNA transcriptome in vivo.

Table A2-10, Cluster 9

Cluster 9 focuses on immune system evasion by the Lyme Disease Spirochete <i>Borrelia burgdorferi</i> (155)
The role of host immune cells and <i>Borrelia burgdorferi</i> antigens in the etiology of Lyme disease.
Immune escape strategies of <i>Borrelia burgdorferi</i> .
The Brilliance of <i>Borrelia</i> : Mechanisms of Host Immune Evasion by Lyme Disease-Causing Spirochetes. [Lyme disease--new findings on its physiopathology, diagnosis, therapy and prevention].
<i>Borrelia burgdorferi</i> complement regulator-acquiring surface protein 2 does not contribute to complement resistance or host infectivity.
Complement evasion strategies of <i>Borrelia burgdorferi</i> sensu lato.
Insights into the biology of <i>Borrelia burgdorferi</i> gained through the application of molecular genetics.
<i>Borrelia burgdorferi</i> sensu lato diversity and its influence on pathogenicity in humans.
Antibody Response to Lyme Disease Spirochetes in the Context of VlsE-Mediated Immune Evasion.
The role of <i>Borrelia burgdorferi</i> outer surface proteins.
Outer surface protein C is a dissemination-facilitating factor of <i>Borrelia burgdorferi</i> during mammalian infection.
Role of the VlsE Lipoprotein in Immune Avoidance by the Lyme Disease Spirochete <i>Borrelia burgdorferi</i> .
The Infectivity Gene <i>bbk13</i> Is Important for Multiple Phases of the <i>Borrelia burgdorferi</i> Enzootic Cycle.
Lipoproteome screening of the Lyme disease agent identifies inhibitors of antibody-mediated complement killing.
Infectious arthritis and immune dysregulation: lessons from Lyme disease.
New Zealand White Rabbits Effectively Clear <i>Borrelia burgdorferi</i> B31 despite the Bacterium's Functional VlsE Antigenic Variation System.
In vivo expression technology identifies a novel virulence factor critical for <i>Borrelia burgdorferi</i> persistence in mice.
<i>Borrelia burgdorferi</i> Keeps Moving and Carries on: A Review of Borrelial Dissemination and Invasion.
Immune evasion of the Lyme disease spirochetes.
Immune evasion by tickborne and host-adapted <i>Borrelia burgdorferi</i> .
The <i>Borrelia burgdorferi</i> VlsE Lipoprotein Prevents Antibody Binding to an Arthritis-Related Surface Antigen.
Binding of the complement inhibitor C4b-binding protein to Lyme disease <i>Borreliae</i> .
Pleomorphic Variants of <i>Borrelia</i> (syn. <i>Borrelia</i> ) <i>burgdorferi</i> Express Evolutionary Distinct Transcriptomes.
Identification of an <i>ospC</i> operator critical for immune evasion of <i>Borrelia burgdorferi</i> .
<i>Borrelia burgdorferi</i> Pathogenesis and the Immune Response.
Travelling between Two Worlds: Complement as a Gatekeeper for an Expanded Host Range of Lyme Disease Spirochetes.
Outer surface protein polymorphisms linked to host-spirochete association in Lyme <i>borreliae</i> .
Diagnosing <i>Borreliosis</i> .
Variable VlsE is critical for host reinfection by the Lyme disease spirochete.
Human neuroglial cells internalize <i>Borrelia burgdorferi</i> by coiling phagocytosis mediated by Daam1.
<i>Borrelia burgdorferi</i> resistance to a major skin antimicrobial peptide is independent of outer surface lipoprotein content.
Multifunctional and Redundant Roles of <i>Borrelia burgdorferi</i> Outer Surface Proteins in Tissue Adhesion, Colonization, and Complement Evasion.
How do Lyme <i>borrelia</i> organisms cause disease? The quest for virulence determinants().

Distinctive Evasion Mechanisms to Allow Persistence of <i>Borrelia burgdorferi</i> in Different Human Cell Lines.
Genome-wide analysis of <i>Borrelia turcica</i> and 'Candidatus <i>Borrelia taylorii</i> ' shows relapsing fever-like genomes with unique genomic links to Lyme disease <i>Borrelia</i> .
Hide and Seek: How Lyme Disease Spirochetes Overcome Complement Attack.
<i>Borrelia burgdorferi</i> Manipulates Innate and Adaptive Immunity to Establish Persistence in Rodent Reservoir Hosts.
Epitope-Specific Evolution of Human B Cell Responses to <i>Borrelia burgdorferi</i> VlsE Protein from Early to Late Stages of Lyme Disease.
Outer surface proteins of <i>Borrelia</i> : peerless immune evasion tools.
Stability of <i>Borrelia burgdorferi</i> outer surface protein C under immune selection pressure.
[New aspects of pathogenesis of Lyme borreliosis].
Complement Evasion Contributes to Lyme Borreliosis-Host Associations.
Lyme borreliosis--an update.
The emergence of Lyme disease.
Structural basis for complement evasion by Lyme disease pathogen <i>Borrelia burgdorferi</i> .
An immune evasion mechanism for spirochetal persistence in Lyme borreliosis.
Cas9-mediated endogenous plasmid loss in <i>Borrelia burgdorferi</i> .
Vls Antigenic Variation Systems of Lyme Disease <i>Borrelia</i> : Eluding Host Immunity through both Random, Segmental Gene Conversion and Framework Heterogeneity.
New Insights Into CRASP-Mediated Complement Evasion in the Lyme Disease Enzootic Cycle.
Further Insights Into the Interaction of Human and Animal Complement Regulator Factor H With Viable Lyme Disease Spirochetes.
Expression and gene sequence of outer surface protein C of <i>Borrelia burgdorferi</i> reisolated from chronically infected mice.
<i>Borrelia burgdorferi</i> regulates expression of complement regulator-acquiring surface protein 1 during the mammal-tick infection cycle.
Plasticity in early immune evasion strategies of a bacterial pathogen.
Blood treatment of Lyme borreliosis demonstrates the mechanism of CspZ-mediated complement evasion to promote systemic infection in vertebrate hosts.
Review of evidence for immune evasion and persistent infection in Lyme disease.
Crystal structure of Lyme disease variable surface antigen VlsE of <i>Borrelia burgdorferi</i> .
YebC regulates variable surface antigen VlsE expression and is required for host immune evasion in <i>Borrelia burgdorferi</i> .
Crystal structure of the membrane attack complex assembly inhibitor BGA71 from the Lyme disease agent <i>Borrelia bavariensis</i> .
Assessment of transcriptional activity of <i>Borrelia burgdorferi</i> and host cytokine genes during early and late infection in a mouse model.
Complement regulator-acquiring surface proteins of <i>Borrelia burgdorferi</i> : Structure, function and regulation of gene expression.
Bactericidal activity of avian complement: a contribution to understand avian-host tropism of Lyme borreliosis.
MyD88 deficiency enhances acquisition and transmission of <i>Borrelia burgdorferi</i> by <i>Ixodes scapularis</i> ticks.
Interactions between host immune response and antigenic variation that control <i>Borrelia burgdorferi</i> population dynamics.

Role of outer membrane architecture in immune evasion by <i>Treponema pallidum</i> and <i>Borrelia burgdorferi</i> .
Host Immune Evasion by Lyme and Relapsing Fever <i>Borreliae</i> : Findings to Lead Future Studies for <i>Borrelia miyamotoi</i> .
Molecular and immunological characterization of the p83/100 protein of various <i>Borrelia burgdorferi</i> sensu lato strains.
Deciphering the role of Toll-like receptors in humoral responses to <i>Borreliae</i> .
Complement Evasion by Lyme Disease Spirochetes.
<i>Borrelia burgdorferi</i> BBK32 Inhibits the Classical Pathway by Blocking Activation of the C1 Complement Complex.
Relapsing fever spirochetes <i>Borrelia recurrentis</i> and <i>B. duttonii</i> acquire complement regulators C4b-binding protein and factor H.
Generality of Post-Antimicrobial Treatment Persistence of <i>Borrelia burgdorferi</i> Strains N40 and B31 in Genetically Susceptible and Resistant Mouse Strains.
Molecular characterization of <i>Borrelia burgdorferi</i> <i>erp</i> promoter/operator elements.
<i>Borrelia burgdorferi</i> hijacks cellular metabolism of immune cells: Consequences for host defense.
In Vivo Imaging Demonstrates That <i>Borrelia burgdorferi</i> <i>ospC</i> Is Uniquely Expressed Temporally and Spatially throughout Experimental Infection.
<i>Borrelia burgdorferi</i> <i>erp</i> genes are expressed at different levels within tissues of chronically infected mammalian hosts.
Outer surface protein <i>OspC</i> is an antiphagocytic factor that protects <i>Borrelia burgdorferi</i> from phagocytosis by macrophages.
Genomic sequence of a Lyme disease spirochaete, <i>Borrelia burgdorferi</i> .
<i>Borrelia burgdorferi</i> inhibits human neutrophil functions.
The role of VlsE antigenic variation in the Lyme disease spirochete: persistence through a mechanism that differs from other pathogens.
Persistence of <i>Borrelia burgdorferi</i> sensu lato in resolved erythema migrans lesions.
Loss of pathogenic potential after cloning of the low-passage <i>Borrelia burgdorferi</i> ZS7 tick isolate: a cautionary note.
Elucidating the Immune Evasion Mechanisms of <i>Borrelia mayonii</i> , the Causative Agent of Lyme Disease.
The Cross-Talk between Spirochetal Lipoproteins and Immunity.
A chromosomally encoded virulence factor protects the Lyme disease pathogen against host-adaptive immunity.
The failure of immune response evasion by linear plasmid 28-1-deficient <i>Borrelia burgdorferi</i> is attributable to persistent expression of an outer surface protein.
Temporal expression analysis of the <i>Borrelia burgdorferi</i> paralogous gene family 54 genes BBA64, BBA65, and BBA66 during persistent infection in mice.
Investigating the potential role of non-vls genes on linear plasmid 28-1 in virulence and persistence by <i>Borrelia burgdorferi</i> .
Delineating the requirement for the <i>Borrelia burgdorferi</i> virulence factor <i>OspC</i> in the mammalian host.
<i>Borrelia miyamotoi</i> FbpA and FbpB Are Immunomodulatory Outer Surface Lipoproteins With Distinct Structures and Functions.
BBA70 of <i>Borrelia burgdorferi</i> is a novel plasminogen-binding protein.
Middle region of the <i>Borrelia burgdorferi</i> surface-located protein 1 ( <i>Lmp1</i> ) interacts with host chondroitin-6-sulfate and independently facilitates infection.



Complement Evasion in <i>Borrelia</i> spirochetes: Mechanisms and Opportunities for Intervention.
Host tropism determination by convergent evolution of immunological evasion in the Lyme disease system.
Lyme Disease Pathogenesis.
Acquisition and subsequent transmission of <i>Borrelia hermsii</i> by the soft tick <i>Ornithodoros hermsi</i> .
Immune evasion of <i>Borrelia burgdorferi</i> : insufficient killing of the pathogens by complement and antibody.
Lyme borreliosis.
A small intergenic region of <i>lpl17</i> is required for evasion of adaptive immunity and induction of pathology by the Lyme disease spirochete.
Versatile roles of CspA orthologs in complement inactivation of serum-resistant Lyme disease spirochetes.
Chronic or late Lyme neuroborreliosis: analysis of evidence compared to chronic or late neurosyphilis.
"Conformational dynamics of C1r inhibitor proteins from Lyme disease and relapsing fever spirochetes".
The spirochete <i>Borrelia crocidurae</i> causes erythrocyte rosetting during relapsing fever.
Interaction of spirochetes with the host fibrinolytic system and potential roles in pathogenesis.
Mechanism of <i>Borrelia</i> immune evasion by FhbA-related proteins.
Conformational dynamics of complement protease C1r inhibitor proteins from Lyme disease- and relapsing fever-causing spirochetes.
Characterization of unique regions of <i>Borrelia burgdorferi</i> surface-located membrane protein 1.
Variations in the <i>ospB</i> gene of <i>Borrelia burgdorferi</i> result in differences in monoclonal antibody reactivity and in production of escape variants.
Evidence that the variable regions of the central domain of VlsE are antigenic during infection with Lyme disease spirochetes.
Antialarmin effect of tick saliva during the transmission of Lyme disease.
Antigenic Variation in the Lyme Spirochete: Insights into Recombinational Switching with a Suggested Role for Error-Prone Repair.
Detailed analysis of sequence changes occurring during <i>vlsE</i> antigenic variation in the mouse model of <i>Borrelia burgdorferi</i> infection.
Molecular analysis of decorin-binding protein A (DbpA) reveals five major groups among European <i>Borrelia burgdorferi sensu lato</i> strains with impact for the development of serological assays and indicates lateral gene transfer of the <i>dbpA</i> gene.
Phylogenomic Diversity Elucidates Mechanistic Insights into Lyme Borreliae-Host Association.
Disordered lymphoid purine metabolism contributes to the pathogenesis of persistent <i>Borrelia garinii</i> infection in mice.
Evidence that the BBA68 protein (BbCRASP-1) of the Lyme disease spirochetes does not contribute to factor H-mediated immune evasion in humans and other animals.
The pathogenesis of Lyme neuroborreliosis: from infection to inflammation.
Inactivation of genes for antigenic variation in the relapsing fever spirochete <i>Borrelia hermsii</i> reduces infectivity in mice and transmission by ticks.
<i>Treponema pallidum</i> Lipoprotein TP0435 Expressed in <i>Borrelia burgdorferi</i> Produces Multiple Surface/Periplasmic Isoforms and mediates Adherence.
Vectors as Epidemiological Sentinels: Patterns of Within-Tick <i>Borrelia burgdorferi</i> Diversity.
The enolase of <i>Borrelia burgdorferi</i> is a plasminogen receptor released in outer membrane vesicles.
Evidence of past recombination events among the genes encoding the Erp antigens of <i>Borrelia burgdorferi</i> .

Characterization of the vls antigenic variation loci of the Lyme disease spirochaetes <i>Borrelia garinii</i> Ip90 and <i>Borrelia afzelii</i> ACAI.
Rrp1, a cyclic-di-GMP-producing response regulator, is an important regulator of <i>Borrelia burgdorferi</i> core cellular functions.
Central role of the Holliday junction helicase RuvAB in vlsE recombination and infectivity of <i>Borrelia burgdorferi</i> .
BosR functions as a repressor of the ospAB operon in <i>Borrelia burgdorferi</i> .
Transcriptome Assessment of Erythema Migrans Skin Lesions in Patients With Early Lyme Disease Reveals Predominant Interferon Signaling.
CD4 T cell responses in persistent <i>Borrelia burgdorferi</i> infection.
Analysis of the ability of spirochete species associated with relapsing fever, avian borreliosis, and epizootic bovine abortion to bind factor H and cleave C3b.
PlzA is a bifunctional c-di-GMP biosensor that promotes tick and mammalian host-adaptation of <i>Borrelia burgdorferi</i> .
Human complement regulators C4b-binding protein and C1 esterase inhibitor interact with a novel outer surface protein of <i>Borrelia recurrentis</i> .
Strain-specific antibodies reduce co-feeding transmission of the Lyme disease pathogen, <i>Borrelia afzelii</i> .
CD55 Facilitates Immune Evasion by <i>Borrelia crocidurae</i> , an Agent of Relapsing Fever.
Mutation and recombination in the upstream homology box-flanked ospE-related genes of the Lyme disease spirochetes result in the development of new antigenic variants during infection.
Spirochetal Lipoproteins in Pathogenesis and Immunity.
Spirochetal Lipoproteins and Immune Evasion.
Stimulated Immune Response by TruCulture(®) Whole Blood Assay in Patients With European Lyme Neuroborreliosis: A Prospective Cohort Study.
Outer surface protein E (OspE) mediates <i>Borrelia burgdorferi</i> sensu stricto strain-specific complement evasion in the eastern fence lizard, <i>Sceloporus undulatus</i> .
Analysis of variable major protein antigenic variation in the relapsing fever spirochete, <i>Borrelia miyamotoi</i> , in response to polyclonal antibody selection pressure.
Antigenic variation with a twist--the <i>Borrelia</i> story.
Analysis of the OspE determinants involved in binding of factor H and OspE-targeting antibodies elicited during <i>Borrelia burgdorferi</i> infection in mice.
Interaction between <i>Borrelia miyamotoi</i> variable major proteins Vlp15/16 and Vlp18 with plasminogen and complement.
Culture of <i>Borrelia persica</i> and its flagellar antigen in vitro.
Laboratory Diagnosis of Lyme Borreliosis.
The <i>Borrelia afzelii</i> outer membrane protein BAPKO_0422 binds human factor-H and is predicted to form a membrane-spanning $\beta$ -barrel.
Vitronectin binding protein, BOM1093, confers serum resistance on <i>Borrelia miyamotoi</i> .
Identification of conserved antigens for early serodiagnosis of relapsing fever <i>Borrelia</i> .
Structure-function investigation of vsp serotypes of the spirochete <i>Borrelia hermsii</i> .
Emerging roles of pathogens in Alzheimer disease.
Activation of a vmp pseudogene in <i>Borrelia hermsii</i> : an alternate mechanism of antigenic variation during relapsing fever.
Author Correction: Immune evasion of <i>Borrelia miyamotoi</i> : CbiA, a novel outer surface protein exhibiting complement binding and inactivating properties.
Lipid binding orientation within CD1d affects recognition of <i>Borrelia burgdorferi</i> antigens by NKT cells.

Expression of the Tick-Associated Vtp Protein of <i>Borrelia hermsii</i> in a Murine Model of Relapsing Fever.
The Putative Role of Viruses, Bacteria, and Chronic Fungal Biotxin Exposure in the Genesis of Intractable Fatigue Accompanied by Cognitive and Physical Disability.
Genomic blueprint of a relapsing fever pathogen in 15th century Scandinavia.
B1b lymphocytes confer T cell-independent long-lasting immunity.

Table A2-11, Cluster 10

Cluster 10 focuses on mechanisms used by African Swine Fever Virus to evade host immune system response and promote viral replication (158)
The African swine fever virus I10L protein inhibits the NF- $\kappa$ B signaling pathway by targeting IKK $\beta$ .
African Swine Fever Virus F317L Protein Inhibits NF- $\kappa$ B Activation To Evade Host Immune Response and Promote Viral Replication.
Deletion of African Swine Fever Virus (ASFV) H240R Gene Attenuates the Virulence of ASFV by Enhancing NLRP3-Mediated Inflammatory Responses.
The African Swine Fever Virus with MGF360 and MGF505 Deleted Reduces the Apoptosis of Porcine Alveolar Macrophages by Inhibiting the NF- $\kappa$ B Signaling Pathway and Interleukin-1 $\beta$ .
I226R Protein of African Swine Fever Virus Is a Suppressor of Innate Antiviral Responses.
Modulation of Host Antiviral Innate Immunity by African Swine Fever Virus: A Review.
African Swine Fever Virus H240R Protein Inhibits the Production of Type I Interferon through Disrupting the Oligomerization of STING.
African Swine Fever Virus EP364R and C129R Target Cyclic GMP-AMP To Inhibit the cGAS-STING Signaling Pathway.
African Swine Fever Virus E120R Protein Inhibits Interferon Beta Production by Interacting with IRF3 To Block Its Activation.
African Swine Fever Virus MGF360-12L Inhibits Type I Interferon Production by Blocking the Interaction of Importin $\alpha$ and NF- $\kappa$ B Signaling Pathway.
Transcriptome Profiling in Swine Macrophages Infected with African Swine Fever Virus (ASFV) Uncovers the Complex and Close Relationship with Host.
ASFV pD345L protein negatively regulates NF- $\kappa$ B signalling by inhibiting IKK kinase activity.
Immune Escape Mechanism and Vaccine Research Progress of African Swine Fever Virus.
Inhibition of cGAS-STING-TBK1 signaling pathway by DP96R of ASFV China 2018/1.
[Characteristics of African swine fever virus and difficulties in vaccine development].
Sensitivity of African swine fever virus to type I interferon is linked to genes within multigene families 360 and 505.
African swine fever virus protein MGF-505-7R promotes virulence and pathogenesis by inhibiting JAK1- and JAK2-mediated signaling.
African swine fever virus MGF360-9L promotes viral replication by degrading the host protein HAX1.
Research progress on live attenuated vaccine against African swine fever virus.
African Swine Fever Virus Ubiquitin-Conjugating Enzyme Is an Immunomodulator Targeting NF- $\kappa$ B Activation.
African Swine Fever Virus E184L Protein Interacts with Innate Immune Adaptor STING to Block IFN Production for Viral Replication and Pathogenesis.
Proteome Analysis of Swine Macrophages after Infection with Two Genotype II African Swine Fever Isolates of Different Pathogenicity.

MGF360-9L Is a Major Virulence Factor Associated with the African Swine Fever Virus by Antagonizing the JAK/STAT Signaling Pathway.
African swine fever virus controls the host transcription and cellular machinery of protein synthesis.
Progress Toward Development of Effective and Safe African Swine Fever Virus Vaccines.
pMGF505-7R determines pathogenicity of African swine fever virus infection by inhibiting IL-1 $\beta$ and type I IFN production.
African Swine Fever Virus L83L Negatively Regulates the cGAS-STING-Mediated IFN-I Pathway by Recruiting Tollip To Promote STING Autophagic Degradation.
Regulation of antiviral immune response by African swine fever virus (ASFV).
African swine fever virus (ASFV) protection mediated by NH/P68 and NH/P68 recombinant live-attenuated viruses.
Suppression of NF- $\kappa$ B Activity: A Viral Immune Evasion Mechanism.
Identification of African swine fever virus MGF505-2R as a potent inhibitor of innate immunity in vitro.
Evolution of African swine fever virus genes related to evasion of host immune response.
African swine fever virus M1249L protein antagonizes type I interferon production via suppressing phosphorylation of TBK1 and degrading IRF3.
African Swine Fever Virus MGF360-14L Negatively Regulates Type I Interferon Signaling by Targeting IRF3.
Identification and utility of innate immune system evasion mechanisms of ASFV.
African Swine Fever Virus Regulates Host Energy and Amino Acid Metabolism To Promote Viral Replication.
Mechanisms of African swine fever virus pathogenesis and immune evasion inferred from gene expression changes in infected swine macrophages.
African Swine Fever Virus: A Review.
African Swine Fever Virus pF778R Attenuates Type I Interferon Response by Impeding STAT1 Nuclear Translocation.
Pseudorabies Virus Infection of Epithelial Cells Leads to Persistent but Aberrant Activation of the NF- $\kappa$ B Pathway, Inhibiting Hallmark NF- $\kappa$ B-Induced Proinflammatory Gene Expression.
African swine fever virus evasion of host defences.
Loss of IKK Subunits Limits NF- $\kappa$ B Signaling in Reovirus-Infected Cells.
The human cytomegalovirus UL26 protein antagonizes NF- $\kappa$ B activation.
African Swine Fever Virus: A Review.
[The E248R protein of African swine fever virus inhibits the cGAS-STING-mediated innate immunity].
Autophagy impairment by African swine fever virus.
African swine fever virus cysteine protease pS273R inhibits pyroptosis by noncanonically cleaving gasdermin D.
African Swine Fever Virus Cysteine Protease pS273R Inhibits Type I Interferon Signaling by Mediating STAT2 Degradation.
African Swine Fever Virus pI215L Negatively Regulates cGAS-STING Signaling Pathway through Recruiting RNF138 to Inhibit K63-Linked Ubiquitination of TBK1.
Combinational Deletions of MGF360-9L and MGF505-7R Attenuated Highly Virulent African Swine Fever Virus and Conferred Protection against Homologous Challenge.
A parapoxviral virion protein inhibits NF- $\kappa$ B signaling early in infection.
Herpes Simplex Virus 1 Ubiquitin-Specific Protease UL36 Abrogates NF- $\kappa$ B Activation in DNA Sensing Signal Pathway.
Herpes Simplex Virus 1 UL2 Inhibits the TNF- $\alpha$ -Mediated NF- $\kappa$ B Activity by Interacting With p65/p50.

A vaccinia virus deletion mutant reveals the presence of additional inhibitors of NF-kappaB.
African Swine Fever Virus pI215L Inhibits Type I Interferon Signaling by Targeting Interferon Regulatory Factor 9 for Autophagic Degradation.
Proteins in pregnant swine serum promote the African swine fever virus replication: an iTRAQ-based quantitative proteomic analysis.
Inhibition of nuclear factor kappaB activation by a virus-encoded IkappaB-like protein.
African swine fever virus infection of porcine aortic endothelial cells leads to inhibition of inflammatory responses, activation of the thrombotic state, and apoptosis.
Deletion of a CD2-like gene, 8-DR, from African swine fever virus affects viral infection in domestic swine.
African Swine Fever Virus HLI/18 CD2v Suppresses Type I IFN Production and IFN-Stimulated Genes Expression through Negatively Regulating cGMP-AMP Synthase-STING and IFN Signaling Pathways.
Inhibition of NF-κB activity by the porcine epidemic diarrhea virus nonstructural protein 1 for innate immune evasion.
Regulation and Evasion of Host Immune Response by African Swine Fever Virus.
Strategies of NF-κB signaling modulation by ectromelia virus in BALB/3T3 murine fibroblasts.
The Reovirus σ3 Protein Inhibits NF-κB-Dependent Antiviral Signaling.
A novel TLR3 inhibitor encoded by African swine fever virus (ASFV).
African Swine Fever Virus MGF505-7R Interacts with Interferon Regulatory Factor 9 to Evade the Type I Interferon Signaling Pathway and Promote Viral Replication.
How to Inhibit Nuclear Factor-Kappa B Signaling: Lessons from Poxviruses.
African swine fever virus MGF505-7R protein interacted with IRF7 and TBK1 to inhibit type I interferon production.
Current status and evolving approaches to African swine fever vaccine development.
Attempts at the Development of a Recombinant African Swine Fever Virus Strain with Abrogated EP402R, 9GL, and A238L Gene Structure using the CRISPR/Cas9 System.
Orf Virus ORF120 Protein Positively Regulates the NF-κB Pathway by Interacting with G3BP1.
African Swine Fever Virus Structural Protein p17 Inhibits cGAS-STING Signaling Pathway Through Interacting With STING.
Blocking antibodies inhibit complete African swine fever virus neutralization.
Porcine transmissible gastroenteritis virus inhibits NF-κB activity via nonstructural protein 3 to evade host immune system.
African swine fever virus pA104R protein acts as a suppressor of type I interferon signaling.
Antagonisms of ASFV towards Host Defense Mechanisms: Knowledge Gaps in Viral Immune Evasion and Pathogenesis.
Recent advances on viral manipulation of NF-κB signaling pathway.
Varicella-zoster virus inhibition of the NF-κB pathway during infection of human dendritic cells: role for open reading frame 61 as a modulator of NF-κB activity.
African Swine Fever Vaccinology: The Biological Challenges from Immunological Perspectives.
Ectromelia virus encodes a BTB/kelch protein, EVM150, that inhibits NF-κB signaling.
The measles virus V protein binds to p65 (RelA) to suppress NF-kappaB activity.
Pseudorabies Virus ICPO Abolishes Tumor Necrosis Factor Alpha-Induced NF-κB Activation by Degrading P65.
Primate lentiviruses use at least three alternative strategies to suppress NF-κB-mediated immune activation.
The hepatitis B e antigen suppresses IL-1β-mediated NF-κB activation in hepatocytes.

African swine fever virus causes microtubule-dependent dispersal of the trans-golgi network and slows delivery of membrane protein to the plasma membrane.
Comparative proteomic analysis reveals different responses in porcine lymph nodes to virulent and attenuated homologous African swine fever virus strains.
Unraveling the Armor of a Killer: Evasion of Host Defenses by African Swine Fever Virus.
MGF360-12L of ASFV-SY18 is an immune-evasion protein that inhibits host type I IFN, NF- $\kappa$ B, and JAK/STAT pathways.
I329L: A Dual Action Viral Antagonist of TLR Activation Encoded by the African Swine Fever Virus (ASFV).
Herpes simplex virus type 2 inhibits TNF- $\alpha$ -induced NF- $\kappa$ B activation through viral protein ICP22-mediated interaction with p65.
Herpes simplex virus 1 DNA polymerase processivity factor UL42 inhibits TNF- $\alpha$ -induced NF- $\kappa$ B activation by interacting with p65/RelA and p50/NF- $\kappa$ B1.
Examination of Immunogenic Properties of Recombinant Antigens Based on p22 Protein from African Swine Fever Virus.
Herpes Simplex Virus Type 2 Glycoprotein D Inhibits NF- $\kappa$ B Activation by Interacting with p65.
African swine fever virus QP383R dampens type I interferon production by promoting cGAS palmitoylation.
Human Bocavirus NS1 and NS1-70 Proteins Inhibit TNF- $\alpha$ -Mediated Activation of NF- $\kappa$ B by targeting p65.
MERS-CoV 4b protein interferes with the NF- $\kappa$ B-dependent innate immune response during infection.
Molluscum Contagiosum Virus Protein MC005 Inhibits NF- $\kappa$ B Activation by Targeting NEMO-Regulated I $\kappa$ B Kinase Activation.
Suppression of the interferon and NF- $\kappa$ B responses by severe fever with thrombocytopenia syndrome virus.
Influenza A virus-encoded NS1 virulence factor protein inhibits innate immune response by targeting IKK.
The African swine fever virus protease pS273R inhibits DNA sensing cGAS-STING pathway by targeting IKK $\epsilon$ .
The role of infectious hematopoietic necrosis virus (IHNV) proteins in the modulation of NF- $\kappa$ B pathway during IHNV infection.
Gaps in African swine fever: Analysis and priorities.
Enterovirus 71 2C Protein Inhibits NF- $\kappa$ B Activation by Binding to RelA(p65).
Vaccinia Virus BBK E3 Ligase Adaptor A55 Targets Importin-Dependent NF- $\kappa$ B Activation and Inhibits CD8(+) T-Cell Memory.
Viral interference with innate immunity by preventing NF- $\kappa$ B activity.
Poxvirus targeting of E3 ligase $\beta$ -TrCP by molecular mimicry: a mechanism to inhibit NF- $\kappa$ B activation and promote immune evasion and virulence.
African swine fever virus ubiquitin-conjugating enzyme pI215L inhibits IFN-I signaling pathway through STAT2 degradation.
The hepatitis C virus protein NS3 suppresses TNF- $\alpha$ -stimulated activation of NF- $\kappa$ B by targeting LUBAC.
NF- $\kappa$ B activation is a turn on for vaccinia virus phosphoprotein A49 to turn off NF- $\kappa$ B activation.
Herpes simplex virus 1 E3 ubiquitin ligase ICP0 protein inhibits tumor necrosis factor alpha-induced NF- $\kappa$ B activation by interacting with p65/RelA and p50/NF- $\kappa$ B1.
Escape from transcriptional shutoff during poliovirus infection: NF- $\kappa$ B-responsive genes I $\kappa$ B $\alpha$ and A20.

Modulation of proinflammatory NF- $\kappa$ B signaling by ectromelia virus in RAW 264.7 murine macrophages.
Herpes Simplex Virus 1 UL24 Abrogates the DNA Sensing Signal Pathway by Inhibiting NF- $\kappa$ B Activation.
Mumps Virus SH Protein Inhibits NF- $\kappa$ B Activation by Interacting with Tumor Necrosis Factor Receptor 1, Interleukin-1 Receptor 1, and Toll-Like Receptor 3 Complexes.
Herpes simplex virus 1 protein kinase US3 hyperphosphorylates p65/RelA and dampens NF- $\kappa$ B activation.
The VSV matrix protein inhibits NF- $\kappa$ B and the interferon response independently in mouse L929 cells.
Putative E3 ubiquitin ligase of human rotavirus inhibits NF- $\kappa$ B activation by using molecular mimicry to target $\beta$ -TrCP.
Middle East Respiratory Syndrome Coronavirus-Encoded Accessory Proteins Impair MDA5- and TBK1-Mediated Activation of NF- $\kappa$ B.
The accessory protein ORF3 of porcine epidemic diarrhea virus inhibits cellular interleukin-6 and interleukin-8 productions by blocking the nuclear factor- $\kappa$ B p65 activation.
Vaccinia virus BTB-Kelch proteins C2 and F3 inhibit NF- $\kappa$ B activation.
NSP1 of human rotaviruses commonly inhibits NF- $\kappa$ B signalling by inducing $\beta$ -TrCP degradation.
miR-615 facilitates porcine epidemic diarrhea virus replication by targeting IRAK1 to inhibit type III interferon expression.
Modulation of NF- $\kappa$ B transcription factor activation by Mollusum contagiosum virus proteins.
The role of the IKK complex in viral infections.
HIV-1 Vpu is a potent transcriptional suppressor of NF- $\kappa$ B-elicited antiviral immune responses.
Vaccination influences the evolution of classical swine fever virus.
The Main DNA Viruses Significantly Affecting Pig Livestock.
Human coronavirus OC43 nucleocapsid protein binds microRNA 9 and potentiates NF- $\kappa$ B activation.
The viral protein A238L inhibits TNF- $\alpha$ expression through a CBP/p300 transcriptional coactivators pathway.
Human T-cell lymphotropic virus: a model of NF- $\kappa$ B-associated tumorigenesis.
Role of innate immunity in pathophysiology of classical swine fever virus infection.
The Npro product of classical swine fever virus interacts with I $\kappa$ B $\alpha$ , the NF- $\kappa$ B inhibitor.
ORF3 of Hepatitis E Virus Inhibits the Expression of Proinflammatory Cytokines and Chemotactic Factors in LPS-Stimulated Human PMA-THP1 Cells by Inhibiting NF- $\kappa$ B Pathway.
Patterns of gene expression in swine macrophages infected with classical swine fever virus detected by microarray.
The ORF61 Protein Encoded by Simian Varicella Virus and Varicella-Zoster Virus Inhibits NF- $\kappa$ B Signaling by Interfering with I $\kappa$ B $\alpha$ Degradation.
Interactions of Vank proteins from Microplitis bicoloratus bracovirus with host Dip3 suppress eIF4E expression.
Inhibition of interferon gene activation by death-effector domain-containing proteins from the mollusum contagiosum virus.
Mollusum Contagiosum Virus MC159 Abrogates cIAP1-NEMO Interactions and Inhibits NEMO Polyubiquitination.
Myxoma virus M013 protein antagonizes NF- $\kappa$ B and inflammasome pathways via distinct structural motifs.
Co-infection of porcine deltacoronavirus and porcine epidemic diarrhea virus induces early TRAF6-mediated NF- $\kappa$ B and IRF7 signaling pathways through TLRs.
IIV-6 Inhibits NF- $\kappa$ B Responses in Drosophila.

Murine gammaherpesvirus 68 evades host cytokine production via replication transactivator-induced RelA degradation.
In vitro infection with classical swine fever virus inhibits the transcription of immune response genes.
The molluscum contagiosum virus death effector domain containing protein MC160 RxDL motifs are not required for its known viral immune evasion functions.
HIV-1 Vpr antagonizes innate immune activation by targeting karyopherin-mediated NF- $\kappa$ B/IRF3 nuclear transport.
Coinfection of Porcine Circovirus 2 and Pseudorabies Virus Enhances Immunosuppression and Inflammation through NF- $\kappa$ B, JAK/STAT, MAPK, and NLRP3 Pathways.
Dengue virus downregulates TNFR1- and TLR3-stimulated NF- $\kappa$ B activation by targeting RIPK1.
A novel mechanism of nuclear factor-kappaB regulation by adenoviral protein 14.7K.
Japanese Encephalitis Virus exploits microRNA-155 to suppress the non-canonical NF- $\kappa$ B pathway in human microglial cells.
Pathogenic Old World arenaviruses inhibit TLR2/Mal-dependent proinflammatory cytokines in vitro.
miR-146a suppresses cellular immune response during Japanese encephalitis virus JaOArS982 strain infection in human microglial cells.
Activation and modulation of antiviral and apoptotic genes in pigs infected with classical swine fever viruses of high, moderate or low virulence.
CD40 induces antigen transporter and immunoproteasome gene expression in carcinomas via the coordinated action of NF-kappaB and of NF-kappaB-mediated de novo synthesis of IRF-1.
Porcine Epidemic Diarrhea Virus nsp13 Protein Downregulates Neonatal Fc Receptor Expression by Causing Promoter Hypermethylation through the NF- $\kappa$ B Signaling Pathway.
miR-146a promotes Borna disease virus 1 replication through IRAK1/TRAF6/NF- $\kappa$ B signaling pathway.
U20 is responsible for human herpesvirus 6B inhibition of tumor necrosis factor receptor-dependent signaling and apoptosis.
Transcriptome Profiling of Vero E6 Cells during Original Parental or Cell-Attenuated Porcine Epidemic Diarrhea Virus Infection.
Exosomes mediate the antibody-resistant intercellular transmission of porcine epidemic diarrhea virus.

Table A2-12, Cluster 11

Cluster 11 focuses on HLA Class 1 expression, emphasizing immune system evasion by cancer resulting from HLA Class 1 loss (278)
Total loss of HLA class I expression on a melanoma cell line after growth in nude mice in absence of autologous antitumor immune response.
Role of altered expression of HLA class I molecules in cancer progression.
Prognostic Value of HLA Class I in Patients with Hepatocellular Carcinoma.
Different mechanisms can lead to the same altered HLA class I phenotype in tumors.
Loss of heterozygosity at 6p21 underlying [corrected] HLA class I downregulation in Chinese primary esophageal squamous cell carcinomas.
HLA class I expression in metastatic melanoma correlates with tumor development during autologous vaccination.
Loss of heterozygosity at 6p21 and HLA class I expression in esophageal squamous cell carcinomas in China.
Loss of heterozygosity in the HLA class I region in human pancreatic cancer.



Cancer Immunology: Immune Escape of Tumors-Expression and Regulation of HLA Class I Molecules and Its Role in Immunotherapies.
Analysis of HLA class I expression in progressing and regressing metastatic melanoma lesions after immunotherapy.
Haplotype loss of HLA class I antigen as an escape mechanism from immune attack in lung cancer.
Targetless T cells in cancer immunotherapy.
High-resolution analysis of HLA class I alterations in colorectal cancer.
The prognostic impact of programmed cell death ligand 1 and human leukocyte antigen class I in pancreatic cancer.
Human Leukocyte Antigen Class I and Programmed Death-Ligand 1 Coexpression Is an Independent Poor Prognostic Factor in Adenocarcinoma of the Lung.
Regressing and progressing metastatic lesions: resistance to immunotherapy is predetermined by irreversible HLA class I antigen alterations.
High frequency of HLA-B44 allelic losses in human solid tumors.
HLA class I expression on the materno-fetal interface.
The Challenges of HLA Class I Loss in Cancer Immunotherapy: Facts and Hopes.
HNPCC versus sporadic microsatellite-unstable colon cancers follow different routes toward loss of HLA class I expression.
Loss of heterozygosity at 6p21.3 underlying HLA class I downregulation in gastric cancer.
Expression of transporters associated with antigen processing and human leucocyte antigen class I in malignant melanoma and its association with prognostic factors.
MUTYH-associated polyposis carcinomas frequently lose HLA class I expression - a common event amongst DNA-repair-deficient colorectal cancers.
The role of classical and non-classical HLA class I antigens in human tumors.
HLA class I antigen processing machinery defects in antitumor immunity and immunotherapy.
[Tissue array analysis of the aberrant expression of HLA class I molecules in human non small cell lung cancer].
Prognostic significance of HLA class I and II expression in patients with diffuse large B cell lymphoma treated with standard chemoimmunotherapy.
Multiple mechanisms generate HLA class I altered phenotypes in laryngeal carcinomas: high frequency of HLA haplotype loss associated with loss of heterozygosity in chromosome region 6p21.
High frequency of homozygosity of the HLA region in melanoma cell lines reveals a pattern compatible with extensive loss of heterozygosity.
Elevated expression of SerpinA1 and SerpinA3 in HLA-positive cervical carcinoma.
Cytokeratin 8-MHC class I interactions: a potential novel immune escape phenotype by a lymph node metastatic carcinoma cell line.
Expression of HLA class I, beta(2)-microglobulin and HLA class II antigens in primary orbital melanoma.
The Escape of Cancer from T Cell-Mediated Immune Surveillance: HLA Class I Loss and Tumor Tissue Architecture.
HLA antigen and NK cell activating ligand expression in malignant cells: a story of loss or acquisition.
HLA expression in uveal melanoma: there is no rule without some exception.
Clinical significance of expression of cancer/testis antigen and down-regulation of HLA class-I in patients with stage I non-small cell lung cancer.
Alterations in classical and nonclassical HLA expression in recurrent and progressive HPV-induced usual vulvar intraepithelial neoplasia and implications for immunotherapy.
Revisiting immune escape in colorectal cancer in the era of immunotherapy.

Down-regulation of HLA class I antigen-processing machinery components in esophageal squamous cell carcinomas: association with disease progression.
Adenovirus expressing $\beta$ 2-microglobulin recovers HLA class I expression and antitumor immunity by increasing T-cell recognition.
Down-regulation of HLA class I antigen processing molecules: an immune escape mechanism of renal cell carcinoma?
Distribution of HLA class I altered phenotypes in colorectal carcinomas: high frequency of HLA haplotype loss associated with loss of heterozygosity in chromosome region 6p21.
Targeting HLA class I expression to increase tumor immunogenicity.
LOH at 6p21.3 region and HLA class I altered phenotypes in bladder carcinomas.
Multiple mechanisms underlie HLA dysregulation in cervical cancer.
Mechanisms of loss of HLA class I expression on colorectal tumor cells.
Molecular mechanisms of HLA class I-mediated immune evasion of human tumors and their role in resistance to immunotherapies.
Downregulation of HLA class II is associated with relapse after allogeneic stem cell transplantation and alters recognition by antigen-specific T cells.
Circulating human papillomavirus type 16 specific T-cells are associated with HLA Class I expression on tumor cells, but not related to the amount of viral oncogene transcripts.
Frequent HLA class I alterations in human prostate cancer: molecular mechanisms and clinical relevance.
Prognostic impact of anticancer immune responses: an introduction.
The expression of histocompatibility-related leukocyte antigens in the pathway to cervical carcinoma.
The HLA crossroad in tumor immunology.
HLA class I expression and chromosomal deletions at 6p and 15q in head and neck squamous cell carcinomas.
HLA Class I Antigen Processing Machinery Defects in Cancer Cells-Frequency, Functional Significance, and Clinical Relevance with Special Emphasis on Their Role in T Cell-Based Immunotherapy of Malignant Disease.
Association of antigen processing machinery and HLA class I defects with clinicopathological outcome in cervical carcinoma.
Low prevalence of selective human leukocyte antigen (HLA)-A and HLA-B epitope losses in early-passage tumor cell lines.
Loss of HLA haplotype and B locus down-regulation in melanoma cell lines.
MHC/HLA Class I Loss in Cancer Cells.
Regulation of HLA class I surface expression requires CD99 and p230/golgin-245 interaction.
Human leukocyte antigen and antigen processing machinery component defects in astrocytic tumors.
HLA class I alterations in breast carcinoma are associated with a high frequency of the loss of heterozygosity at chromosomes 6 and 15.
Genome-wide differential genetic profiling characterizes colorectal cancers with genetic instability and specific routes to HLA class I loss and immune escape.
Copy Neutral LOH Affecting the Entire Chromosome 6 Is a Frequent Mechanism of HLA Class I Alterations in Cancer.
HLA Class I Downregulation in Progressing Metastases of Melanoma Patients Treated With Ipilimumab.
HLA class I loss and PD-L1 expression in lung cancer: impact on T-cell infiltration and immune escape.
Myeloma cells resistance to NK cell lysis mainly involves an HLA class I-dependent mechanism.
Comparison of surface HLA class I levels in squamous cell carcinoma cell lines of the head and neck.

The absence of HLA class I expression in non-small cell lung cancer correlates with the tumor tissue structure and the pattern of T cell infiltration.
Role of Human Leukocyte Antigen System as A Predictive Biomarker for Checkpoint-Based Immunotherapy in Cancer Patients.
PD-L1 and HLA Class I Antigen Expression and Clinical Course of the Disease in Intrahepatic Cholangiocarcinoma.
Transporter associated protein expression in uveal melanoma.
Reduced human leukocyte antigen expression in advanced-stage Ewing sarcoma: implications for immune recognition.
Cancer immune escape: MHC expression in primary tumours versus metastases.
Down-regulation of locus-specific human lymphocyte antigen class I expression in Epstein-Barr virus-associated gastric cancer: implication for viral-induced immune evasion.
Immune escape pathways from the HBV core(18-27) CD8 T cell response are driven by individual HLA class I alleles.
The SPPL3-Defined Glycosphingolipid Repertoire Orchestrates HLA Class I-Mediated Immune Responses.
HLA class I antigen loss, tumor immune escape and immune selection.
Less correlation between mismatch repair proteins deficiency and decreased expression of HLA class I molecules in endometrial carcinoma: a different propensity from colorectal cancer.
HLA class I loss in colorectal cancer: implications for immune escape and immunotherapy.
Frequent loss of heterozygosity in the $\beta$ 2-microglobulin region of chromosome 15 in primary human tumors.
Identification of a lung cancer antigen evading CTL attack due to loss of human leukocyte antigen (HLA) class I expression.
Human leukocyte antigen class II expression is a good prognostic factor in adult T-cell leukemia/lymphoma.
Immune escape associated with functional defects in antigen-processing machinery in head and neck cancer.
Down-regulation of human leukocyte antigens class I on peripheral T lymphocytes and NK cells from subjects in region of high-incidence gastrointestinal tumor.
Multiple structural and epigenetic defects in the human leukocyte antigen class I antigen presentation pathway in a recurrent metastatic melanoma following immunotherapy.
Immunological and clinical significance of HLA class I antigen processing machinery component defects in malignant cells.
The low rate of HLA class I molecules on the human embryonic stem cell line HS293 is associated with the APM components' expression level.
Human cytomegalovirus and natural killer-mediated surveillance of HLA class I expression: a paradigm of host-pathogen adaptation.
Identification of different tumor escape mechanisms in several metastases from a melanoma patient undergoing immunotherapy.
c-Myc and EBV-LMP1: two opposing regulators of the HLA class I antigen presentation machinery in epithelial cells.
A mechanism for evasion of CTL immunity by altered O-glycosylation of HLA class I.
Tumor Escape Phenotype in Bladder Cancer Is Associated with Loss of HLA Class I Expression, T-Cell Exclusion and Stromal Changes.
Identification of microRNAs Targeting the Transporter Associated with Antigen Processing TAP1 in Melanoma.

Evidence for whole chromosome 6 loss and duplication of the remaining chromosome in acute lymphoblastic leukemia.
Involvement of HLA class I molecules in the immune escape of urologic tumors.
Quantification of classical HLA class I mRNA by allele-specific, real-time polymerase chain reaction for most Han individuals.
HLA dependent immune escape mechanisms in B-cell lymphomas: Implications for immune checkpoint inhibitor therapy?
Decreased expression of B7 costimulatory molecules and major histocompatibility complex class-I in human hepatocellular carcinoma.
The transition from HLA-I positive to HLA-I negative primary tumors: the road to escape from T-cell responses.
Loss of heterozygosity, a frequent but a non-exclusive mechanism responsible for HLA dysregulation in non-Hodgkin's lymphomas.
Immune escape of cancer cells with beta2-microglobulin loss over the course of metastatic melanoma.
Mismatch Repair Protein Deficiency Is a Risk Factor for Aberrant Expression of HLA Class I Molecules: A Putative "Adaptive Immune Escape" Phenomenon.
HLA expression at the maternal-fetal interface.
Loss of expression of antigen-presenting molecules in human pancreatic cancer and pancreatic cancer cell lines.
MHC class-I downregulation in PD-1/PD-L1 inhibitor refractory Merkel cell carcinoma and its potential reversal by histone deacetylase inhibition: a case series.
Loss of HLA class I and mismatch repair protein expression in sporadic endometrioid endometrial carcinomas.
Total loss of MHC class I in colorectal tumors can be explained by two molecular pathways: beta2-microglobulin inactivation in MSI-positive tumors and LMP7/TAP2 downregulation in MSI-negative tumors.
Characterization of the immune escape phenotype of human gastric cancers with and without high-frequency microsatellite instability.
Locus-specific analysis of human leukocyte antigen class I expression in melanoma cell lines.
Pan-Cancer HLA Gene-Mediated Tumor Immunogenicity and Immune Evasion.
Genetic alterations of HLA-class II in ovarian cancer.
A Novel Tongue Squamous Cell Carcinoma Cell Line Escapes from Immune Recognition due to Genetic Alterations in HLA Class I Complex.
Association of defective HLA-I expression with antigen processing machinery and their association with clinicopathological characteristics in Kazak patients with esophageal cancer.
STAT1-Induced HLA Class I Upregulation Enhances Immunogenicity and Clinical Response to Anti-EGFR mAb Cetuximab Therapy in HNC Patients.
Human leucocyte antigen class I and class II antigen expression in malignant fibrous histiocyteoma, fibrosarcoma and dermatofibrosarcoma protuberans is significantly downregulated.
Up-regulate HLA class I expression following hepatitis B virus transfection in a hepatocellular carcinoma cell line BEL7405.
Promiscuous binding of invariant chain-derived CLIP peptide to distinct HLA-I molecules revealed in leukemic cells.
Tumor Microenvironment, HLA Class I and APM Expression in HPV-Negative Oral Squamous Cell Carcinoma.

Tumor-specific cytotoxic T lymphocyte responses against chondrosarcoma with HLA haplotype loss restricted by the remaining HLA class I allele.
Extensive genetic alterations of the HLA region, including homozygous deletions of HLA class II genes in B-cell lymphomas arising in immune-privileged sites.
Epithilone B enhances Class I HLA and HLA-A2 surface molecule expression in ovarian cancer cells.
HLA class I loss in metachronous metastases prevents continuous T cell recognition of mutated neoantigens in a human melanoma model.
Tapasin and human leukocyte antigen class I dysregulation correlates with survival in glioblastoma multiforme.
Pervasiveness of HLA allele-specific expression loss across tumor types.
HLA class I is most tightly linked to levels of tapasin compared with other antigen-processing proteins in glioblastoma.
Flow Cytometry Analysis to Detect Lapatinib-Induced Modulation of Constitutive and IFN- $\gamma$ -Induced HLA Class I Expression in HER2-Positive Breast Cancer Cells.
Low frequency of HLA haplotype loss associated with loss of heterozygosity in chromosome region 6p21 in clear renal cell carcinomas.
Loss of Human Leukocyte Antigen Class I Expression Is Associated with Poor Prognosis in Patients with Advanced Breast Cancer.
Somatic HLA Class I Loss Is a Widespread Mechanism of Immune Evasion Which Refines the Use of Tumor Mutational Burden as a Biomarker of Checkpoint Inhibitor Response.
Involvement of the chaperone tapasin in HLA-B44 allelic losses in colorectal tumors.
Landscape of helper and regulatory antitumour CD4(+) T cells in melanoma.
Down-regulation of HLA class I antigen in human papillomavirus type 16 E7 expressing HaCaT cells: correlate with TAP-1 expression.
Methylation of the promoter of human leukocyte antigen class I in human esophageal squamous cell carcinoma and its histopathological characteristics.
HLA antigen changes in malignant cells: epigenetic mechanisms and biologic significance.
Natural history of HLA expression during tumour development.
TEIPP antigens for T-cell based immunotherapy of immune-edited HLA class I(low) cancers.
Identification of Novel HLA Class II-Restricted Neoantigens Derived from Driver Mutations.
BMI1 induces ubiquitination and protein degradation of Nod-like receptor family CARD domain containing 5 and suppresses human leukocyte antigen class I expression to induce immune escape in non-small cell lung cancer.
Differential expression of melanoma-associated antigens and molecules involved in antigen processing and presentation in three cell lines established from a single patient.
Melanoma Lesions Independently Acquire T-cell Resistance during Metastatic Latency.
Natural and cryptic peptides dominate the immunopeptidome of atypical teratoid rhabdoid tumors.
Defective HLA Class I Expression and Patterns of Lymphocyte Infiltration in Chordoma Tumors.
Human leukocyte antigen expression in paired primary lung tumors and brain metastases in non-small cell lung cancer.
Lack of HLA class II antigen expression in microsatellite unstable colorectal carcinomas is caused by mutations in HLA class II regulatory genes.
beta2-Microglobulin mutations, HLA class I antigen loss, and tumor progression in melanoma.
Expression of endoplasmic reticulum aminopeptidases in EBV-B cell lines from healthy donors and in leukemia/lymphoma, carcinoma, and melanoma cell lines.

Immune selection of hot-spot beta 2-microglobulin gene mutations, HLA-A2 allospecificity loss, and antigen-processing machinery component down-regulation in melanoma cells derived from recurrent metastases following immunotherapy.
Molecular basis for lack of expression of HLA class I antigens in human small-cell lung carcinoma cell lines.
Probable HLA-mediated immunoediting of JAK2 V617F-driven oncogenesis.
WHO grade associated downregulation of MHC class I antigen-processing machinery components in human astrocytomas: does it reflect a potential immune escape mechanism?
Identification of non-mutated neoantigens presented by TAP-deficient tumors.
Natural variation of the expression of HLA and endogenous antigen modulates CTL recognition in an in vitro melanoma model.
Analysis of the expression of HLA class I, proinflammatory cytokines and chemokines in primary tumors from patients with localized and metastatic renal cell carcinoma.
HLA class II antigen presentation by prostate cancer cells.
Characterization of human lymphocyte antigen class I antigen-processing machinery defects in renal cell carcinoma lesions with special emphasis on transporter-associated with antigen-processing down-regulation.
Loss of human leucocyte antigen class I and gain of class II expression are early events in carcinogenesis: clues from a study of Barrett's oesophagus.
Role of gene methylation in antitumor immune response: implication for tumor progression.
Post-transcriptional and epigenetic regulation of antigen processing machinery (APM) components and HLA-I in cervical cancers from Uighur women.
Identification of multiple antigens recognized by tumor-infiltrating lymphocytes from a single patient: tumor escape by antigen loss and loss of MHC expression.
Complex pattern of immune evasion in MSI colorectal cancer.
Frequent HLA-DR loss on hematopoietic stem progenitor cells in patients with cyclosporine-dependent aplastic anemia carrying HLA-DR15.
The prevalence of HLA-I LOH in Chinese pan-cancer patients and genomic features of patients harboring HLA-I LOH.
HLA genotyping meets response to immune checkpoint inhibitors prediction: A story just started.
Identification of miR-200a-5p targeting the peptide transporter TAP1 and its association with the clinical outcome of melanoma patients.
The coincidence of chromosome 15 aberrations and beta2-microglobulin gene mutations is causative for the total loss of human leucocyte antigen class I expression in melanoma.
Sex-specific differences in immunogenomic features of response to immune checkpoint blockade.
Epstein-Barr virus latent membrane protein 2A mediated activation of Sonic Hedgehog pathway induces HLA class Ia downregulation in gastric cancer cells.
Relapse Following Allogeneic Hematopoietic Cell Transplantation for Acute Myeloid Leukemia Apparently Due to Somatic Cell Evolution via Epigenetic Variation and Immune Selection.
Permanent, lowered HLA class I expression using lentivirus vectors with shRNA constructs: Averting cytotoxicity by alloreactive T lymphocytes.
Human Leukocyte Antigen Class I Deficiency in Gastric Carcinoma: An Adaptive Immune Evasion Strategy Most Common in Microsatellite Instable Tumors.
HLA expression as a risk factor for metastases of cutaneous squamous-cell carcinoma in organ-transplant recipients.
Pathogenicity and impact of HLA class I alleles in aplastic anemia patients of different ethnicities.

ALK and RET Inhibitors Promote HLA Class I Antigen Presentation and Unmask New Antigens within the Tumor Immunopeptidome.
High level of aneuploidy of chromosome 6 by FISH analysis of head and neck squamous cell carcinoma: limited applicability of LOH analysis to define HLA loss.
$\gamma\delta$ T cells unveil invisible tumors.
HLA-A, -B, -C expression in colon carcinoma mimics that of the normal colonic mucosa and is prognostically relevant.
Disruption of HLA class II antigen presentation in Burkitt lymphoma: implication of a 47,000 MW acid labile protein in CD4+ T-cell recognition.
Regression of metastatic Merkel cell carcinoma following transfer of polyomavirus-specific T cells and therapies capable of re-inducing HLA class-I.
Abrogation of HLA surface expression using CRISPR/Cas9 genome editing: a step toward universal T cell therapy.
Molecular landscape of immune pressure and escape in aplastic anemia.
Expression of transporter associated with antigen processing 1 and 2 (TAP1/2) in malignant melanoma cell lines.
B7-H abnormalities in melanoma and clinical relevance.
Clonal evolution including partial loss of human leukocyte antigen genes favoring extramedullary acute myeloid leukemia relapse after matched related allogeneic hematopoietic stem cell transplantation.
Elevated neoantigen levels in tumors with somatic mutations in the HLA-A, HLA-B, HLA-C and B2M genes.
Integration of tumor extrinsic and intrinsic features associates with immunotherapy response in non-small cell lung cancer.
Predominance of TH2 cells and plasma cells in polyoma virus nephropathy: a role for humoral immunity?
Prognostic impact of human leukocyte antigen class I expression and association of platinum resistance with immunologic profiles in epithelial ovarian cancer.
Sequential immune escape and shifting of T cell responses in a long-term survivor of melanoma.
HLA Class II Loss and JAK1/2 Deficiency Coevolve in Melanoma Leading to CD4 T-cell and IFN $\gamma$ Cross-Resistance.
Presentation of an immunodominant immediate-early CD8+ T cell epitope resists human cytomegalovirus immunoevasion.
Programmed death ligand-1, tumor infiltrating lymphocytes and HLA expression in Chinese extrahepatic cholangiocarcinoma patients: Possible immunotherapy implications.
TGF- $\beta$ and EGF induced HLA-I downregulation is associated with epithelial-mesenchymal transition (EMT) through upregulation of snail in prostate cancer cells.
Expression of human leukocyte antigen class I and $\beta$ 2-microglobulin in colorectal cancer and its prognostic impact.
Effect of platinum-based chemotherapy on the expression of natural killer group 2 member D ligands, programmed cell death-1 ligand 1 and HLA class I in non-small cell lung cancer.
Antitumor activity of human papillomavirus type 16 E7-specific T cells against virally infected squamous cell carcinoma of the head and neck.
KIR2DS3 is associated with protection against acute myeloid leukemia.
Assessment of Impact of Human Leukocyte Antigen-Type and Cytokine-Type Responses on Outcomes after Targeted Therapy Currently Used to Treat Chronic Lymphocytic Leukemia.

Alterations in HLA Class I-Presented Immunoepitome and Class I-Interactome upon Osimertinib Resistance in EGFR Mutant Lung Adenocarcinoma.
Elevation of c-MYC disrupts HLA class II-mediated immune recognition of human B cell tumors.
HPV-16 E5 down-regulates expression of surface HLA class I and reduces recognition by CD8 T cells.
Association of human papillomavirus type 16 E7 and HLA class I antigen expression in cervical premalignant and malignant lesions.
Integrated Multiomic Profiling Identifies the Epigenetic Regulator PRC2 as a Therapeutic Target to Counteract Leukemia Immune Escape and Relapse.
An expanded peripheral T cell population to a cytotoxic T lymphocyte (CTL)-defined, melanocyte-specific antigen in metastatic melanoma patients impacts on generation of peptide-specific CTLs but does not overcome tumor escape from immune surveillance in metastatic lesions.
HLA class I antigen processing machinery component expression and intratumoral T-Cell infiltrate as independent prognostic markers in ovarian carcinoma.
Purification of soluble HLA class I complexes from human serum or plasma deliver high quality immuno peptidomes required for biomarker discovery.
From brain to testis: immune escape and clonal selection in a B cell lymphoma with selective outgrowth in two immune sanctuaries [correction of sanctuaries].
IL-2 enhanced MHC class I expression in papillary thyroid cancer with Hashimoto's thyroiditis overcomes immune escape in vitro.
The mutation in the ATP-binding region of JAK1, identified in human uterine leiomyosarcomas, results in defective interferon-gamma inducibility of TAP1 and LMP2.
Expression of HLA class I, beta(2)-microglobulin, TAP1 and IL-10 in Epstein-Barr virus-associated nasal NK/T-cell lymphoma: Implications for tumor immune escape mechanism.
HLA Class I Analysis Provides Insight Into the Genetic and Epigenetic Background of Immune Evasion in Colorectal Cancer With High Microsatellite Instability.
Deregulation of HLA-I in cancer and its central importance for immunotherapy.
MYCN: from oncoprotein to tumor-associated antigen.
Tumor genetic alterations and features of the immune microenvironment drive myelodysplastic syndrome escape and progression.
Immunoproteasome deficiency is a feature of non-small cell lung cancer with a mesenchymal phenotype and is associated with a poor outcome.
Immune escape mechanism behind resistance to anti-PD-1 therapy in gastrointestinal tract metastasis in malignant melanoma patients with multiple metastases.
Antigen presentation in cancer: insights into tumour immunogenicity and immune evasion.
Adenosine augments the production of IL-10 in cervical cancer cells through interaction with the A(2B) adenosine receptor, resulting in protection against the activity of cytotoxic T cells.
Genetic immune escape landscape in primary and metastatic cancer.
Recurrent hepatocellular carcinoma cells with stem cell-like properties: possible targets for immunotherapy.
SHP2 is overexpressed and inhibits pSTAT1-mediated APM component expression, T-cell attracting chemokine secretion, and CTL recognition in head and neck cancer cells.
TEIPP peptides: exploration of unTAPped cancer antigens.
Epigenetic changes within the promoter regions of antigen processing machinery family genes in Kazakh primary esophageal squamous cell carcinoma.
Acquired IFN $\gamma$ resistance impairs anti-tumor immunity and gives rise to T-cell-resistant melanoma lesions.
Anomalous expression of the HLA-DR alpha and beta chains in ovarian and other cancers.



Class II transactivator (CIITA) isoform expression and activity in melanoma.
Immunity to cancer: attack and escape in T lymphocyte-tumor cell interaction.
Leukemia relapse via genetic immune escape after allogeneic hematopoietic cell transplantation.
In vitro evidence of the presence of mesenchymal stromal cells in cervical cancer and their role in protecting cancer cells from cytotoxic T cell activity.
Immune Modulatory microRNAs Involved in Tumor Attack and Tumor Immune Escape.
Tryptophan metabolism promotes immune evasion in human pancreatic $\beta$ cells.
Genetically engineered fixed K562 cells: potent "off-the-shelf" antigen-presenting cells for generating virus-specific T cells.
Regulation of multiple myeloma survival and progression by CD1d.
Absence of class II-associated invariant chain peptide on leukemic blasts of patients promotes activation of autologous leukemia-reactive CD4+ T cells.
Generation of Retinal Pigment Epithelial Cells Derived from Human Embryonic Stem Cells Lacking Human Leukocyte Antigen Class I and II.
Novel HLA-Cw8-restricted T cell epitopes derived from tyrosinase-related protein-2 and gp100 melanoma antigens.
Burkitt lymphoma: pathogenesis and immune evasion.
A robust and scalable TCR-based reporter cell assay to measure HIV-1 Nef-mediated T cell immune evasion.
Melanomas with concordant loss of multiple melanocytic differentiation proteins: immune escape that may be overcome by targeting unique or undefined antigens.
High numbers of PDCD1 (PD-1)-positive T cells and B2M mutations in microsatellite-unstable colorectal cancer.
Comparative expression profiling for human endoplasmic reticulum-resident aminopeptidases 1 and 2 in normal kidney versus distinct renal cell carcinoma subtypes.
Key Aspects of the Immunobiology of Haploidentical Hematopoietic Cell Transplantation.
Microsatellite instability derived JAK1 frameshift mutations are associated with tumor immune evasion in endometrioid endometrial cancer.
Activation of ER $\alpha$ signaling differentially modulates IFN- $\gamma$ induced HLA-class II expression in breast cancer cells.
Characterization of small spheres derived from various solid tumor cell lines: are they suitable targets for T cells?
Control of HIV-1 immune escape by CD8 T cells expressing enhanced T-cell receptor.
Epigenetic Regulation to Enhance Graft-versus-Leukemia Activity.
IL-15 is produced by a subset of human melanomas, and is involved in the regulation of markers of melanoma progression through juxtacrine loops.
Implications of altered O-glycosylation in tumour immune evasion.
Tumor immune escape in acute myeloid leukemia: Class II-associated invariant chain peptide expression as result of deficient antigen presentation.
Immune regulation in multiple myeloma: the host-tumour conflict.
Immune escape mechanisms in malignant melanoma.
Immune escape from NY-ESO-1-specific T-cell therapy via loss of heterozygosity in the MHC.
Engineering universal cells that evade immune detection.
Exosomes released by EBV-infected nasopharyngeal carcinoma cells convey the viral latent membrane protein 1 and the immunomodulatory protein galectin 9.
HTLV-1 infection promotes excessive T cell activation and transformation into adult T cell leukemia/lymphoma.

Selection for beta 2-microglobulin mutation in mismatch repair-defective colorectal carcinomas.
Recruitment of Oligoclonal Viral-Specific T cells to Kill Human Tumor Cells Using Single-Chain Antibody-Peptide-HLA Fusion Molecules.
A Case Report of Aggressive Fumarate Hydrase-deficient Renal Cell Carcinoma With Loss of HLA Antigens.
T-Cell Transfer Therapy Targeting Mutant KRAS in Cancer.
Lack of B7 expression, not human leukocyte antigen expression, facilitates immune evasion by human malignant gliomas.
Mechanisms of escape from CD8+ T-cell clones specific for the HER-2/neu proto-oncogene expressed in ovarian carcinomas: related and unrelated to decreased MHC class 1 expression.
[The innate immune system in oropharyngeal squamous cell carcinoma : Immune modulation by HPV].
Phenylbutyrate induces cell differentiation and modulates Epstein-Barr virus gene expression in Burkitt's lymphoma cells.
Epidrugs in the Immunotherapy of Cutaneous and Uveal Melanoma.
C-myc activation impairs the NF-kappaB and the interferon response: implications for the pathogenesis of Burkitt's lymphoma.
Therapeutic targeting of the BCR-associated protein CD79b in a TCR-based approach is hampered by aberrant expression of CD79b.
Results of a randomized, double-blind phase II clinical trial of NY-ESO-1 vaccine with ISCOMATRIX adjuvant versus ISCOMATRIX alone in participants with high-risk resected melanoma.
LncRNA USP30-AS1 promotes the survival of acute myeloid leukemia cells by cis-regulating USP30 and ANKRD13A.
Establishment and characterization of HROC69 - a Crohn's related colonic carcinoma cell line and its matched patient-derived xenograft.
Endoplasmic Reticulum Aminopeptidase 2, a common immunological link to adverse pregnancy outcomes and cancer clearance?
Enhancement of human melanoma antigen expression by IFN-beta.
The role of endoplasmic reticulum aminopeptidase 2 in modulating immune detection of choriocarcinoma.
The superantigens SpeC and TSST-1 specifically activate TRBV12-3/12-4(+) memory T cells.
Immunobiological and experimental aspects of malignant astrocytoma.
Deficiency of activated STAT1 in head and neck cancer cells mediates TAP1-dependent escape from cytotoxic T lymphocytes.
High class II-associated invariant chain peptide expression on residual leukemic cells is associated with increased relapse risk in acute myeloid leukemia.
Class II-associated invariant chain peptide as predictive immune marker in minimal residual disease in acute myeloid leukemia.

Table A2-13, Cluster 12

Cluster 12 focuses on immune escape of the SARS-CoV-2 Omicron variant (510)
Insight into SARS-CoV-2 Omicron variant immune escape possibility and variant independent potential therapeutic opportunities.
Humoral and Cellular Immune Responses of COVID-19 vaccines against SARS-Cov-2 Omicron variant: a systemic review.

Role of previous infection with SARS-CoV-2 in protecting against omicron reinfections and severe complications of COVID-19 compared to pre-omicron variants: a systematic review.
Evolution of the SARS-CoV-2 omicron variants BA.1 to BA.5: Implications for immune escape and transmission.
Structural Analysis of the SARS-CoV-2 Omicron Variant Proteins.
An omicron-based vaccine booster elicits potent neutralizing antibodies against emerging SARS-CoV-2 variants in adults.
SARS-CoV-2 Omicron variant: recent progress and future perspectives.
Considerable escape of SARS-CoV-2 Omicron to antibody neutralization.
The emergence and epidemic characteristics of the highly mutated SARS-CoV-2 Omicron variant.
Emerging SARS-CoV-2 variants: Why, how, and what's next?
Sensitivity to Vaccines, Therapeutic Antibodies, and Viral Entry Inhibitors and Advances To Counter the SARS-CoV-2 Omicron Variant.
SARS-CoV-2 bivalent mRNA vaccine with broad protection against variants of concern.
A Heterologous Challenge Rescues the Attenuated Immunogenicity of SARS-CoV-2 Omicron BA.1 Variant in Syrian Hamster Model.
SARS-CoV-2 Omicron variant (B.1.1.529): A concern with immune escape.
Omicron variant (B.1.1.529) and its sublineages: What do we know so far amid the emergence of recombinant variants of SARS-CoV-2?
Characterization of the novel SARS-CoV-2 Omicron (B.1.1.529) variant of concern and its global perspective.
COVID-19: Omicron - the latest, the least virulent, but probably not the last variant of concern of SARS-CoV-2.
Challenges of SARS-CoV-2 Omicron Variant and appropriate countermeasures.
Severity of the Omicron SARS-CoV-2 variant compared with the previous lineages: A systematic review.
SARS-CoV-2 Omicron variant: Characteristics and prevention.
SARS-CoV-2 Omicron Variant of Concern: Everything You Wanted to Know about Omicron but Were Afraid to Ask.
Comprehensive structural analysis reveals broad-spectrum neutralizing antibodies against SARS-CoV-2 Omicron variants.
Human serum from SARS-CoV-2-vaccinated and COVID-19 patients shows reduced binding to the RBD of SARS-CoV-2 Omicron variant.
Spike and nsp6 are key determinants of SARS-CoV-2 Omicron BA.1 attenuation.
A Preliminary Genomic Analysis of the Omicron Variants of SARS-CoV-2 in Central India During the third wave of the COVID-19 Pandemic.
Different Neutralization Profiles After Primary SARS-CoV-2 Omicron BA.1 and BA.2 Infections.
SARS-CoV-2 Omicron variant: Immune escape and vaccine development.
Vaccination with the Omicron spike RBD boosts broadly neutralizing antibody levels and confers sustained protection even after acquiring immunity to the original antigen.
Vaccination After SARS-CoV-2 Infection Increased Antibody Avidity Against the Omicron Variant Compared to Vaccination Alone.
"Is Omicron mild"? Testing this narrative with the mutational landscape of its three lineages and response to existing vaccines and therapeutic antibodies.
Omicron extensively but incompletely escapes Pfizer BNT162b2 neutralization.
Is Omicron the end of pandemic or start of a new innings?
The Biological Functions and Clinical Significance of SARS-CoV-2 Variants of Concern.

Divergent SARS-CoV-2 Omicron-reactive T and B cell responses in COVID-19 vaccine recipients.
Determinants of Spike infectivity, processing, and neutralization in SARS-CoV-2 Omicron subvariants BA.1 and BA.2.
Sub-lineages of the SARS-CoV-2 Omicron variants: Characteristics and prevention.
Global emerging Omicron variant of SARS-CoV-2: Impacts, challenges and strategies.
The humoral and cellular immune evasion of SARS-CoV-2 Omicron and sub-lineages.
Genetic diversity and evolutionary dynamics of the Omicron variant of SARS-CoV-2 in Morocco.
Immune evasion of neutralizing antibodies by SARS-CoV-2 Omicron.
Omicron (B.1.1.529) - variant of concern - molecular profile and epidemiology: a mini review.
Advances in the Omicron variant development.
Mutational Pattern, Impacts and Potential Preventive Strategies of Omicron SARS-CoV-2 Variant Infection.
Genome Evolution and Early Introductions of the SARS-CoV-2 Omicron Variant in Mexico.
Duration of COVID-19 PCR positivity for Omicron vs earlier variants.
Neutralization and Stability of SARS-CoV-2 Omicron Variant.
Drastic decline in sera neutralization against SARS-CoV-2 Omicron variant in Wuhan COVID-19 convalescents.
SARS-CoV-2 Omicron-B.1.1.529 leads to widespread escape from neutralizing antibody responses.
Covid-19 variants: Impact on transmissibility and virulence.
Role of spike in the pathogenic and antigenic behavior of SARS-CoV-2 BA.1 Omicron.
Characterization of Entry Pathways, Species-Specific Angiotensin-Converting Enzyme 2 Residues Determining Entry, and Antibody Neutralization Evasion of Omicron BA.1, BA.1.1, BA.2, and BA.3 Variants.
Live-virus neutralization of the omicron variant in children and adults 14 months after SARS-CoV-2 wild-type infection.
Omicron Genetic and Clinical Peculiarities That May Overturn SARS-CoV-2 Pandemic: A Literature Review.
Activity of convalescent and vaccine serum against SARS-CoV-2 Omicron.
Dynamics of SARS-CoV-2 VOC Neutralization and Novel mAb Reveal Protection against Omicron.
Molecular basis of receptor binding and antibody neutralization of Omicron.
Broadly neutralizing antibodies overcome SARS-CoV-2 Omicron antigenic shift.
Effectiveness of mRNA-1273 against SARS-CoV-2 Omicron and Delta variants.
Outpatient treatment options to address the SARS-CoV-2 variant Omicron.
Omicron: What Makes the Latest SARS-CoV-2 Variant of Concern So Concerning?
The importance of booster vaccination in the context of Omicron wave.
SARS-CoV-2 Omicron Variant Genomic Sequences and Their Epidemiological Correlates Regarding the End of the Pandemic: In Silico Analysis.
Increased risk of infection with SARS-CoV-2 Omicron BA.1 compared with Delta in vaccinated and previously infected individuals, the Netherlands, 22 November 2021 to 19 January 2022.
The relative prevalence of the Omicron variant within SARS-CoV-2 infected cohorts in different countries: A systematic review.
Imprinted antibody responses against SARS-CoV-2 Omicron sublineages.
The significant immune escape of pseudotyped SARS-CoV-2 variant Omicron.
In Silico Genome Analysis Reveals the Evolution and Potential Impact of SARS-CoV-2 Omicron Structural Changes on Host Immune Evasion and Antiviral Therapeutics.
SARS-CoV-2 Omicron Variant Genomic and Phylogenetic Analysis in Iraqi Kurdistan Region.

Household transmission of the SARS-CoV-2 Omicron variant in Denmark.
Global challenge with the SARS-CoV-2 omicron BA.2 (B.1.1.529.2) subvariant: Should we be concerned?
Neutralization capacity of antibodies elicited through homologous or heterologous infection or vaccination against SARS-CoV-2 VOCs.
Baseline Sequencing Surveillance of Public Clinical Testing, Hospitals, and Community Wastewater Reveals Rapid Emergence of SARS-CoV-2 Omicron Variant of Concern in Arizona, USA.
Intradermal administration of DNA vaccine targeting Omicron SARS-CoV-2 via pyro-drive jet injector provides the prolonged neutralizing antibody production via germinal center reaction.
Assessing COVID-19 vaccine effectiveness against Omicron subvariants: Report from a meeting of the World Health Organization.
Analysis of mRNA vaccination-elicited RBD-specific memory B cells reveals strong but incomplete immune escape of the SARS-CoV-2 Omicron variant.
Sequence analysis of the emerging SARS-CoV-2 variant Omicron in South Africa.
Omicron-specific mRNA vaccine induced cross-protective immunity against ancestral SARS-CoV-2 infection with low neutralizing antibodies.
Omicron infection following vaccination enhances a broad spectrum of immune responses dependent on infection history.
SARS-CoV-2 vaccine breakthrough infections (VBI) by Omicron variant (B.1.1.529) and consequences in structural and functional impact.
Structural diversity of the SARS-CoV-2 Omicron spike.
Omicron: a drug developer's perspective.
Omicron Variant (B.1.1.529): Infectivity, Vaccine Breakthrough, and Antibody Resistance.
Structures of Omicron spike complexes and implications for neutralizing antibody development.
SARS-CoV-2 Omicron spike mediated immune escape and tropism shift.
The Omicron variant is highly resistant against antibody-mediated neutralization: Implications for control of the COVID-19 pandemic.
Vaccine effectiveness against hospitalization among adolescent and pediatric SARS-CoV-2 cases between May 2021 and January 2022 in Ontario, Canada: A retrospective cohort study.
Clinical virology and effect of Covid-19 vaccination and monoclonal antibodies against highly infectious SARS-CoV-2 omicron sub variant BF.7 (BA.5.2.1.7): A systematic review.
A Detailed Overview of SARS-CoV-2 Omicron: Its Sub-Variants, Mutations and Pathophysiology, Clinical Characteristics, Immunological Landscape, Immune Escape, and Therapies.
Diminished neutralization responses towards SARS-CoV-2 Omicron VoC after mRNA or vector-based COVID-19 vaccinations.
Omicron infection-associated T- and B-cell immunity in antigen-naïve and triple-COVID-19-vaccinated individuals.
Cryo-EM structure of a SARS-CoV-2 omicron spike protein ectodomain.
Increased Receptor Affinity and Reduced Recognition by Specific Antibodies Contribute to Immune Escape of SARS-CoV-2 Variant Omicron.
Origin, virological features, immune evasion and intervention of SARS-CoV-2 Omicron sublineages.
Structural evolution of SARS-CoV-2 omicron in human receptor recognition.
SARS-CoV-2 Omicron variant replication in human bronchus and lung ex vivo.
Atlas of currently available human neutralizing antibodies against SARS-CoV-2 and escape by Omicron sub-variants BA.1/BA.1.1/BA.2/BA.3.
Neutralization of Omicron sublineages and Deltacron SARS-CoV-2 by three doses of BNT162b2 vaccine or BA.1 infection.

SARS-CoV-2 omicron BA.1.1 is highly resistant to antibody neutralization of convalescent serum from the origin strain.
The Delta and Omicron Variants of SARS-CoV-2: What We Know So Far.
mRNA booster immunization elicits potent neutralizing serum activity against the SARS-CoV-2 Omicron variant.
Rapidly shifting immunologic landscape and severity of SARS-CoV-2 in the Omicron era in South Africa.
BA.2 and BA.5 omicron differ immunologically from both BA.1 omicron and pre-omicron variants.
Comparative Pharmacological Efficacy of COVID-19 Vaccines against the Variants of Concerns (VOCs) of SARS-CoV-2: Recent Clinical Studies on Booster Dose.
Comparison of the replication and neutralization of different SARS-CoV-2 Omicron subvariants in vitro.
Characteristic analysis of Omicron-included SARS-CoV-2 variants of concern.
Potent antibodies against immune invasive SARS-CoV-2 Omicron subvariants.
Parallel profiling of antigenicity alteration and immune escape of SARS-CoV-2 Omicron and other variants.
SARS-CoV-2 Omicron variant shows less efficient replication and fusion activity when compared with Delta variant in TMPRSS2-expressed cells.
Omicron (B.1.1.529): Infectivity, vaccine breakthrough, and antibody resistance.
The Increased Amyloidogenicity of Spike RBD and pH-Dependent Binding to ACE2 May Contribute to the Transmissibility and Pathogenic Properties of SARS-CoV-2 Omicron as Suggested by In Silico Study.
Omicron, the great escape artist.
Analyzing the interaction of human ACE2 and RBD of spike protein of SARS-CoV-2 in perspective of Omicron variant.
Evolution of Immune Evasion and Host Range Expansion by the SARS-CoV-2 B.1.1.529 (Omicron) Variant.
Immune Escape Associated with RBD Omicron Mutations and SARS-CoV-2 Evolution Dynamics.
The Impact of Evolving SARS-CoV-2 Mutations and Variants on COVID-19 Vaccines.
Vaccination-infection interval determines cross-neutralization potency to SARS-CoV-2 Omicron after breakthrough infection by other variants.
Omicron-specific mRNA vaccination alone and as a heterologous booster against SARS-CoV-2.
Omicron escapes the majority of existing SARS-CoV-2 neutralizing antibodies.
Dynamics of competing SARS-CoV-2 variants during the Omicron epidemic in England.
A Systematic Review on the Emergence of Omicron Variant and Recent Advancement in Therapies.
Immune Evasion of SARS-CoV-2 Omicron Subvariants.
A tabulated summary of the evidence on humoral and cellular responses to the SARS-CoV-2 Omicron VOC, as well as vaccine efficacy against this variant.
SARS-CoV-2 Omicron is an immune escape variant with an altered cell entry pathway.
Efficacy of the neutralizing antibodies after the booster dose on SARS-CoV-2 Omicron variant and a two-year longitudinal antibody study on Wild Type convalescents.
Cross-reactivity of eight SARS-CoV-2 variants rationally predicts immunogenicity clustering in sarbecoviruses.
Effects of boosted mRNA and adenoviral-vectored vaccines on immune responses to omicron BA.1 and BA.2 following the heterologous CoronaVac/AZD1222 vaccination.
Limited Neutralization of Omicron by Antibodies from the BNT162b2 Vaccination against SARS-CoV-2.
Synthetic multiantigen MVA vaccine COH04S1 and variant-specific derivatives protect Syrian hamsters from SARS-CoV-2 Omicron subvariants.

Antibody escape and cryptic cross-domain stabilization in the SARS-CoV-2 Omicron spike protein.
A bias of Asparagine to Lysine mutations in SARS-CoV-2 outside the receptor binding domain affects protein flexibility.
More effective vaccines and oral antivirals: Keys for the battle against Omicron.
Epistasis at the SARS-CoV-2 Receptor-Binding Domain Interface and the Propitiously Boring Implications for Vaccine Escape.
SARS-CoV-2 Omicron variant: Antibody evasion and cryo-EM structure of spike protein-ACE2 complex.
Omicron: Call for updated vaccines.
Omicron variant of SARS-CoV-2 imposes a new challenge for the global public health.
Neutralizing antibody activity against 21 SARS-CoV-2 variants in older adults vaccinated with BNT162b2.
Neutralization of SARS-CoV-2 Omicron sub-lineages BA.1, BA.1.1, and BA.2.
Omicron-specific mRNA vaccination alone and as a heterologous booster against SARS-CoV-2.
Omicron variant showed lower neutralizing sensitivity than other SARS-CoV-2 variants to immune sera elicited by vaccines after boost.
The emergence of SARS-CoV-2 Omicron subvariants: current situation and future trends.
Increased household transmission and immune escape of the SARS-CoV-2 Omicron compared to Delta variants.
Broadly neutralizing antibodies overcome SARS-CoV-2 Omicron antigenic shift.
Progressive loss of conserved spike protein neutralizing antibody sites in Omicron sublineages is balanced by preserved T cell immunity.
Booster dose of BNT162b2 after two doses of CoronaVac improves neutralization of SARS-CoV-2 Omicron variant.
The SARS-CoV-2 Delta-Omicron Recombinant Lineage (XD) Exhibits Immune-Escape Properties Similar to the Omicron (BA.1) Variant.
Spike protein of SARS-CoV-2 Omicron variant: An in-silico study evaluating spike interactions and immune evasion.
3Sb5 antibody potentially neutralizes SARS-CoV-2 Omicron by disrupting the N-glycan switch via a conserved spike epitope.
A booster dose of Delta × Omicron hybrid mRNA vaccine produced broadly neutralizing antibody against Omicron and other SARS-CoV-2 variants.
Omicron: A Heavily Mutated SARS-CoV-2 Variant Exhibits Stronger Binding to ACE2 and Potently Escapes Approved COVID-19 Therapeutic Antibodies.
A recombinant spike-XBB.1.5 protein vaccine induces broad-spectrum immune responses against XBB.1.5-included Omicron variants of SARS-CoV-2.
Pulmonary lesions following inoculation with the SARS-CoV-2 Omicron BA.1 (B.1.1.529) variant in Syrian golden hamsters.
Neutralizing antibodies against the SARS-CoV-2 Delta and Omicron variants following heterologous CoronaVac plus BNT162b2 booster vaccination.
Rapid epidemic expansion of the SARS-CoV-2 Omicron variant in southern Africa.
SARS-CoV-2 Omicron Variant is Expected to Retain Most of the Spike Protein Specific Dominant T-Cell Epitopes Presented by COVID-19 Vaccines - Worldwide, 2021.
Steep Decline in Binding Capability of SARS-CoV-2 Omicron Variant (B.1.1.529) RBD to the Antibodies in Early COVID-19 Convalescent Sera and Inactivated Vaccine Sera.
An RBD bispecific antibody effectively neutralizes a SARS-CoV-2 Omicron variant.
Development of New SNP Genotyping Assays to Discriminate the Omicron Variant of SARS-CoV-2.
Comparative analysis of SARS-CoV-2 Omicron BA.2.12.1 and BA.5.2 variants.

Investigating SARS-CoV-2 breakthrough infections per variant and vaccine type.
Epidemiologic Characteristics of SARS-CoV-2 Omicron BA.5.1.3 Variant and the Protection Provided By Inactivated Vaccination.
Antibody Responses to the SARS-CoV-2 Ancestral Strain and Omicron Variants in Moderna mRNA-1273 Vaccinated Active-Duty US Navy Sailors and Marines.
SARS-CoV-2 Omicron variants: burden of disease, impact on vaccine effectiveness and need for variant-adapted vaccines.
Phylogeny and evolution of SARS-CoV-2 during Delta and Omicron variant waves in India.
Significant neutralizing escapes of Omicron and its sublineages in SARS-CoV-2-infected individuals vaccinated with inactivated vaccines.
BA.2.12.1, BA.4 and BA.5 escape antibodies elicited by Omicron infection.
Omicron variant as nature's solution to the COVID-19 pandemic.
Progress of the COVID-19: Persistence, Effectiveness, and Immune Escape of the Neutralizing Antibody in Convalescent Serum.
The outbreak of SARS-CoV-2 Omicron lineages, immune escape, and vaccine effectivity.
Striking antibody evasion manifested by the Omicron variant of SARS-CoV-2.
Serum Neutralization of SARS-CoV-2 Omicron BA.1 and BA.2 after BNT162b2 Booster Vaccination.
Omicron BA.2 lineage predominance in severe acute respiratory syndrome coronavirus 2 positive cases during the third wave in North India.
Omicron-B.1.1.529 leads to widespread escape from neutralizing antibody responses.
Prevalence of symptoms, comorbidities, and reinfections in individuals infected with Wild-Type SARS-CoV-2, Delta, or Omicron variants: a comparative study in western Mexico.
Protective antibodies and T cell responses to Omicron variant after the booster dose of BNT162b2 vaccine.
Omicron: the highly mutational COVID-19 variant with immune escape.
[Progress in research of 2019-nCoV Omicron variant].
Durability of neutralization against Omicron subvariants after vaccination and breakthrough infection.
Potent monoclonal antibodies neutralize Omicron sublineages and other SARS-CoV-2 variants.
The influence of new SARS-CoV-2 variant Omicron (B.1.1.529) on vaccine efficacy, its correlation to Delta variants: A computational approach.
Neutralizing SARS-CoV-2 Spike Antibodies against Omicron in Paired Samples after Two or Three Doses of mRNA Vaccine.
Insights on the mutational landscape of the SARS-CoV-2 Omicron variant receptor-binding domain.
Reduced neutralization against Delta, Gamma, Mu, and Omicron BA.1 variants of SARS-CoV-2 from previous non-Omicron infection.
An Early SARS-CoV-2 Omicron Outbreak in a Dormitory in Saint Petersburg, Russia.
Importation of SARS-CoV-2 Omicron variant in Beijing, China.
Antibodies induced by an ancestral SARS-CoV-2 strain that cross-neutralize variants from Alpha to Omicron BA.1.
Low SARS-CoV-2 viral load among vaccinated individuals infected with Delta B.1.617.2 and Omicron BA.1.1.529 but not with Omicron BA.1.1 and BA.2 variants.
A self-assembled trimeric protein vaccine induces protective immunity against Omicron variant.
Spherical neutralizing aptamer suppresses SARS-CoV-2 Omicron escape.
Cryo-EM structure of the SARS-CoV-2 Omicron spike.
Comparison of total and neutralizing SARS-CoV-2 spike antibodies against omicron and other variants in paired samples after two or three doses of mRNA vaccine.



The spike gene is a major determinant for the SARS-CoV-2 Omicron-BA.1 phenotype.
Neohesperidin and spike RBD interaction in omicron and its sub-variants: In silico, structural and simulation studies.
Molecular mechanisms responsible for SARS-CoV-2 antibody waning and vaccine escape in Omicron sublineages BA.4 and BA.5.
The rapid replacement of the SARS-CoV-2 Delta variant by Omicron (B.1.1.529) in England.
Chimeric mRNA-based COVID-19 vaccine induces protective immunity against Omicron and Delta variants.
SARS-CoV-2 vaccination of convalescents boosts neutralization capacity against Omicron subvariants BA.1, BA.2 and BA.5 and can be predicted by anti-S antibody concentrations in serological assays.
Does the COVID-19 XBB Omicron subvariant signal the beginning of the end of the pandemic?
Three exposures to the spike protein of SARS-CoV-2 by either infection or vaccination elicit superior neutralizing immunity to all variants of concern.
Anti-human ACE2 antibody neutralizes and inhibits virus production of SARS-CoV-2 variants of concern.
Structural basis of Omicron immune evasion: A comparative computational study.
RETRACTED: Gradual emergence followed by exponential spread of the SARS-CoV-2 Omicron variant in Africa.
Immediate reinfection with Omicron variant after clearance of a previous SARS-CoV-2 infection.
A case of SARS-CoV-2 Omicron reinfection resulting in a significant immunity boost in a paediatric patient affected by B-cell acute lymphoblastic leukemia.
Detection of the Omicron (B.1.1.529) variant of SARS-CoV-2 in aircraft wastewater.
Analysis of anti-SARS-CoV-2 Omicron-neutralizing antibody titers in different vaccinated and unvaccinated convalescent plasma sources.
Cysteamine exerts in vitro antiviral activity against the SARS-CoV-2 Delta and Omicron variants.
Evolution of antibody immunity following Omicron BA.1 breakthrough infection.
Genomic Surveillance for SARS-CoV-2 Variants: Predominance of the Delta (B.1.617.2) and Omicron (B.1.1.529) Variants - United States, June 2021-January 2022.
Differential Cell Line Susceptibility to the SARS-CoV-2 Omicron BA.1.1 Variant of Concern.
Structural and functional characteristics of the SARS-CoV-2 Omicron subvariant BA.2 spike protein.
Structural basis of SARS-CoV-2 Omicron immune evasion and receptor engagement.
The Spread of SARS-CoV-2 Variant Omicron with a Doubling Time of 2.0-3.3 Days Can Be Explained by Immune Evasion.
Long term anti-SARS-CoV-2 antibody kinetics and correlate of protection against Omicron BA.1/BA.2 infection.
Omicron mutations increase interdomain interactions and reduce epitope exposure in the SARS-CoV-2 spike.
Antigenic cartography using sera from sequence-confirmed SARS-CoV-2 variants of concern infections reveals antigenic divergence of Omicron.
The rapid rise of SARS-CoV-2 Omicron subvariants with immune evasion properties: XBB.1.5 and BQ.1.1 subvariants.
Editorial: The XBB.1.5 ('Kraken') Subvariant of Omicron SARS-CoV-2 and its Rapid Global Spread.
Effectiveness of BNT162b2 and Sinovac vaccines against the transmission of SARS-CoV-2 during Omicron-predominance in Hong Kong: A retrospective cohort study of COVID-19 cases.
Structural and functional characteristics of SARS-CoV-2 Omicron subvariant BA.2 spike.
Persistent SARS-COV-2 infection in vaccinated individual with three doses of COVID-19 vaccine.
Developing Pseudovirus-Based Neutralization Assay against Omicron-Included SARS-CoV-2 Variants.

A structural dynamic explanation for observed escape of SARS-CoV-2 BA.2 variant mutation S371L/F.
Do We Really Need Omicron Spike-Based Updated COVID-19 Vaccines? Evidence and Pipeline.
Omicron SARS-CoV-2 variant of concern: A review on its transmissibility, immune evasion, reinfection, and severity.
Omicron SARS-CoV-2 mutations stabilize spike up-RBD conformation and lead to a non-RBM-binding monoclonal antibody escape.
Humoral immune response to authentic circulating severe acute respiratory syndrome coronavirus 2 variants elicited by booster vaccination with distinct receptor-binding domain subunits in mice.
SARS-CoV-2 Omicron BA.4/BA.5 Mutations in Spike Leading to T Cell Escape in Recently Vaccinated Individuals.
Autochthonous Outbreak of SARS-CoV-2 Omicron Variant in Booster-Vaccinated (3 Doses) Healthcare Workers in Southern Italy: Just the Tip of the Iceberg?
Retained avidity despite reduced cross-binding and cross-neutralizing antibody levels to Omicron after SARS-CoV-2 wild-type infection or mRNA double vaccination.
Literature Review of Omicron: A Grim Reality Amidst COVID-19.
Structural Basis for the Enhanced Infectivity and Immune Evasion of Omicron Subvariants.
The Omicron Variant Reinfection Risk among Individuals with a Previous SARS-CoV-2 Infection within One Year in Shanghai, China: A Cross-Sectional Study.
Early Genomic, Epidemiological, and Clinical Description of the SARS-CoV-2 Omicron Variant in Mexico City.
Longitudinal Tracking of Immune Responses in COVID-19 Convalescents Reveals Absence of Neutralization Activity Against Omicron and Staggered Impairment to Other SARS-CoV-2 Variants of Concern.
An Early and Preliminary Assessment of the Clinical Severity of the Emerging SARS-CoV-2 Omicron Variants in Maharashtra, India.
Emergence of SARS-CoV-2 Omicron (B.1.1.529) variant, salient features, high global health concerns and strategies to counter it amid ongoing COVID-19 pandemic.
Evolution of antibody immunity following Omicron BA.1 breakthrough infection.
SARS-CoV-2 Omicron sublineages exhibit distinct antibody escape patterns.
Booster dose of BNT162b2 in a CoronaVac primary vaccination protocol improves neutralization of SARS-CoV-2 Omicron variant.
What do we know about pathological mechanism and pattern of lung injury related to SARS-CoV-2 Omicron variant?
Rapid displacement of SARS-CoV-2 variant Delta by Omicron revealed by allele-specific PCR in wastewater.
VH3-53/66-Class RBD-Specific Human Monoclonal Antibody iB20 Displays Cross-Neutralizing Activity against Emerging SARS-CoV-2 Lineages.
Evasion of vaccine-induced humoral immunity by emerging sub-variants of SARS-CoV-2.
Reduced replication but increased interferon resistance of SARS-CoV-2 Omicron BA.1.
Neutralization Activity against SARS-CoV-2 Variants after Booster Vaccination in Populations without COVID-19: A Meta-Analysis.
Cross-variant protection against SARS-CoV-2 infection in hamsters immunized with monovalent and bivalent inactivated vaccines.
Comparative analysis of the risks of hospitalisation and death associated with SARS-CoV-2 omicron (B.1.1.529) and delta (B.1.617.2) variants in England: a cohort study.
Clinical characteristics and host immunity responses of SARS-CoV-2 Omicron variant BA.2 with deletion of ORF7a, ORF7b and ORF8.

The rapid and efficient strategy for SARS-CoV-2 Omicron transmission control: analysis of outbreaks at the city level.
A Synthetic SARS-CoV-2-Derived T-Cell and B-Cell Peptide Cocktail Elicits Full Protection against Lethal Omicron BA.1 Infection in H11-K18-hACE2 Mice.
Evaluation of immune evasion in SARS-CoV-2 Delta and Omicron variants.
[Progress in research of etiology and epidemiology of 2019-nCoV Omicron variant].
Omicron variants escape the persistent SARS-CoV-2-specific antibody response in 2-year COVID-19 convalescents regardless of vaccination.
Sub-optimal neutralisation of omicron (B.1.1.529) variant by antibodies induced by vaccine alone or SARS-CoV-2 Infection plus vaccine (hybrid immunity) post 6-months.
SARS-CoV-2 evolves to reduce but not abolish neutralizing action.
Structural and functional characterizations of infectivity and immune evasion of SARS-CoV-2 Omicron.
SARS-CoV-2 in relation to global vaccination and booster doses: what is the future of vaccination in the battle against COVID-19?
Sequencing and mutations analysis of the first recorded SARS-CoV-2 Omicron variant during the fourth wave of pandemic in Iraq.
Correlation of Binding and Neutralizing Antibodies against SARS-CoV-2 Omicron Variant in Infection-Naïve and Convalescent BNT162b2 Recipients.
Potent cross-reactive antibodies following Omicron breakthrough in vaccinees.
Emerging Omicron subvariants evade neutralizing immunity elicited by vaccine or BA.1/BA.2 infection.
COVID-19 infection after SARS-CoV-2 mRNA vaccination in Multiple Sclerosis, AQP4-antibody NMOSD and MOGAD patients during the Omicron subvariant BA.1/2 wave in Singapore.
SARS CoV-2 reinfection rate is higher in the Omicron variant than in the Alpha and Delta variants.
Neutralization against Omicron subvariants after BA.5/BF.7 breakthrough infection weakened as virus evolution and aging despite repeated prototype-based vaccination.
A systems approach evaluating the impact of SARS-CoV-2 variant of concern mutations on CD8+ T cell responses.
Co-incidence of BA.1 and BA.2 at the start of Singapore's Omicron wave revealed by Community and University Campus wastewater surveillance.
Introduction, Spread and Impact of the SARS-CoV-2 Omicron Variants BA.1 and BA.2 in Cyprus.
Development of Bivalent mRNA Vaccines against SARS-CoV-2 Variants.
Impact of Reinfection with SARS-CoV-2 Omicron Variants in Previously Infected Hamsters.
Omicron BA.1 Mutations in SARS-CoV-2 Spike Lead to Reduced T-Cell Response in Vaccinated and Convalescent Individuals.
Receptor binding and complex structures of human ACE2 to spike RBD from omicron and delta SARS-CoV-2.
Molecular recognition of SARS-CoV-2 spike protein with three essential partners: exploring possible immune escape mechanisms of viral mutants.
BNT162b2 Protection against the Omicron Variant in Children and Adolescents.
Shared N417-Dependent Epitope on the SARS-CoV-2 Omicron, Beta, and Delta Plus Variants.
Deep Structural Analysis of Myriads of Omicron Sub-Variants Revealed Hotspot for Vaccine Escape Immunity.
A comprehensive review on Covid-19 Omicron (B.1.1.529) variant.
The landscape of antibody binding affinity in SARS-CoV-2 Omicron BA.1 evolution.
Immune Escape Mechanisms of SARS-CoV-2 Delta and Omicron Variants against Two Monoclonal Antibodies That Received Emergency Use Authorization.

Identification of key mutations responsible for the enhancement of receptor-binding affinity and immune escape of SARS-CoV-2 Omicron variant.
A Fourth Dose of COVID-19 Vaccine Does Not Induce Neutralization of the Omicron Variant Among Solid Organ Transplant Recipients With Suboptimal Vaccine Response.
Unique mutations in SARS-CoV-2 Omicron subvariants' non-spike proteins: Potential impacts on viral pathogenesis and host immune evasion.
Rapid evaluation of heterologous chimeric RBD-dimer mRNA vaccine for currently-epidemic Omicron sub-variants as booster shot after inactivated vaccine.
Ending the COVID-19 pandemic: We still have a long way to go.
Analysis of a SARS-CoV-2 convalescent cohort identified a common strategy for escape of vaccine-induced anti-RBD antibodies by Beta and Omicron variants.
Combined Antibodies Evusheld against the SARS-CoV-2 Omicron Variants BA.1.1 and BA.5: Immune Escape Mechanism from Molecular Simulation.
Omicron (B.1.1.529) variant of SARS-CoV-2: Concerns, challenges, and recent updates.
Vaccination and Transmission Risk during the Outbreak of B.1.1.529 (Omicron).
Population immunity to pre-Omicron and Omicron SARS-CoV-2 variants in US states and counties through December 1, 2021.
Immunogenicity of BNT162b2 Vaccination against SARS-CoV-2 Omicron Variant and Attitudes toward a COVID-19 Booster Dose among Healthy Thai Adolescents.
An imported human case with the SARS-CoV-2 Omicron subvariant BA.2.75 in Yunnan Province, China.
Viral load of SARS-CoV-2 Omicron is not high despite its high infectivity.
Epistasis at the SARS-CoV-2 RBD Interface and the Propitiously Boring Implications for Vaccine Escape.
Long-term variations and potency of neutralizing antibodies against Omicron subvariants after CoronaVac-inactivated booster: A 7-month follow-up study.
An updated atlas of antibody evasion by SARS-CoV-2 Omicron sub-variants including BQ.1.1 and XBB.
Coronavirus Disease 19 (COVID-19) Vaccine Effectiveness Against Symptomatic Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) Infection During Delta-Dominant and Omicron-Dominant Periods in Japan: A Multicenter Prospective Case-control Study (Factors Associated with SARS-CoV-2 Infection and the Effectiveness of COVID-19 Vaccines Study).
Structural delineation and computational design of SARS-CoV-2-neutralizing antibodies against Omicron subvariants.
Rational strategies for enhancing mAb binding to SARS-CoV-2 variants through CDR diversification and antibody-escape prediction.
Immunoinformatics and MD-simulation data suggest that Omicron spike epitopes are more interacting to IgG via better MHC recognition than Delta variant.
Effectiveness of a Third Dose of mRNA Vaccines Against COVID-19-Associated Emergency Department and Urgent Care Encounters and Hospitalizations Among Adults During Periods of Delta and Omicron Variant Predominance - VISION Network, 10 States, August 2021-January 2022.
Delta-Omicron recombinant escapes therapeutic antibody neutralization.
Can the New BA.2.75 Sub-variant Lead To One More COVID-19 Wave?
Imprinted SARS-CoV-2 humoral immunity induces convergent Omicron RBD evolution.
The immune evasion ability of Delta variant is comparable to that of Beta variant in South Africa.

Novel pro-and eukaryotic expression plasmid expressing omicron antigens delivered via Salmonella elicited MHC class I and II based protective immunity.
Biophysical Fitness Landscape of the SARS-CoV-2 Delta Variant Receptor Binding Domain.
Resistance of Omicron subvariants BA.2.75.2, BA.4.6 and BQ.1.1 to neutralizing antibodies.
Mutation-induced changes in the receptor-binding interface of the SARS-CoV-2 Delta variant B.1.617.2 and implications for immune evasion.
Circulation of SARS-CoV-2 Omicron sub-lineages revealed by multiplex genotyping RT-qPCR assays for sewage surveillance.
Strain wars 5: Gibbs energies of binding of BA.1 through BA.4 variants of SARS-CoV-2.
Methylene blue, Mycophenolic acid, Posaconazole, and Niclosamide inhibit SARS-CoV-2 Omicron variant BA.1 infection of human airway epithelial organoids.
Impact of SARS-CoV-2 Mu variant on vaccine effectiveness: A comparative genomics study at the peak of the third wave in Bogota, Colombia.
Urticaria after breakthrough Omicron BA.5.1 severe acute respiratory syndrome coronavirus 2 infection in a triple-vaccinated (Pfizer) patient: a case report.
Investigating the role of glycans in Omicron sub-lineages XBB.1.5 and XBB.1.16 binding to host receptor using molecular dynamics and binding free energy calculations.
Neutralization Titers in Vaccinated Patients with SARS-CoV-2 Delta Breakthrough Infections.
Structural Characteristics of Heparin Binding to SARS-CoV-2 Spike Protein RBD of Omicron Sub-Lineages BA.2.12.1, BA.4 and BA.5.
Heterologous boosting with third dose of coronavirus disease recombinant subunit vaccine increases neutralizing antibodies and T cell immunity against different severe acute respiratory syndrome coronavirus 2 variants.
Boosting with Multiple Doses of mRNA Vaccine after Priming with Two Doses of Protein Subunit Vaccine MVC-COV1901 Elicited Robust Humoral and Cellular Immune Responses against Emerging SARS-CoV-2 Variants.
Sera from Breakthrough Infections with SARS-CoV-2 BA.5 or BF.7 Showed Lower Neutralization Activity against XBB.1.5 and CH.1.1.
Enhanced evasion of neutralizing antibody response by Omicron XBB.1.5, CH.1.1, and CA.3.1 variants.
An ecological study on reinfection rates using a large dataset of RT-qPCR tests for SARS-CoV-2 in Santiago of Chile.
Clinical and genomic signatures of SARS-CoV-2 Delta breakthrough infections in New York.
Genetic and Structural Data on the SARS-CoV-2 Omicron BQ.1 Variant Reveal Its Low Potential for Epidemiological Expansion.
Genomic Tracking of SARS-CoV-2 Variants in Myanmar.
A delicate balance between antibody evasion and ACE2 affinity for Omicron BA.2.75.
Emergence of COVID-19 Variants: An Update.
Investigating the structural impact of Omicron RBD mutation on antibody escape and receptor management.
Immune evasion and chronological decrease in titer of neutralizing antibody against SARS-CoV-2 and its variants of concerns in COVID-19 patients.
Clinical and genomic signatures of rising SARS-CoV-2 Delta breakthrough infections in New York.
When the Dust Has Settled: Calculation of Binding Affinities from First Principles for SARS-CoV-2 Variants with Quantitative Accuracy.
Immunodominant antibody responses directed to SARS-CoV-2 hotspot mutation sites and risk of immune escape.
Characterization of the enhanced infectivity and antibody evasion of Omicron BA.2.75.

The allocation of COVID-19 vaccines and antivirals against emerging SARS-CoV-2 variants of concern in East Asia and Pacific region: A modelling study.
Understanding the dynamic relation between wastewater SARS-CoV-2 signal and clinical metrics throughout the pandemic.
Early Detection of SARS-CoV-2 Omicron BA.4 and BA.5 in German Wastewater.
Weighted gene co-expression network-based identification of genetic effect of mRNA vaccination and previous infection on SARS-CoV-2 infection.
Trends in Confirmed COVID-19 Cases in the Korean Military Before and After the Emergence of the Omicron Variant.
Reproduction Number of the Omicron Variant Triples That of the Delta Variant.
A Critical Analysis of the Use of Cilgavimab plus Tixagevimab Monoclonal Antibody Cocktail (Evusheld™) for COVID-19 Prophylaxis and Treatment.
Coarse-Grained Molecular Simulations and Ensemble-Based Mutational Profiling of Protein Stability in the Different Functional Forms of the SARS-CoV-2 Spike Trimers: Balancing Stability and Adaptability in BA.1, BA.2 and BA.2.75 Variants.
Tracking B Cell Memory to SARS-CoV-2 Using Rare Cell Analysis System.
Oral high-dose acetylcysteine: Effective against the Omicron variant of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)?
Omicron Sub-Lineage BA.5 and Recombinant XBB Evasion from Antibody Neutralisation in BNT162b2 Vaccine Recipients.
Early Estimates of Bivalent mRNA Vaccine Effectiveness in Preventing COVID-19-Associated Hospitalization Among Immunocompetent Adults Aged ≥65 Years - IVY Network, 18 States, September 8-November 30, 2022.
Autophagy and evasion of the immune system by SARS-CoV-2. Structural features of the non-structural protein 6 from wild type and Omicron viral strains interacting with a model lipid bilayer.
New Omicron strains may portend big COVID-19 waves.
Neutralizing monoclonal antibodies elicited by mosaic RBD nanoparticles bind conserved sarbecovirus epitopes.
Immunogenicity of BNT162b2 vaccine against the Alpha and Delta variants in immunocompromised patients with systemic inflammatory diseases.
Neutralizing antibody activity against the B.1.617.2 (delta) variant 8 months after two-dose vaccination with BNT162b2 in health care workers.
Bivalent Omicron BA.1-Adapted BNT162b2 Booster in Adults Older than 55 Years.
Rapid transmission and tight bottlenecks constrain the evolution of highly transmissible SARS-CoV-2 variants.
XBB.1.16 Omicron subvariant rise to a variant of interest: Implications for global alertness and preparedness.
Twice evasions of Omicron variants explain the temporal patterns in six Asian and Oceanic countries.
Fourth Dose of BNT162b2 Vaccine for Patients with Autoimmune Rheumatic Diseases in a Nationwide Setting.
Severe breakthrough COVID-19 with a heavily mutated variant in a multiple myeloma patient 10 weeks after vaccination.
Omicron subvariant BA.5 is highly contagious but containable: Successful experience from Macau.
The Biological Properties of the SARS-CoV-2 Cameroon Variant Spike: An Intermediate between the Alpha and Delta Variants.
Adjudicating the logistics of COVID-19 vaccine boosters from a global perspective.

Variable neutralizing antibody responses to 10 SARS-CoV-2 variants in natural infection with wild-type (B.1) virus, Kappa (B.1.617.1), and Delta (B.1.617.2) variants and COVISHIELD vaccine immunization in India: utility of the MSD platform.
Myopericarditis following both BNT162b2 and NVX-CoV2373.
Rapid investigation of BA.4/BA.5 cases in France.
A noncompeting pair of human neutralizing antibodies block COVID-19 virus binding to its receptor ACE2.
COVID-19 vaccine breakthrough infections.
BNT162b2 vaccine effectiveness was marginally affected by the SARS-CoV-2 beta variant in fully vaccinated individuals.
Elucidating causes of COVID-19 infection and related deaths after vaccination.
Neutralizing Antibody Activity Against the B.1.617.2 (delta) Variant Before and After a Third BNT162b2 Vaccine Dose in Hemodialysis Patients.
Waning Immunity Against XBB.1.5 Following Bivalent mRNA Boosters.
Mobility was a significant determinant of reported COVID-19 incidence during the Omicron Surge in the most populous U.S. Counties.
Fully understanding the efficacy profile of the COVID-19 vaccination and its associated factors in multiple real-world settings.
New Omicron begins to take over, despite late start.
Post-Vaccination Coronavirus Disease 2019: A Case-Control Study and Genomic Analysis of 119 Breakthrough Infections in Partially Vaccinated Individuals.
Nano-Enabled COVID-19 Vaccines: Meeting the Challenges of Durable Antibody Plus Cellular Immunity and Immune Escape.
A review of the safety and efficacy of current COVID-19 vaccines.
How Immune-Evasive Omicron Offspring and a Lack of Mitigation Measures Could Shape a COVID-19 Winter Wave.
After Omicron, some scientists foresee 'a period of quiet'.
Human airway and nasal organoids reveal escalating replicative fitness of SARS-CoV-2 emerging variants.
Humoral immunity and transcriptome differences of COVID-19 inactivated vaccine and protein subunit vaccine as third booster dose in human.
Modelling the effects of social distancing, antiviral therapy, and booster shots on mitigating Omicron spread.
Evolution of SARS-CoV-2-specific CD4(+) T cell epitopes.
A spatial vaccination strategy to reduce the risk of vaccine-resistant variants.
The Mechanism of bnAb Production and Its Application in Mutable Virus Broad-Spectrum Vaccines: Inspiration from HIV-1 Broad Neutralization Research.
Vaccination strategies when vaccines are scarce: on conflicts between reducing the burden and avoiding the evolution of escape mutants.
[Progress in research of epidemiology of 2019-nCoV reinfection].
Wastewater-based modeling, reconstruction, and prediction for COVID-19 outbreaks in Hungary caused by highly immune evasive variants.
Epidemiological and evolutionary considerations of SARS-CoV-2 vaccine dosing regimes.
The assessment of Twitter discourse on the new COVID-19 variant, XBB.1.5, through social network analysis.
Real-world effectiveness of Yindan Jiedu granules-based treatment on patients infected with the SARS-CoV-2 Omicron variants BA.2 combined with high-risk factors: A cohort study.

Protective effect conferred by prior infection and vaccination on COVID-19 in a healthcare worker cohort in South India.
Management experience of a designated hospital for children with coronavirus disease 2019.
Dosing interval strategies for two-dose COVID-19 vaccination in 13 middle-income countries of Europe: Health impact modelling and benefit-risk analysis.
Protection Conferred by Delta and BA.1/BA.2 Infection Against BA.4/BA.5 Infection and Hospitalization: A Retrospective Cohort Study.
The role of booster vaccination and ongoing viral evolution in seasonal circulation of SARS-CoV-2.
Allogeneic hematopoietic stem cell transplantation in the COVID-19 era.
The XBB.1.5 slightly increase the binding affinity for host receptor ACE2 and exhibit strongest immune escaping features: molecular modeling and free energy calculation.
Time trends in social contacts of individuals according to comorbidity and vaccination status, before and during the COVID-19 pandemic.
Vaccination deep into a pandemic wave potential mechanisms for a "third wave" and the impact of vaccination.

Table A2-14, Cluster 13

Cluster 13 focuses on hepatocellular carcinoma, emphasizing the potential role of immunotherapies in countering tumor microenvironment-mediated immune system evasion (222)
From bench to bed: the tumor immune microenvironment and current immunotherapeutic strategies for hepatocellular carcinoma.
Current immunotherapeutic strategies in hepatocellular carcinoma: recent advances and future directions.
Immunotherapy for hepatocellular carcinoma.
Immunotherapy of hepatocellular carcinoma.
Immunotherapeutic Approaches for Treating Hepatocellular Carcinoma.
Tumor microenvironment-mediated immune evasion in hepatocellular carcinoma.
New Therapeutics for HCC: Does Tumor Immune Microenvironment Matter?
Targeting Immune Cells in the Tumor Microenvironment of HCC: New Opportunities and Challenges.
Immunological Microenvironment of Hepatocellular Carcinoma and Its Clinical Implication.
Immunotherapy in hepatocellular carcinoma: Primed to make a difference?
[Research progress of AFP in the diagnosis and therapy of hepatocellular carcinoma].
Recent progress in understanding mitokines as diagnostic and therapeutic targets in hepatocellular carcinoma.
The Mechanism and Clinical Significance of Circular RNAs in Hepatocellular Carcinoma.
Epigenetic mechanisms of liver tumor resistance to immunotherapy.
Hepatocellular Carcinoma Immune Landscape and the Potential of Immunotherapies.
PIGU promotes hepatocellular carcinoma progression through activating NF- $\kappa$ B pathway and increasing immune escape.
Immune responses in hepatocellular carcinoma.
Immunotherapy for Hepatocellular Carcinoma: Current Status and Future Prospects.
Integrated bioinformatics analyses of key genes involved in hepatocellular carcinoma immunosuppression.
Elevated MPP6 expression correlates with an unfavorable prognosis, angiogenesis and immune evasion in hepatocellular carcinoma.



Immunological landscape and immunotherapy of hepatocellular carcinoma.
SEPHS1 promotes SMAD2/3/4 expression and hepatocellular carcinoma cells invasion.
Circular RNA circTRIM33-12 acts as the sponge of MicroRNA-191 to suppress hepatocellular carcinoma progression.
HERC2 promotes inflammation-driven cancer stemness and immune evasion in hepatocellular carcinoma by activating STAT3 pathway.
Six-Transmembrane Epithelial Antigen of Prostate 4: An Indicator of Prognosis and Tumor Immunity in Hepatocellular Carcinoma.
Dual role of ANGPTL8 in promoting tumor cell proliferation and immune escape during hepatocarcinogenesis.
Annexin A2 promotion of hepatocellular carcinoma tumorigenesis via the immune microenvironment.
Immunomodulatory TGF- $\beta$ Signaling in Hepatocellular Carcinoma.
Immunotherapy for Hepatocellular Carcinoma: New Prospects for the Cancer Therapy.
Glycolysis-related gene dihydrolipoamide acetyltransferase promotes poor prognosis in hepatocellular carcinoma through the Wnt/ $\beta$ -catenin and PI3K/Akt signaling pathways.
Role of Noncoding RNAs in the Tumor Immune Microenvironment of Hepatocellular Carcinoma.
Highly expressed carbohydrate sulfotransferase 11 correlates with unfavorable prognosis and immune evasion of hepatocellular carcinoma.
Immune checkpoint blockade in hepatocellular carcinoma: current progress and future directions.
Long non-coding RNA LINC00992 promotes hepatocellular carcinoma cell proliferation, metastasis, and invasiveness by downregulating MicroRNA miR-361-5p expression to increase levels of the transcription factor twist1.
Xanthine dehydrogenase as a prognostic biomarker related to tumor immunology in hepatocellular carcinoma.
Elevated FBXL6 expression in hepatocytes activates VRK2-transketolase-ROS-mTOR-mediated immune evasion and liver cancer metastasis in mice.
The Role of Immunotherapy in a Tolerogenic Environment: Current and Future Perspectives for Hepatocellular Carcinoma.
Radiation therapy in the era of immune treatment for hepatocellular carcinoma.
Immunotherapy and the Combination with Targeted Therapies for Advanced Hepatocellular Carcinoma.
TKIs in combination with immunotherapy for hepatocellular carcinoma.
Immunotherapy in hepatocellular carcinoma.
Mitochondrial TSPO Promotes Hepatocellular Carcinoma Progression through Ferroptosis Inhibition and Immune Evasion.
Combinatorial targeting of immune checkpoints and epigenetic regulators for hepatocellular carcinoma therapy.
An Integrated Analysis of the Identified PRPF19 as an Onco-immunological Biomarker Encompassing the Tumor Microenvironment, Disease Progression, and Prognoses in Hepatocellular Carcinoma.
A comprehensive study based on exosome-related immunosuppression genes and tumor microenvironment in hepatocellular carcinoma.
Insights into an Immunotherapeutic Approach to Combat Multidrug Resistance in Hepatocellular Carcinoma.
Immunotherapies for hepatocellular carcinoma.
Novel risk model based on angiogenesis-related lncRNAs for prognosis prediction of hepatocellular carcinoma.

The unique characteristic in peripheral immune cells in patients with advanced hepatocellular carcinoma.
Loss of SIRT5 promotes bile acid-induced immunosuppressive microenvironment and hepatocarcinogenesis.
Cooperation Between Distinct Cancer Driver Genes Underlies Intertumor Heterogeneity in Hepatocellular Carcinoma.
Functional long non-coding RNAs in hepatocellular carcinoma.
Upregulation of B7-H1 expression is associated with macrophage infiltration in hepatocellular carcinomas.
Epigenetic modification-related mechanisms of hepatocellular carcinoma resistance to immune checkpoint inhibition.
Development of Prognostic Evaluation Model to Predict the Overall Survival and Early Recurrence of Hepatocellular Carcinoma.
Indoleamine 2,3-dioxygenase: As a potential prognostic marker and immunotherapeutic target for hepatocellular carcinoma.
Immunotherapy for targeting cancer stem cells in hepatocellular carcinoma.
Sorafenib combined with STAT3 knockdown triggers ER stress-induced HCC apoptosis and cGAS-STING-mediated anti-tumor immunity.
Liver immunotolerance and hepatocellular carcinoma: Patho-physiological mechanisms and therapeutic perspectives.
Significant correlation between HSPA4 and prognosis and immune regulation in hepatocellular carcinoma.
The yin and yang of evasion and immune activation in HCC.
The clinical and prognostic significance of CD14(+)HLA-DR(-/low) myeloid-derived suppressor cells in hepatocellular carcinoma patients receiving radiotherapy.
HSP70-eIF4G Interaction Promotes Protein Synthesis and Cell Proliferation in Hepatocellular Carcinoma.
Large scale identification of human hepatocellular carcinoma-associated antigens by autoantibodies.
Recent advances in immunotherapy for hepatocellular carcinoma.
Emerging roles and the regulation of aerobic glycolysis in hepatocellular carcinoma.
DCK is a promising prognostic biomarker and correlated with immune infiltrates in hepatocellular carcinoma.
Clinical Significance of Glycolytic Metabolic Activity in Hepatocellular Carcinoma.
Prognostic stratification based on m(5)C regulators acts as a novel biomarker for immunotherapy in hepatocellular carcinoma.
HIF inhibitor 32-134D eradicates murine hepatocellular carcinoma in combination with anti-PD1 therapy.
ERRF1 induces apoptosis of hepatocellular carcinoma cells in response to tryptophan deficiency.
Immunogenic landscape and risk score prediction based on unfolded protein response (UPR)-related molecular subtypes in hepatocellular carcinoma.
Prognostic and Immunological Significance of FUNDC1 in Hepatocellular Carcinoma: A Study on TCGA Mining.
Novel hypoxia features with appealing implications in discriminating the prognosis, immune escape and drug responses of 947 hepatocellular carcinoma patients.
Immune checkpoint inhibitors for hepatocellular carcinoma.
Tumor-Intrinsic Mechanisms Regulating Immune Exclusion in Liver Cancers.
Immunomodulation in hepatocellular cancer.

SLC7A2 deficiency promotes hepatocellular carcinoma progression by enhancing recruitment of myeloid-derived suppressors cells.
Next Generation of Preclinical Liver Cancer Models.
Evaluation of the possible association of PDCD-1 and LAG3 gene polymorphisms with hepatocellular carcinoma risk.
MicroRNAs: Regulators of immunological reactions in hepatocellular carcinoma.
Comprehensive analysis of the tumor-promoting effect and immune infiltration correlation MAZ from pan-cancer to hepatocellular carcinoma.
MAT1A Suppression by the CTBP1/HDAC1/HDAC2 Transcriptional Complex Induces Immune Escape and Reduces Ferroptosis in Hepatocellular Carcinoma.
LOXL2 serves as a prognostic biomarker for hepatocellular carcinoma by mediating immune infiltration and vasculogenic mimicry.
Clinico-histological and molecular features of hepatocellular carcinoma from nonalcoholic fatty liver disease.
M7G methylated core genes (METTL1 and WDR4) and associated RNA risk signatures are associated with prognosis and immune escape in HCC.
Construction and Validation of an Epigenetic Regulator Signature as A Novel Biomarker For Prognosis, Immunotherapy, And Chemotherapy In Hepatocellular Carcinoma.
Construction of a novel immune-related lncRNA signature and its potential to predict the immune status of patients with hepatocellular carcinoma.
4Mu Decreases CD47 Expression on Hepatic Cancer Stem Cells and Primes a Potent Antitumor T Cell Response Induced by Interleukin-12.
Immunotherapy for hepatocellular carcinoma.
A new perspective on the immune escape mechanism in HCC: onco-foetal reprogramming.
Non-coding RNA-associated competitive endogenous RNA regulatory networks: Novel diagnostic and therapeutic opportunities for hepatocellular carcinoma.
Current Strategies for the Treatment of Hepatocellular Carcinoma by Modulating the Tumor Microenvironment via Nano-Delivery Systems: A Review.
Signature construction and molecular subtype identification based on immune-related genes for better prediction of prognosis in hepatocellular carcinoma.
Role of Virus-Related Chronic Inflammation and Mechanisms of Cancer Immune-Suppression in Pathogenesis and Progression of Hepatocellular Carcinoma.
Histone deacetylase 2 regulates STAT1-dependent upregulation of atypical chemokine receptor 3 to induce M2 macrophage migration and immune escape in hepatocellular carcinoma.
Interferon Gamma-Induced Interferon Regulatory Factor 1 Activates Transcription of HHLA2 and Induces Immune Escape of Hepatocellular Carcinoma Cells.
Prospect of the use of checkpoint inhibitors in hepatocellular cancer treatments.
Hepatocellular Carcinoma: Current Therapeutic Algorithm for Localized and Advanced Disease.
The Significance of Circulating Tumor Cells in Patients with Hepatocellular Carcinoma: Real-Time Monitoring and Moving Targets for Cancer Therapy.
Identification of m6A- and ferroptosis-related lncRNA signature for predicting immune efficacy in hepatocellular carcinoma.
Immunotherapy in hepatocellular carcinoma: the complex interface between inflammation, fibrosis, and the immune response.
High numbers of myeloid derived suppressor cells in peripheral blood and ascitic fluid of cirrhotic and HCC patients.
Prognostic Implication of a Novel Metabolism-Related Gene Signature in Hepatocellular Carcinoma.

Hepatocellular carcinoma: Prospects for natural killer cell immunotherapy.
Crosstalk between 5-methylcytosine and N(6)-methyladenosine machinery defines disease progression, therapeutic response and pharmacogenomic landscape in hepatocellular carcinoma.
Angiogenic inflammation and formation of necrosis in the tumor microenvironment influence patient survival after radical surgery for de novo hepatocellular carcinoma in non-cirrhosis.
Biology, Significance and Immune Signaling of Mucin 1 in Hepatocellular Carcinoma.
CCL15 Recruits Suppressive Monocytes to Facilitate Immune Escape and Disease Progression in Hepatocellular Carcinoma.
Protein glycosylation alterations in hepatocellular carcinoma: function and clinical implications.
Hepatic stellate cells: central modulators of hepatic carcinogenesis.
Immunotherapies for hepatocellular carcinoma and intrahepatic cholangiocarcinoma: Current and developing strategies.
Establishment of a lncRNA-Based Prognostic Gene Signature Associated With Altered Immune Responses in HCC.
A Novel hepatocellular carcinoma specific hypoxic related signature for predicting prognosis and therapeutic responses.
Elevated FBXW10 drives hepatocellular carcinoma tumorigenesis via AR-VRK2 phosphorylation-dependent GAPDH ubiquitination in male transgenic mice.
Metabolic Reprogramming of HCC: A New Microenvironment for Immune Responses.
MARCKS on Tumor-Associated Macrophages is Correlated with Immune Infiltrates and Poor Prognosis in Hepatocellular Carcinoma.
Role of TGFβ3-Smads-Sp1 axis in DcR3-mediated immune escape of hepatocellular carcinoma.
Emerging Role of Cancer-Associated Fibroblasts in Progression and Treatment of Hepatocellular Carcinoma.
Molecular subtype identification and signature construction based on Golgi apparatus-related genes for better prediction prognosis and immunotherapy response in hepatocellular carcinoma.
Targeting HNRNPM Inhibits Cancer Stemness and Enhances Antitumor Immunity in Wnt-activated Hepatocellular Carcinoma.
The TGF-β Pathway: A Pharmacological Target in Hepatocellular Carcinoma?
Intrinsic activation of β-catenin signaling by CRISPR/Cas9-mediated exon skipping contributes to immune evasion in hepatocellular carcinoma.
Down-regulation of beta-actin might be involved in dendritic cells dysfunction and subsequent hepatocellular carcinoma immune escape: a proteomic study.
A bispecific antibody targeting GPC3 and CD47 induced enhanced antitumor efficacy against dual antigen-expressing HCC.
An Immune Gene Expression Signature Associated With Development of Human Hepatocellular Carcinoma Identifies Mice That Respond to Chemopreventive Agents.
B7-H1 up-regulated expression in human hepatocellular carcinoma tissue: correlation with tumor interleukin-10 levels.
An Immunogram for an Individualized Assessment of the Antitumor Immune Response in Patients With Hepatocellular Carcinoma.
Identification of prognostic markers for hepatocellular carcinoma based on the epithelial-mesenchymal transition-related gene BIRC5.
A novel risk score based on immune-related genes for hepatocellular carcinoma as a reliable prognostic biomarker and correlated with immune infiltration.
[Changes in the immune function of dendritic cells (DC) derived from HBV-related hepatocellular carcinoma (HCC) patient's peripheral blood monocytes (PBMC) pulsed with tumor antigen].

HERC5 is a prognostic biomarker for post-liver transplant recurrent human hepatocellular carcinoma.
A novel immune classification reveals distinct immune escape mechanism and genomic alterations: implications for immunotherapy in hepatocellular carcinoma.
The dual checkpoint blockade in unresectable hepatocellular carcinoma: opportunities emerging in clinical trials.
High Indoleamine 2,3-Dioxygenase Expression along with Low Bridging Integrator-1 Expression in Hepatocellular Carcinoma Patients.
Inflamed and non-inflamed classes of HCC: a revised immunogenomic classification.
Nonalcoholic steatohepatitis-related hepatocellular carcinoma: pathogenesis and treatment.
M2-like tumor-associated macrophage-related biomarkers to construct a novel prognostic signature, reveal the immune landscape, and screen drugs in hepatocellular carcinoma.
Glucose metabolism reprogramming promotes immune escape of hepatocellular carcinoma cells.
Spatial transcriptomics analysis of neoadjuvant cabozantinib and nivolumab in advanced hepatocellular carcinoma identifies independent mechanisms of resistance and recurrence.
Heterogeneous immunogenomic features and distinct escape mechanisms in multifocal hepatocellular carcinoma.
Link of sorafenib resistance with the tumor microenvironment in hepatocellular carcinoma: Mechanistic insights.
Development and Verification of a Combined Immune- and Metabolism-Related Prognostic Signature for Hepatocellular Carcinoma.
Loss of CD155 expression predicts poor prognosis in hepatocellular carcinoma.
Promotion effects of DEHP on hepatocellular carcinoma models: up-regulation of PD-L1 by activating the JAK2/STAT3 pathway.
Crocetin imparts antiproliferative activity via inhibiting STAT3 signaling in hepatocellular carcinoma.
Intrahepatic metastases may be specific to hepatocellular carcinoma due to the coagulation and fibrinolytic systems (Review).
Therapeutic modulators of hepatic stellate cells for hepatocellular carcinoma.
PKC $\alpha$ /ZFP64/CSF1 axis resets the tumor microenvironment and fuels anti-PD1 resistance in hepatocellular carcinoma.
Overcoming resistance to immune checkpoint inhibitors in hepatocellular carcinoma: Challenges and opportunities.
Possible mechanisms associated with immune escape and apoptosis on anti-hepatocellular carcinoma effect of Mu Ji Fang granules.
Liver Cancer Cell of Origin, Molecular Class, and Effects on Patient Prognosis.
Early Hepatic Lesions Display Immature Tertiary Lymphoid Structures and Show Elevated Expression of Immune Inhibitory and Immunosuppressive Molecules.
Dual Targeting of Sorafenib-Resistant HCC-Derived Cancer Stem Cells.
Molecular biology of liver cancer stem cells.
Blocking siglec-10(hi) tumor-associated macrophages improves anti-tumor immunity and enhances immunotherapy for hepatocellular carcinoma.
Noninvasive evaluation of neutrophil extracellular traps signature predicts clinical outcomes and immunotherapy response in hepatocellular carcinoma.
TNF- $\alpha$ -mediated epithelial-to-mesenchymal transition regulates expression of immune checkpoint molecules in hepatocellular carcinoma.
Genes responsible for the characteristics of primary cultured invasive phenotype hepatocellular carcinoma cells.

Oncogenic miR-93-5p/Gal-9 axis drives CD8 (+) T-cell inactivation and is a therapeutic target for hepatocellular carcinoma immunotherapy.
A cuproptosis random forest cox score model-based evaluation of prognosis, mutation characterization, immune infiltration, and drug sensitivity in hepatocellular carcinoma.
Identification of a novel signature based on macrophage-related marker genes to predict prognosis and immunotherapeutic effects in hepatocellular carcinoma.
Receptor-Interacting Protein Kinase 3 Deficiency Recruits Myeloid-Derived Suppressor Cells to Hepatocellular Carcinoma Through the Chemokine (C-X-C Motif) Ligand 1-Chemokine (C-X-C Motif) Receptor 2 Axis.
Progress in clinical oncolytic virus-based therapy for hepatocellular carcinoma.
DNASE1L3 as a Prognostic Biomarker Associated with Immune Cell Infiltration in Cancer.
Modulation of chemotactic and pro-inflammatory activities of endothelial progenitor cells by hepatocellular carcinoma.
A novel hepatocellular carcinoma-specific mTORC1-related signature for anticipating prognosis and immunotherapy.
CCL23 in Balancing the Act of Endoplasmic Reticulum Stress and Antitumor Immunity in Hepatocellular Carcinoma.
Gut Microbiota Promotes Obesity-Associated Liver Cancer through PGE(2)-Mediated Suppression of Antitumor Immunity.
Role of 5-methylcytosine in determining the prognosis, tumor microenvironment, and applicability of precision medicine in patients with hepatocellular carcinoma.
Comprehensive Analysis of HHLA2 as a Prognostic Biomarker and Its Association With Immune Infiltrates in Hepatocellular Carcinoma.
A cellular senescence-related classifier based on a tumorigenesis- and immune infiltration-guided strategy can predict prognosis, immunotherapy response, and candidate drugs in hepatocellular carcinoma.
Development of a Costimulatory Molecule Signature to Predict Prognosis, Immune Landscape, and Response to Immune Therapy for Hepatocellular Carcinoma.
Insights on the CXCL12-CXCR4 axis in hepatocellular carcinoma carcinogenesis.
Profiling of hepatocellular carcinoma neoantigens reveals immune microenvironment and clonal evolution related patterns.
High expression of B7-H2 or B7-H3 is associated with poor prognosis in hepatocellular carcinoma.
Retinoblastoma protein potentiates the innate immune response in hepatocytes: significance for hepatocellular carcinoma.
Deregulated bile acids may drive hepatocellular carcinoma metastasis by inducing an immunosuppressive microenvironment.
Research on Predicting the Occurrence of Hepatocellular Carcinoma Based on Notch Signal-Related Genes Using Machine Learning Algorithms.
Role of $\beta$ -Catenin Activation in the Tumor Immune Microenvironment and Immunotherapy of Hepatocellular Carcinoma.
Hepatic Stellate Cell Modulates the Immune Microenvironment in the Progression of Hepatocellular Carcinoma.
Cystathionine $\beta$ -synthase mediated PRRX2/IL-6/STAT3 inactivation suppresses Tregs infiltration and induces apoptosis to inhibit HCC carcinogenesis.
Over-expression of MiR-122 promotes apoptosis of hepatocellular carcinoma via targeting TLR4.
Overexpression of proteinase inhibitor 9 is associated with poor prognosis in human hepatocellular carcinoma and with proliferation and apoptosis in HepG2 cells in vitro.

Homotypic Cell Membrane-Cloaked Biomimetic Nanocarrier for the Targeted Chemotherapy of Hepatocellular Carcinoma.
Hepatitis B-Induced IL8 Promotes Hepatocellular Carcinoma Venous Metastasis and Intrahepatic Treg Accumulation.
Analysis of gene expression profiles as a tool to uncover tumor markers of liver cancer progression in a rat model.
The Role of Cancer-Associated Fibroblasts in Hepatocellular Carcinoma and the Value of Traditional Chinese Medicine Treatment.
Unique molecular characteristics of NAFLD-associated liver cancer accentuate $\beta$ -catenin/TNFRSF19-mediated immune evasion.
Alpha fetoprotein promotes polarization of macrophages towards M2-like phenotype and inhibits macrophages to phagocytize hepatoma cells.
A RIPK3-independent role of MLKL in suppressing parthanatos promotes immune evasion in hepatocellular carcinoma.
Alpha-fetoprotein impairs APC function and induces their apoptosis.
The Identification and Validation of Two Heterogenous Subtypes and a Risk Signature Based on Ferroptosis in Hepatocellular Carcinoma.
Extracellular Vesicle-Based Therapeutic Targeting of $\beta$ -Catenin to Modulate Anticancer Immune Responses in Hepatocellular Cancer.
3,5,3'-Triiodothyronine-Loaded Liposomes Inhibit Hepatocarcinogenesis Via Inflammation-Associated Macrophages.
Phase I study of MRX34, a liposomal miR-34a mimic, administered twice weekly in patients with advanced solid tumors.
Long non-coding RNA cox-2 prevents immune evasion and metastasis of hepatocellular carcinoma by altering M1/M2 macrophage polarization.
MET Inhibitors Promote Liver Tumor Evasion of the Immune Response by Stabilizing PDL1.
Chemical and genetic inhibition of STAT3 sensitizes hepatocellular carcinoma cells to sorafenib induced cell death.
Expression of TNF-related apoptosis-inducing ligand in human hepatocellular carcinoma.
Multi-omics Analysis of Microenvironment Characteristics and Immune Escape Mechanisms of Hepatocellular Carcinoma.
Dissecting spatial heterogeneity and the immune-evasion mechanism of CTCs by single-cell RNA-seq in hepatocellular carcinoma.
Suppression of toll-like receptor 2 expression inhibits the bioactivity of human hepatocellular carcinoma.
Intratumoral CD45(+)CD71(+) erythroid cells induce immune tolerance and predict tumor recurrence in hepatocellular carcinoma.
The portrait of liver cancer is shaped by mitochondrial genetics.
H(2)S suppresses indoleamine 2, 3-dioxygenase 1 and exhibits immunotherapeutic efficacy in murine hepatocellular carcinoma.
Granulin-epithelin precursor interacts with 78-kDa glucose-regulated protein in hepatocellular carcinoma.
In vivo assessment of angioarchitecture and microcirculation in experimental liver cancer: a new model in rats.
Loss of CD95 expression is linked to most but not all p53 mutants in European hepatocellular carcinoma.

Regulation of alpha-fetoprotein by nuclear factor-kappaB protects hepatocytes from tumor necrosis factor-alpha cytotoxicity during fetal liver development and hepatic oncogenesis.
Increased levels of interleukin-10 in serum from patients with hepatocellular carcinoma correlate with profound numerical deficiencies and immature phenotype of circulating dendritic cell subsets. [Relation between changes of dendritic cell function and down-regulation of beta-actin in hepatocellular carcinoma].
Alcohol reshapes a liver premetastatic niche for cancer by extra- and intrahepatic crosstalk-mediated immune evasion.
The role of miR-155 on liver diseases by modulating immunity, inflammation and tumorigenesis.
Multi-pathway network analysis of mammalian epithelial cell responses in inflammatory environments.
The Roles of circRNAs in Liver Cancer Immunity.
CCL23 suppresses liver cancer progression through the CCR1/AKT/ESR1 feedback loop.
Salvage therapy for hepatocellular carcinoma with thalidomide.
C-C chemokine receptor type 2 promotes epithelial-to-mesenchymal transition by upregulating matrix metalloproteinase-2 in human liver cancer.
WITHDRAWN:PPAR $\alpha$ Suppresses PD-L1-Mediated Immune Escape by Down-regulating SPP1 in Human Hepatocellular Carcinoma.
Glycolysis in Chronic Liver Diseases: Mechanistic Insights and Therapeutic Opportunities.
ALDH2 Hampers Immune Escape in Liver Hepatocellular Carcinoma through ROS/Nrf2-mediated Autophagy.

Table A2-15, Cluster 14

Cluster 14 focuses on Non-Small-Cell Lung Cancer, emphasizing regulation of PD-L1 expression on tumor cells to minimize tumor immune evasion which occurs by inhibiting functional activity of cytotoxic lymphocytes (350)
Potential Role of Tumor-Derived Exosomes in Non-Small-Cell Lung Cancer in the Era of Immunotherapy (350)
Development of PD-1/PD-L1 Pathway in Tumor Immune Microenvironment and Treatment for Non-Small Cell Lung Cancer.
Mechanisms of immune evasion and current status of checkpoint inhibitors in non-small cell lung cancer.
Targeting the PD-1/PD-L1 axis for the treatment of non-small-cell lung cancer.
Non-small cell lung cancer in China.
An Update on Predictive Biomarkers for Treatment Selection in Non-Small Cell Lung Cancer.
Lung cancer: potential targets for immunotherapy.
Implementation of immunotherapy in the treatment of advanced non-small cell lung cancer (NSCLC).
Emerging immunotherapies in the treatment of non-small cell lung cancer (NSCLC): the role of immune checkpoint inhibitors.
Current immune therapeutic strategies in advanced or metastatic non-small cell lung cancer.
[Recent Advances in Immunotherapy for Non-Small Cell Lung Cancer].
Immunotherapy in non-small-cell lung cancer: a bridge between research and clinical practice.
PD-1/PD-L1 immune checkpoint blockade in non-small cell lung cancer.
PD-1 and PD-L1 as immunotherapy targets and biomarkers in non-small cell lung cancer.
SNF5 promotes cell proliferation and immune evasion in non-small cell lung cancer.



Immune checkpoint inhibitors: the new frontier in non-small-cell lung cancer treatment.
SChLAP1 contributes to non-small cell lung cancer cell progression and immune evasion through regulating the AUF1/PD-L1 axis.
The progress and confusion of anti-PD1/PD-L1 immunotherapy for patients with advanced non-small cell lung cancer.
Expression landscapes in non-small cell lung cancer shaped by the thyroid transcription factor 1.
The B7-H4 gene induces immune escape partly via upregulating the PD-1/Stat3 pathway in non-small cell lung cancer.
LINC00665 knockdown confers sensitivity in irradiated non-small cell lung cancer cells through the miR-582-5p/Uchl3/AhR axis.
Current status of immunotherapy for non-small cell lung cancer.
[Advances of PD-1/PD-L1 signaling pathway in immune escape and treatment for <sup>(P)</sup> <sub>SEP</sub> non-small cell lung cancer].
Mechanistic Insights of Anti-Immune Evasion by Nobiletin through Regulating miR-197/STAT3/PD-L1 Signaling in Non-Small Cell Lung Cancer (NSCLC) Cells.
Immunotherapy for ALK-Rearranged Non-Small Cell Lung Cancer: Challenges Inform Promising Approaches.
Elevated circASCC3 limits antitumor immunity by sponging miR-432-5p to upregulate C5a in non-small cell lung cancer.
Biomarkers of resistance to immune checkpoint inhibitors in non-small-cell lung cancer: myth or reality?
Chemotherapy-induced immunomodulation in non-small-cell lung cancer: a rationale for combination chemioimmunotherapy.
The significance of Siglec-15 expression in resectable non-small cell lung cancer.
Estradiol initiates the immune escape of non-small cell lung cancer cells via ERβ/SIRT1/FOXO3a/PD-L1 axis.
MUC3A induces PD-L1 and reduces tyrosine kinase inhibitors effects in EGFR-mutant non-small cell lung cancer.
[Expression of PD-1 and PD-L1 in the peripheral blood of advanced non-small-cell lung cancer patients and its implications].
KLHL18 inhibits the proliferation, migration, and invasion of non-small cell lung cancer by inhibiting PI3K/PD-L1 axis activity.
CDK7 inhibitor THZ1 enhances antiPD-1 therapy efficacy via the p38α/MYC/PD-L1 signaling in non-small cell lung cancer.
Unravelling the Role of miR-20b-5p, CCNB1, HMGA2 and E2F7 in Development and Progression of Non-Small Cell Lung Cancer (NSCLC).
Circ_0092012 knockdown restrains non-small cell lung cancer progression by inhibiting cell malignant phenotype and immune escape through microRNA-635/programmed death ligand 1 axis.
Circular RNA circFGFR1 promotes progression and anti-PD-1 resistance by sponging miR-381-3p in non-small cell lung cancer cells.
CYP4F2-Catalyzed Metabolism of Arachidonic Acid Promotes Stromal Cell-Mediated Immunosuppression in Non-Small Cell Lung Cancer.
Diagnostic and Therapeutic Implications of microRNAs in Non-Small Cell Lung Cancer.
Targeting YAP/TAZ in Combination with PD-L1 Immune Checkpoint Inhibitors in Non-Small Cell Lung Cancer (NSCLC).
Advances in immunotherapy for non-small cell lung cancer.

A Review About Pembrolizumab in First-Line Treatment of Advanced NSCLC: Focus on KEYNOTE Studies.
Programmed Cell Death Protein 1/Programmed Cell Death Protein Ligand 1 Immunosuppressants in Advanced Non-Small Cell Lung Cancer Research Progress in Treatment.
CircDENND2D Inhibits PD-L1-Mediated Non-Small Cell Lung Cancer Metastasis and Immune Escape by Regulating miR-130b-3p/STK11 Axis.
Prospects and feasibility of synergistic therapy with radiotherapy, immunotherapy, and DNA methyltransferase inhibitors in non-small cell lung cancer.
CircFOXK2 enhances tumorigenesis and immune evasion in non-small cell lung cancer by miR-485-5p/PD-L1 axis.
Current Immunotherapeutic Strategies Targeting the PD-1/PD-L1 Axis in Non-Small Cell Lung Cancer with Oncogenic Driver Mutations.
Combination of immunotherapy with chemotherapy and radiotherapy in lung cancer: is this the beginning of the end for cancer?
Effectiveness of immunological agents in non-small cell lung cancer.
Combination immunotherapy strategies in advanced non-small cell lung cancer (NSCLC): Does biological rationale meet clinical needs?
Exploring the potential of immuno-oncology-based treatment for patients with non-small cell lung cancer.
Targeting Immunometabolism Mediated by CD73 Pathway in EGFR-Mutated Non-small Cell Lung Cancer: A New Hope for Overcoming Immune Resistance.
Different biomarkers in non-small cell lung cancer in blood vessel invasion.
[Immunotherapy in non-small cell lung cancer: evolution of knowledge and clinical advances].
Clinical significance of programmed death-1 ligand-1 expression in patients with non-small cell lung cancer: a 5-year-follow-up study.
PD-L1 and epithelial-mesenchymal transition in circulating tumor cells from non-small cell lung cancer patients: A molecular shield to evade immune system?
Immune checkpoint modulation for non-small cell lung cancer.
MicroRNA-138-5p Suppresses Non-small Cell Lung Cancer Cells by Targeting PD-L1/PD-1 to Regulate Tumor Microenvironment.
Therapeutic vaccines in non-small cell lung cancer.
[ $\beta$ -Catenin Expression in Non-Small Cell Lung Cancer and Therapeutic Effect of Immune Checkpoint Inhibitors].
Tumor Immune Microenvironment and Immunotherapy in Non-Small Cell Lung Cancer: Update and New Challenges.
Immunotherapy for Advanced Lung Cancer.
The Combination of New Immunotherapy and Radiotherapy: A New Potential Treatment for Locally Advanced Non-Small Cell Lung Cancer.
EGFR-Mutated Non-Small Cell Lung Cancer and Resistance to Immunotherapy: Role of the Tumor Microenvironment.
Immunotherapy in Non-Small Cell Lung Cancer: Biological Principles and Future Opportunities.
Hyperprogressive disease in non-small cell lung cancer after PD-1/PD-L1 inhibitors immunotherapy: underlying killer.
Programmed death-1 upregulation is correlated with dysfunction of tumor-infiltrating CD8 <sup>+</sup> T lymphocytes in human non-small cell lung cancer.
Immunotherapy in extensive small cell lung cancer.

Immunomodulatory functions of the circ_001678/miRNA-326/ZEB1 axis in non-small cell lung cancer via the regulation of PD-1/PD-L1 pathway.
Pembrolizumab in the treatment of metastatic non-small-cell lung cancer: patient selection and perspectives.
Clinical Characterization of the Expression of Insulin-Like Growth Factor Binding Protein 1 and Tumor Immunosuppression Caused by Ferroptosis of Neutrophils in Non-Small Cell Lung Cancer.
Increased expression of the immunosuppressive interleukin-35 in patients with non-small cell lung cancer.
Molecular basis and rationale for combining immune checkpoint inhibitors with chemotherapy in non-small cell lung cancer.
Circ-HSP90A expedites cell growth, stemness, and immune evasion in non-small cell lung cancer by regulating STAT3 signaling and PD-1/PD-L1 checkpoint.
Hsa_circ_0003528 promotes cell malignant transformation and immune escape via increasing oncogene PDL1 through sponging miR-511-3p in non-small cell lung cancer.
Immuno-targeted combinations in oncogene-addicted non-small cell lung cancer.
Systematic analysis of IL-6 as a predictive biomarker and desensitizer of immunotherapy responses in patients with non-small cell lung cancer.
Immunotherapy: is a minor god yet in the pantheon of treatments for lung cancer?
[Non-small cell lung cancer: news from immunotherapy].
Determinants of immunological evasion and immunecheckpoint inhibition response in non-small cell lung cancer: the genetic front.
The Combination of Immune Checkpoint Blockade and Angiogenesis Inhibitors in the Treatment of Advanced Non-Small Cell Lung Cancer.
LncRNA SOX2-OT/miR-30d-5p/PDK1 Regulates PD-L1 Checkpoint Through the mTOR Signaling Pathway to Promote Non-small Cell Lung Cancer Progression and Immune Escape.
circMET promotes NSCLC cell proliferation, metastasis, and immune evasion by regulating the miR-145-5p/CXCL3 axis.
Immunotherapy for non-small-cell lung cancer.
Synergizing systemic responses by combining immunotherapy with radiotherapy in metastatic non-small cell lung cancer: The potential of the abscopal effect.
Ubiquitin C-terminal hydrolase L1 promotes expression of programmed cell death-ligand 1 in non-small-cell lung cancer cells.
Emerging therapeutic strategies for enhancing sensitivity and countering resistance to programmed cell death protein 1 or programmed death-ligand 1 inhibitors in non-small cell lung cancer.
Heterogeneity of PD-L1 Expression and Relationship with Biology of NSCLC.
Immune checkpoint therapy for non-small-cell lung cancer: an update.
Non-Small-Cell Lung Cancer Signaling Pathways, Metabolism, and PD-1/PD-L1 Antibodies.
Differential miRNA expressions in peripheral blood mononuclear cells for diagnosis of lung cancer.
Atezolizumab: A novel PD-L1 inhibitor in cancer therapy with a focus in bladder and non-small cell lung cancers.
Has programmed cell death ligand-1 MET an accomplice in non-small cell lung cancer?-a narrative review.
EZH2 regulates PD-L1 expression via HIF-1 $\alpha$ in non-small cell lung cancer cells.
Prognostic factors in non-small cell lung carcinoma.
Untangling the evolutionary roots of lung cancer.
Inhibition of c-MET upregulates PD-L1 expression in lung adenocarcinoma.

IL-9 Producing Tumor-Infiltrating Lymphocytes and Treg Subsets Drive Immune Escape of Tumor Cells in Non-Small Cell Lung Cancer.
Circular RNA drives resistance to anti-PD-1 immunotherapy by regulating the miR-30a-5p/SOX4 axis in non-small cell lung cancer.
Small molecule TKI inhibitors affect the development of non-small cell carcinoma through HIPPO/YAP/PD-L1.
Proteogenomics of non-small cell lung cancer reveals molecular subtypes associated with specific therapeutic targets and immune evasion mechanisms.
Beyond PD-L1 testing-emerging biomarkers for immunotherapy in non-small cell lung cancer.
Alterations of immune response of Non-Small Cell Lung Cancer with Azacytidine.
[IoNECO trial: Immune neoadjuvant therapy in early stage non-small cell lung cancer].
Circular RNA circ-CPA4/ let-7 miRNA/PD-L1 axis regulates cell growth, stemness, drug resistance and immune evasion in non-small cell lung cancer (NSCLC).
Potential immune escape mechanisms underlying the distinct clinical outcome of immune checkpoint blockades in small cell lung cancer.
The clinical relevance of the miR-197/CKS1B/STAT3-mediated PD-L1 network in chemoresistant non-small-cell lung cancer.
Small cell transformation of non-small cell lung cancer on immune checkpoint inhibitors: uncommon or under-recognized?
Prognostic Factors and Biomarkers of Responses to Immune Checkpoint Inhibitors in Lung Cancer.
New strategies in immunotherapy for lung cancer: beyond PD-1/PD-L1.
Tumour Microenvironment and Immune Evasion in EGFR Addicted NSCLC: Hurdles and Possibilities.
Enhanced Acid Sphingomyelinase Activity Drives Immune Evasion and Tumor Growth in Non-Small Cell Lung Carcinoma.
Immune subgroup analysis for non-small cell lung cancer may be a good choice for evaluating therapeutic efficacy and prognosis.
Programmed Death-Ligand 1 Expression Predicts Tyrosine Kinase Inhibitor Response and Better Prognosis in a Cohort of Patients With Epidermal Growth Factor Receptor Mutation-Positive Lung Adenocarcinoma.
Understanding the checkpoint blockade in lung cancer immunotherapy.
A Phase 2 Study of Docetaxel, Ramucirumab, and Pembrolizumab for Patients With Metastatic or Recurrent Non-Small-Cell Lung Cancer (NSCLC) who Progressed on Platinum-Doublet and PD-1/PD-L1 Blockade.
MUC1-C integrates PD-L1 induction with repression of immune effectors in non-small-cell lung cancer.
Macrophage infiltration in human non-small-cell lung cancer: the role of CC chemokines.
Therapeutic value of EGFR inhibition in CRC and NSCLC: 15 years of clinical evidence.
Circular RNA circ_0101675 Promotes NSCLC Cell Proliferation, Migration, Invasion, Angiogenesis and Immune Evasion by Sponging miR-607/PDL1 Axis.
[Advances of the Role of Lung Cancer Driver Gene and PD-1/PD-L1 Pathway Interaction in the Tumorigenesis and Progression of Non-small Cell Lung Cancer].
Artesunate promoted anti-tumor immunity and overcame EGFR-TKI resistance in non-small-cell lung cancer by enhancing oncogenic TAZ degradation.
Immunosuppressive TREM2(+) macrophages are associated with undesirable prognosis and responses to anti-PD-1 immunotherapy in non-small cell lung cancer.
Development of Novel CD47-Specific ADCs Possessing High Potency Against Non-Small Cell Lung Cancer in vitro and in vivo.

PD-L1 confers resistance to EGFR mutation-independent tyrosine kinase inhibitors in non-small cell lung cancer via upregulation of YAP1 expression.
A Novel mTORC1/2 Inhibitor (MTI-31) Inhibits Tumor Growth, Epithelial-Mesenchymal Transition, Metastases, and Improves Antitumor Immunity in Preclinical Models of Lung Cancer.
Flotillin-1 enhances radioresistance through reducing radiation-induced DNA damage and promoting immune escape via STING signaling pathway in non-small cell lung cancer.
Osimertinib for Front-Line Treatment of Locally Advanced or Metastatic EGFR-Mutant NSCLC Patients: Efficacy, Acquired Resistance and Perspectives for Subsequent Treatments.
Novel genetic variants of KIR3DL2 and PVR involved in immunoregulatory interactions are associated with non-small cell lung cancer survival.
Targeting programmed cell death protein 1 (PD-1) for treatment of non-small-cell lung carcinoma (NSCLC); the recent advances.
Combined Consideration of Tumor-Associated Immune Cell Density and Immune Checkpoint Expression in the Peritumoral Microenvironment for Prognostic Stratification of Non-Small-Cell Lung Cancer Patients.
Diagnosis value of serum soluble B7-H4 expression in non-small cell lung cancer.
Platelet PD-L1 reflects collective intratumoral PD-L1 expression and predicts immunotherapy response in non-small cell lung cancer.
Immunoregulation and Clinical Implications of ANGPT2/TIE2(+) M-MDSC Signature in Non-Small Cell Lung Cancer.
Failure of Immunotherapy-The Molecular and Immunological Origin of Immunotherapy Resistance in Lung Cancer.
Antimetabolite pemetrexed primes a favorable tumor microenvironment for immune checkpoint blockade therapy.
EGFR-TKI resistance promotes immune escape in lung cancer via increased PD-L1 expression.
FoxP3 genetic variants and risk of non-small cell lung cancer in the Chinese Han population.
N(6)-methyladenosine-modified circIGF2BP3 inhibits CD8(+) T-cell responses to facilitate tumor immune evasion by promoting the deubiquitination of PD-L1 in non-small cell lung cancer.
Locally advanced lung cancer: an optimal setting for vaccines and other immunotherapies.
Emerging drugs for EGFR-mutated non-small cell lung cancer.
Eosinophil granulocytes account for indoleamine 2,3-dioxygenase-mediated immune escape in human non-small cell lung cancer.
Thalidomide suppresses angiogenesis and immune evasion via lncRNA FGD5-AS1/miR-454-3p/ZEB1 axis-mediated VEGFA expression and PD-1/PD-L1 checkpoint in NSCLC.
Morphine-3-glucuronide upregulates PD-L1 expression via TLR4 and promotes the immune escape of non-small cell lung cancer.
Expression of tumor necrosis factor receptor 2 in human non-small cell lung cancer and its role as a potential prognostic biomarker.
Circular RNA hsa_circ_0000190 Facilitates the Tumorigenesis and Immune Evasion by Upregulating the Expression of Soluble PD-L1 in Non-Small-Cell Lung Cancer.
Clinical significance of the induction of macrophage differentiation by the costimulatory molecule B7-H3 in human non-small cell lung cancer.
The Interplay Between Programmed Death Ligand 1 and Vimentin in Advanced Non-Small-Cell Lung Cancer.
Case Report: Circulating Myeloid-Derived Suppressive-Like Cells and Exhausted Immune Cells in Non-Small Cell Lung Cancer Patients Treated With Three Immune Checkpoint Inhibitors.
Expression of Three Clones of PD-L1 in Lung Cancer: A Single-center Experience.

Angiotensin II contributes to intratumoral immunosuppression via induction of PD-L1 expression in non-small cell lung carcinoma.
Circular RNA hsa_circ_0068252 Functions in Cisplatin Resistance and Immune Response via miR-1304-5p/PD-L1 Axis in Non-Small Cell Lung Cancer.
MRTF-A-NF- $\kappa$ B/p65 axis-mediated PDL1 transcription and expression contributes to immune evasion of non-small-cell lung cancer via TGF- $\beta$ .
Study on the Expression Levels and Clinical Significance of PD-1 and PD-L1 in Plasma of NSCLC Patients.
Continuous targeted kinase inhibitors treatment induces upregulation of PD-L1 in resistant NSCLC.
Therapeutic and Systemic Adverse Events of Immune Checkpoint Inhibitors Targeting the PD-1/PD-L1 axis for Clinical Management of NSCLC.
A Bayesian Network Meta-Analysis of First-Line Treatments for Non-Small Cell Lung Cancer with High Programmed Death Ligand-1 Expression.
The Expression of Programmed Death Ligand 1 and Vimentin in Resected Non-Metastatic Non-Small-Cell Lung Cancer: Interplay and Prognostic Effects.
An increase in BAG-1 by PD-L1 confers resistance to tyrosine kinase inhibitor in non-small cell lung cancer via persistent activation of ERK signalling.
MEK and EGFR inhibition demonstrate synergistic activity in EGFR-dependent NSCLC.
Immune Escape Mechanisms in Non Small Cell Lung Cancer.
Modulating Tumor Microenvironment: A Review on STK11 Immune Properties and Predictive vs Prognostic Role for Non-small-cell Lung Cancer Immunotherapy.
BIN1 reverses PD-L1-mediated immune escape by inactivating the c-MYC and EGFR/MAPK signaling pathways in non-small cell lung cancer.
Gefitinib enhances the anti-tumor immune response against EGFR-mutated NSCLC by upregulating B7H5 expression and activating T cells via CD28H.
Epithelial-to-mesenchymal transition promotes immune escape by inducing CD70 in non-small cell lung cancer.
Immunotherapy in the treatment of non-small cell lung cancer.
Optimization of the Enrichment of Circulating Tumor Cells for Downstream Phenotypic Analysis in Patients with Non-Small Cell Lung Cancer Treated with Anti-PD-1 Immunotherapy.
Inhibition of Non-Small Cell Lung Cancer Cells by Oxy210, an Oxysterol-Derivative that Antagonizes TGF $\beta$ and Hedgehog Signaling.
Analysis of the immunological microenvironment at the tumor site in patients with non-small cell lung cancer.
FOXP3 and TLR4 protein expression are correlated in non-small cell lung cancer: implications for tumor progression and escape.
Safety and activity of anti-PD-L1 antibody in patients with advanced cancer.
Tumor immune microenvironment in epidermal growth factor receptor-mutated non-small cell lung cancer before and after epidermal growth factor receptor tyrosine kinase inhibitor treatment: a narrative review.
Wnt/ $\beta$ -Catenin Signaling and Resistance to Immune Checkpoint Inhibitors: From Non-Small-Cell Lung Cancer to Other Cancers.
The radiological appearances of lung cancer treated with immunotherapy.
Discovery of Potent OTUB1/USP8 Dual Inhibitors Targeting Proteostasis in Non-Small-Cell Lung Cancer.
B7-H3 in combination with regulatory T cell is associated with tumor progression in primary human non-small cell lung cancer.

Mean platelet volume, thrombocytosis, and survival in non-small cell lung cancer patients treated with first-line pembrolizumab alone or with chemotherapy.
OPA1 supports mitochondrial dynamics and immune evasion to CD8(+) T cell in lung adenocarcinoma.
Pharmacodynamics of Pre-Operative PD1 checkpoint blockade and receptor activator of NFkB ligand (RANKL) inhibition in non-small cell lung cancer (NSCLC): study protocol for a multicentre, open-label, phase 1B/2, translational trial (POPCORN).
Arginase 2 is expressed by human lung cancer, but it neither induces immune suppression, nor affects disease progression.
[Up-regulation of TIM-3 on CD4+ tumor infiltrating lymphocytes predicts poor prognosis in human non-small-cell lung cancer].
Alteration of circulating natural autoantibodies to CD25-derived peptide antigens and FOXP3 in non-small cell lung cancer.
Clinical outcomes of immune checkpoint blockades and the underlying immune escape mechanisms in squamous and adenocarcinoma NSCLC.
Obesity accelerates immune evasion of non-small cell lung carcinoma via TFEB-dependent upregulation of Siglec-15 and glycolytic reprogramming.
Epigenetic Therapy Ties MYC Depletion to Reversing Immune Evasion and Treating Lung Cancer.
Molecular mechanisms of acquired resistance to third-generation EGFR-TKIs in EGFR T790M-mutant lung cancer.
PARP1 inhibition elicits immune responses against non-small cell lung cancer.
Corrigendum: LncRNA SOX2-OT/miR-30d-5p/PDK1 Regulates PD-L1 Checkpoint Through the mTOR Signaling Pathway to Promote Non-Small Cell Lung Cancer Progression and Immune Escape.
PD-L1 expression and its effect on clinical outcomes of EGFR-mutant NSCLC patients treated with EGFR-TKIs.
A novel bifunctional anti-PD-L1/TGF-β Trap fusion protein (M7824) efficiently reverts mesenchymalization of human lung cancer cells.
PD-L1/p-STAT3 promotes the progression of NSCLC cells by regulating TAM polarization.
[Operability and Pathological Response of Non-Small Cell Lung Cancer (NSCLC) after Neoadjuvant Therapy with Immune Checkpoint Inhibition].
Analysis of non-small cell lung cancer microenvironment indicates preponderance of T cell exhaustion marker expression.
Distinct Functional Metagenomic Markers Predict the Responsiveness to Anti-PD-1 Therapy in Chinese Non-Small Cell Lung Cancer Patients.
Anti-PD-(L)1 therapy of non-small cell lung cancer-A summary of clinical trials and current progresses.
The Future of Immunotherapy in the Treatment of Small Cell Lung Cancer.
ILT3 promotes tumor cell motility and angiogenesis in non-small cell lung cancer.
Role of immunotherapy in metastatic EGFRm NSCLC: Is it relevant?
Interleukin-10-regulated tumour tolerance in non-small cell lung cancer.
Targeting the STAT5A/IDO1 axis overcomes radioresistance and reverses the immunosuppressive tumor microenvironment in NSCLC.
[The in situ analysis of cytokine mRNA expression in immunological microenvironment about non-small cell lung cancer].
Progressive changes in composition of lymphocytes in lung tissues from patients with non-small-cell lung cancer.
Current status and challenges of immunotherapy in ALK rearranged NSCLC.
Correlation Study on the Expression of INSR, IRS-1, and PD-L1 in Nonsmall Cell Lung Cancer.

Baseline extracellular vesicle TGF- $\beta$ is a predictive biomarker for response to immune checkpoint inhibitors and survival in non-small cell lung cancer.
Circulating tumor cell methylation profiles reveal the classification and evolution of non-small cell lung cancer.
KRAS G12V mutation upregulates PD-L1 expression via TGF- $\beta$ /EMT signaling pathway in human non-small-cell lung cancer.
Orthotopic model of lung cancer: isolation of bone micro-metastases after tumor escape from Osimertinib treatment.
Recombinant human erythropoietin accelerated the proliferation of non-small cell lung cancer cell lines and reduced the expression of VEGF, HIF-1 $\alpha$ , and PD-L1 under a simulated hypoxic environment in vitro.
Why has active immunotherapy not worked in lung cancer?
Silencing of circCRIM1 Drives IGF2BP1-Mediated NSCLC Immune Evasion.
Clinical relevance of galectin-1 expression in non-small cell lung cancer patients.
Increase in soluble PD-1 is associated with prolonged survival in patients with advanced EGFR-mutated non-small cell lung cancer treated with erlotinib.
EGFR may participate in immune evasion through regulation of B7-H5 expression in non-small cell lung carcinoma.
Biological therapies in nonsmall cell lung cancer.
Impaired Tumor-Infiltrating T Cells in Patients with Chronic Obstructive Pulmonary Disease Impact Lung Cancer Response to PD-1 Blockade.
An overview of the crosstalk between YAP and cGAS-STING signaling in non-small cell lung cancer: it takes two to tango.
Clinical significance of B7-H4 expression in matched non-small cell lung cancer brain metastases and primary tumors.
Characterizing heterogeneity of non-small cell lung tumour microenvironment to identify signature prognostic genes.
Systems Immunology Analysis Reveals an Immunomodulatory Effect of Snail-p53 Binding on Neutrophil- and T Cell-Mediated Immunity in KRAS Mutant Non-Small Cell Lung Cancer.
SBRT combined with PD-1/PD-L1 inhibitors in NSCLC treatment: a focus on the mechanisms, advances, and future challenges.
[Vascular endothelial growth factor inhibits dendritic cells from patients with non-small cell lung carcinoma].
The targeting of indoleamine 2,3 dioxygenase -mediated immune escape in cancer.
Neoantigen-directed immune escape in lung cancer evolution.
Recent Advances in Lung Cancer Immunotherapy: Input of T-Cell Epitopes Associated With Impaired Peptide Processing.
Head to head evaluation of second generation ALK inhibitors brigatinib and alectinib as first-line treatment for ALK+ NSCLC using an in silico systems biology-based approach.
The role of spatial interplay patterns between PD-L1-positive tumor cell and T cell in recurrence of locally advanced non-small cell lung cancer.
A Novel Antagonist of the Immune Checkpoint Protein Adenosine A2a Receptor Restores Tumor-Infiltrating Lymphocyte Activity in the Context of the Tumor Microenvironment.
A scoping review on the significance of programmed death-ligand 1-inhibiting microRNAs in non-small cell lung treatment: A single-cell RNA sequencing-based study.
Analyzing the percentage of different PD-1(+) T cell subsets in peripheral blood and bronchoalveolar lavage fluid of small cell lung cancer patients: A prospective study.



Immune-related adverse events associated with programmed cell death protein-1 and programmed cell death ligand 1 inhibitors for non-small cell lung cancer: a PRISMA systematic review and meta-analysis.
Commensal microbiota contributes to predicting the response to immune checkpoint inhibitors in non-small-cell lung cancer patients.
Upregulation of PD-L1 by EGFR Activation Mediates the Immune Escape in EGFR-Driven NSCLC: Implication for Optional Immune Targeted Therapy for NSCLC Patients with EGFR Mutation.
Molecular mechanism of lncRNA SNHG12 in immune escape of non-small cell lung cancer through the HuR/PD-L1/USP8 axis.
Human lung carcinomas synthesize immunoregulatory glucocorticoids.
[bcl-2 expression in small cell lung cancer: a mechanism for apoptosis antagonism and immune evasion].
CD47 Expression in Circulating Tumor Cells and Circulating Tumor Microemboli from Non-Small Cell Lung Cancer Patients Is a Poor Prognosis Factor.
TKI-resistant ALK-rearranged lung adenocarcinoma with secondary CTNNB1 p.S45V and tertiary ALK p.I1171N mutations.
Management of Small Cell Lung Cancer: Progress and Updates.
Immunotherapy in NSCLC patients with brain metastases. Understanding brain tumor microenvironment and dissecting outcomes from immune checkpoint blockade in the clinic.
Na, K-ATPase $\alpha 1$ cooperates with its endogenous ligand to reprogram immune microenvironment of lung carcinoma and promotes immune escape.
EGFR inhibition in EGFR-mutant lung cancer cells perturbs innate immune signaling pathways in the tumor microenvironment.
Targeting the PD1/PD-L1 axis in melanoma: biological rationale, clinical challenges and opportunities.
Addressing resistance to PD-1/PD-(L)1 pathway inhibition: considerations for combinatorial clinical trial designs.
Expression Analysis and Significance of PD-1, LAG-3, and TIM-3 in Human Non-Small Cell Lung Cancer Using Spatially Resolved and Multiparametric Single-Cell Analysis.
Digital gene expression analysis of NSCLC-patients reveals strong immune pressure, resulting in an immune escape under immunotherapy.
FBW7-mediated ubiquitination and destruction of PD-1 protein primes sensitivity to anti-PD-1 immunotherapy in non-small cell lung cancer.
Down-regulation of ERAP1 mRNA expression in non-small cell lung cancer.
The immune checkpoint, HVEM may contribute to immune escape in non-small cell lung cancer lacking PD-L1 expression.
Activation of the PD-1 pathway contributes to immune escape in EGFR-driven lung tumors.
Tenascin-C, over expressed in lung cancer down regulates effector functions of tumor infiltrating lymphocytes.
Concomitant Mycobacterium tuberculosis infection promotes lung tumor growth through enhancing Treg development.
Immune regulatory function of cancer-associated fibroblasts in non-small cell lung cancer.
STK11/LKB1 Deficiency Promotes Neutrophil Recruitment and Proinflammatory Cytokine Production to Suppress T-cell Activity in the Lung Tumor Microenvironment.
Strategies for co-targeting the PI3K/AKT/mTOR pathway in NSCLC.
Effect of cyclo-oxygenase inhibitor use during checkpoint blockade immunotherapy in patients with metastatic melanoma and non-small cell lung cancer.

Inactivation of AMPK Leads to Attenuation of Antigen Presentation and Immune Evasion in Lung Adenocarcinoma.
Circ_0010235 facilitates lung cancer development and immune escape by regulating miR-636/PDL1 axis.
Biological insights in non-small cell lung cancer.
Acquired DNA damage repairs deficiency-driven immune evolution and involved immune factors of local versus distant metastases in non-small cell lung cancer.
PPAR $\gamma$ inhibited tumor immune escape by inducing PD-L1 autophagic degradation.
The superior efficacy of anti-PD-1/PD-L1 immunotherapy in KRAS-mutant non-small cell lung cancer that correlates with an inflammatory phenotype and increased immunogenicity.
STAT3-targeted treatment with silibinin overcomes the acquired resistance to crizotinib in ALK-rearranged lung cancer.
Dysregulation of circulating CDC42 and its correlation with demographic characteristics, comorbidities, tumor features, chemotherapeutic regimen and survival profile in non-small-cell lung cancer patients.
The status of immunosuppression in patients with stage IIIB or IV non-small-cell lung cancer correlates with the clinical characteristics and response to chemotherapy.
ILT4 inhibition prevents TAM- and dysfunctional T cell-mediated immunosuppression and enhances the efficacy of anti-PD-L1 therapy in NSCLC with EGFR activation.
Phase II Study Evaluating the Mechanisms of Resistance on Tumor Tissue and Liquid Biopsy in Patients With EGFR-mutated Non-pretreated Advanced Lung Cancer Receiving Osimertinib Until and Beyond Radiologic Progression: The MELROSE Trial.
Tumor $\beta$ -catenin expression is associated with immune evasion in non-small cell lung cancer with high tumor mutation burden.
A new role for circulating T follicular helper cells in humoral response to anti-PD-1 therapy.
Circ-METTL15 contributes to the proliferation, metastasis, immune escape and restrains apoptosis in lung cancer by regulating miR-1299/PDL1 axis.
High-dose irradiation in combination with toll-like receptor 9 agonist CpG oligodeoxynucleotide 7909 downregulates PD-L1 expression via the NF- $\kappa$ B signaling pathway in non-small cell lung cancer cells.
Self-associated molecular patterns mediate cancer immune evasion by engaging Siglecs on T cells.
SKIL facilitates tumorigenesis and immune escape of NSCLC via upregulating TAZ/autophagy axis.
Dendritic cells infiltrating human non-small cell lung cancer are blocked at immature stage.
Upregulation of PD-L1 by EML4-ALK fusion protein mediates the immune escape in ALK positive NSCLC: Implication for optional anti-PD-1/PD-L1 immune therapy for ALK-TKIs sensitive and resistant NSCLC patients.
Atezolizumab for use in PD-L1-positive unresectable, locally advanced or metastatic triple-negative breast cancer.
[Research Progress on the Mechanism and Clinical Data of Cereblon <sup>(P)</sup> <sub>SEP</sub> in Reversing the Resistance of Lung Cancer to PD-1 Antibody by T cells].
Beyond the PD-L1 horizon: In search for a good biomarker to predict success of immunotherapy in gastric and esophageal adenocarcinoma.
Myeloid-derived suppressor cells infiltration in non-small-cell lung cancer tumor and MAGE-A4 and NY-ESO-1 expression.
Indoleamine 2,3-dioxygenase (IDO) expression in lung cancer.
Interleukin-10 haplotype may predict survival and relapse in resected non-small cell lung cancer.
Intrinsic and Extrinsic Regulation of PD-L2 Expression in Oncogene-Driven Non-Small Cell Lung Cancer.
Immunotherapy strategy of EGFR mutant lung cancer.

MiR-192 and miR-662 enhance chemoresistance and invasiveness of squamous cell lung carcinoma.
Neutrophil expansion defines an immunoinhibitory peripheral and intratumoral inflammatory milieu in resected non-small cell lung cancer: a descriptive analysis of a prospectively immunoprofiled cohort.
Tissue micro array analysis of ganglioside N-glycolyl GM3 expression and signal transducer and activator of transcription (STAT)-3 activation in relation to dendritic cell infiltration and microvessel density in non-small cell lung cancer.
[Expression of PD-1/PD-L1 in peripheral blood mononuclear cells in lung cancer patients and its biological significance].
Induced expression of B7-H4 on the surface of lung cancer cell by the tumor-associated macrophages: a potential mechanism of immune escape.
CXCR4 Inhibition Counteracts Immunosuppressive Properties of Metastatic NSCLC Stem Cells.
Nivolumab and pembrolizumab: Monoclonal antibodies against programmed cell death-1 (PD-1) that are interchangeable.
Neutralization of complement regulatory proteins CD55 and CD59 augments therapeutic effect of herceptin against lung carcinoma cells.
Evading the STING: LKB1 Loss Leads to STING Silencing and Immune Escape in KRAS-Mutant Lung Cancers.
Immunotherapy for Lung Cancer.
The Interaction of the IFN $\gamma$ /JAK/STAT1 and JAK/STAT3 Signalling Pathways in EGFR-Mutated Lung Adenocarcinoma Cells.
Circular RNA CHST15 Sponges miR-155-5p and miR-194-5p to Promote the Immune Escape of Lung Cancer Cells Mediated by PD-L1.
Preliminary mechanisms of regulating PD-L1 expression in non-small cell lung cancer during the EMT process.
Narrative review of immunotherapy in thymic malignancies.
[Intervention effect of Feiji Recipe on immune escape of lung cancer].
Prognostic impact of LILRB4 expression on tumor-infiltrating cells in resected non-small cell lung cancer.
A subset of VEGFR-TKIs activates AMPK in LKB1-mutant lung cancer.
Successful Treatment of a Patient With Multiple-Line Relapsed Extensive-Stage Small-Cell Lung Cancer Receiving Penpulimab Combined With Anlotinib: A Case Report.
Wide Expression and Significance of Alternative Immune Checkpoint Molecules, B7x and HHLA2, in PD-L1-Negative Human Lung Cancers.
Afatinib restrains K-RAS-driven lung tumorigenesis.
Deep and Prolonged Response to Aurora A Kinase Inhibitor and Subsequently to Nivolumab in MYCL1-Driven Small-Cell Lung Cancer: Case Report and Literature Review.
The combination of anti-KIR monoclonal antibodies with anti-PD-1/PD-L1 monoclonal antibodies could be a critical breakthrough in overcoming tumor immune escape in NSCLC.
High plasma levels of soluble programmed cell death ligand 1 are prognostic for reduced survival in advanced lung cancer.
Association between IDO activity and prognosis in patients with non-small cell lung cancer after radiotherapy.
Characteristics of T-Cell Receptor Repertoire and Correlation With EGFR Mutations in All Stages of Lung Cancer.

Metastatic pulmonary pleomorphic carcinoma replaced by a granulomatous lesion after spontaneous regression and PD-1 blockade-induced regression: can epithelioid granuloma be a histological hallmark of cancer immunity?
Expression and prognostic significance in lung cancer of human tumor-associated antigen RCAS1.
Nivolumab monotherapy or combination therapy with ipilimumab for lung cancer: a systemic review and meta-analysis.
CD73 Inhibitor Oleclumab Plus Osimertinib in Previously Treated Patients With Advanced T790M-Negative EGFR-Mutated NSCLC: A Brief Report.
CRACD loss promotes small cell lung cancer tumorigenesis via EZH2-mediated immune evasion.
Tumor glycolytic profiling through (18)F-FDG PET/CT predicts immune checkpoint inhibitor efficacy in advanced NSCLC.
B2M gene expression shapes the immune landscape of lung adenocarcinoma and determines the response to immunotherapy.
CD47-blocking immunotherapies stimulate macrophage-mediated destruction of small-cell lung cancer.
CDK5 Activates Hippo Signaling to Confer Resistance to Radiation Therapy Via Upregulating TAZ in Lung Cancer.
XIST/miR-34a-5p/PDL1 axis regulated the development of lung cancer cells and the immune function of CD8(+) T cells.
Interleukin-10 (IL-10) promoter genotypes are associated with lung cancer risk in Taiwan males and smokers.
The TLR4/ERK/PD-L1 axis may contribute to NSCLC initiation.
Managing Hyperprogressive Disease in the Era of Programmed Cell Death Protein 1/Programmed Death-Ligand 1 Blockade: A Case Discussion and Review of the Literature.
Mevalonate improves anti-PD-1/PD-L1 efficacy by stabilizing CD274 mRNA.
Quantitative Image Analysis of Fibrillar Collagens Reveals Novel Diagnostic and Prognostic Biomarkers and Histotype-Dependent Aberrant Mechanobiology in Lung Cancer.
Expression of cancer-testis antigens MAGE-A3/6 and NY-ESO-1 in non-small-cell lung carcinomas and their relationship with immune cell infiltration.
Feiji Recipe inhibits the growth of lung cancer by modulating T-cell immunity through indoleamine-2,3-dioxygenase pathway in an orthotopic implantation model.
Somatic mutations and immune checkpoint biomarkers.
Afatinib plus Cetuximab Delays Resistance Compared to Single-Agent Erlotinib or Afatinib in Mouse Models of TKI-Naïve EGFR L858R-Induced Lung Adenocarcinoma.
A Novel Epitope Quality-Based Immune Escape Mechanism Reveals Patient's Suitability for Immune Checkpoint Inhibition.
Elevated NOX4 promotes tumorigenesis and acquired EGFR-TKIs resistance via enhancing IL-8/PD-L1 signaling in NSCLC.
PDL1 Regulation by p53 via miR-34.
Autoantibody panel on small extracellular vesicles for the early detection of lung cancer.
Increased serum kynurenine/tryptophan ratio correlates with disease progression in lung cancer.
Sex-Based Dimorphism of Anticancer Immune Response and Molecular Mechanisms of Immune Evasion.
BTLA-HVEM Couple in Health and Diseases: Insights for Immunotherapy in Lung Cancer.
Blockading a new NSCLC immunosuppressive target by pluripotent autologous tumor vaccines magnifies sequential immunotherapy.
Suppression of STING Associated with LKB1 Loss in KRAS-Driven Lung Cancer.

Histologic assessment of tumor-associated CD45(+) cell numbers is an independent predictor of prognosis in small cell lung cancer.
Downregulation of NKG2DLs by TGF- $\beta$ in human lung cancer cells.
Design, synthesis, and biological activity of dual monoamine oxidase A and heat shock protein 90 inhibitors, N-Methylpropargylamine-conjugated 4-isopropylresorcinol for glioblastoma.
Expression of tumor-associated antigen RCAS1 correlates significantly with poor prognosis in nonsmall cell lung carcinoma.
SEC61G assists EGFR-amplified glioblastoma to evade immune elimination.
The Correlation Between SPP1 and Immune Escape of EGFR Mutant Lung Adenocarcinoma Was Explored by Bioinformatics Analysis.
Integrated Clinical, Molecular and Immunological Characterization of Pulmonary Sarcomatoid Carcinomas Reveals an Immune Escape Mechanism That May Influence Therapeutic Strategies.
Prognostic value of soluble H7-B4 in pleural effusion associated with lung cancer.
Targeting USP22 with miR-30-5p to inhibit the hypoxia-induced expression of PD-L1 in lung adenocarcinoma cells.
Homeobox A1 Facilitates Immune Escape and Alleviates Oxidative Stress in Lung Adenocarcinoma.
The M1/M2 spectrum and plasticity of malignant pleural effusion-macrophage in advanced lung cancer.
E2F1 induces apoptosis and sensitizes human lung adenocarcinoma cells to death-receptor-mediated apoptosis through specific downregulation of c-FLIP(short).
Granzyme B in peripheral blood mononuclear cells as a measure of cell-mediated immune response in paraneoplastic neurological syndromes and malignancy.
Evidence of Epstein-Barr virus heterogeneous gene expression in adult lung transplant recipients with posttransplant lymphoproliferative disorder.

Table A2-16, Cluster 15

Cluster 15 focuses on cancer immunotherapy using IDO1 inhibitors that will transform the tumor microenvironment from tolerogenic to normal immunogenic by interfering with the IDO1-kynurenine-aryl hydrocarbon receptor pathway (159)
Indoleamine 2,3-Dioxygenase (IDO) Activity: A Perspective Biomarker for Laboratory Determination in Tumor Immunotherapy.
Indoleamine 2,3-Dioxygenase 1: A Promising Therapeutic Target in Malignant Tumor.
Indoleamine 2,3-dioxygenase 1 (IDO1) inhibitors in clinical trials for cancer immunotherapy.
Discovery of a Novel Scaffold as an Indoleamine 2,3-Dioxygenase 1 (IDO1) Inhibitor Based on the Pyrrolpiperazinone Alkaloid, Longamide B.
Indoleamine 2,3-Dioxygenase and Its Therapeutic Inhibition in Cancer.
IDO1: An important immunotherapy target in cancer treatment.
Synthesis, Docking and Biological Evaluation of a Novel Class of Imidazothiazoles as IDO1 Inhibitors.
Recent discovery of indoleamine-2,3-dioxygenase 1 inhibitors targeting cancer immunotherapy.
The Advances of the Structure and Function of Indoleamine 2, 3- dioxygenase 1 and Its Inhibitors.
IDO1 in cancer: a Gemini of immune checkpoints.
An updated patent review of IDO1 inhibitors for cancer (2018-2022).
Hydroxyamidine inhibitors of indoleamine-2,3-dioxygenase potently suppress systemic tryptophan catabolism and the growth of IDO-expressing tumors.

4,5-Disubstituted 1,2,3-triazoles: Effective Inhibition of Indoleamine 2,3-Dioxygenase 1 Enzyme Regulates T cell Activity and Mitigates Tumor Growth.
IDO1 plays an immunosuppressive role in 2,4,6-trinitrobenzene sulfate-induced colitis in mice.
Design and Synthesis of Indoleamine 2,3-Dioxygenase 1 Inhibitors and Evaluation of Their Use as Anti-Tumor Agents.
A patent review of IDO1 inhibitors for cancer.
The potential of targeting indoleamine 2,3-dioxygenase for cancer treatment.
Tryptophan: A Rheostat of Cancer Immune Escape Mediated by Immunosuppressive Enzymes IDO1 and TDO.
Carbidopa, an activator of aryl hydrocarbon receptor, suppresses IDO1 expression in pancreatic cancer and decreases tumor growth.
Immunological and nonimmunological effects of indoleamine 2,3-dioxygenase on breast tumor growth and spontaneous metastasis formation.
Cell-Based Identification of New IDO1 Modulator Chemotypes.
Limitations and Off-Target Effects of Tryptophan-Related IDO Inhibitors in Cancer Treatment.
Discovery of 5-(pyridin-3-yl)-1H-indole-4,7-diones as indoleamine 2,3-dioxygenase 1 (IDO1) inhibitors.
Tumor immune escape mediated by indoleamine 2,3-dioxygenase.
N-Benzyl/Aryl Substituted Tryptanthrin as Dual Inhibitors of Indoleamine 2,3-Dioxygenase and Tryptophan 2,3-Dioxygenase.
Role of indoleamine 2,3-dioxygenase in acute myeloid leukemia.
The interplay between indoleamine 2,3-dioxygenase 1 (IDO1) and cyclooxygenase (COX)-2 in chronic inflammation and cancer.
Indoleamine 2,3-dioxygenase in cancer: targeting pathological immune tolerance with small-molecule inhibitors.
Microenvironment Tumor Metabolic Interactions Highlighted by qMSI: Application to the Tryptophan-Kynurenine Pathway in Immuno-Oncology.
Targeting immunometabolism mediated by the IDO1 Pathway: A new mechanism of immune resistance in endometrial cancer.
Modulating Tumor Immunology by Inhibiting Indoleamine 2,3-Dioxygenase (IDO): Recent Developments and First Clinical Experiences.
Indoleamine 2,3-dioxygenase in T-cell tolerance and tumoral immune escape.
Indoleamine 2,3-dioxygenase, an emerging target for anti-cancer therapy.
Design, synthesis and biological evaluation of 2,5-dimethylfuran-3-carboxylic acid derivatives as potential IDO1 inhibitors.
In vivo metabolism of tryptophan in meningiomas is mediated by indoleamine 2,3-dioxygenase 1.
Important Hydrogen Bond Networks in Indoleamine 2,3-Dioxygenase 1 (IDO1) Inhibitor Design Revealed by Crystal Structures of Imidazoleisoindole Derivatives with IDO1.
Preclinical Characterization of Linrodostat Mesylate, a Novel, Potent, and Selective Oral Indoleamine 2,3-Dioxygenase 1 Inhibitor.
Indoleamine 2,3-dioxygenase expression in human cancers: clinical and immunologic perspectives.
The expanding roles of 1-methyl-tryptophan (1-MT): in addition to inhibiting kynurenine production, 1-MT activates the synthesis of melatonin in skin cells.
Design, synthesis and biological evaluation of bicyclic carboxylic acid derivatives as IDO1 inhibitors.
Turn off the IDO: will clinical trials be successful?
IDO induces expression of a novel tryptophan transporter in mouse and human tumor cells.
IDO Targeting in Sarcoma: Biological and Clinical Implications.

Evidence for a tumoral immune resistance mechanism based on tryptophan degradation by indoleamine 2,3-dioxygenase.
Inhibition of indoleamine 2,3-dioxygenase 1 expression alters immune response in colon tumor microenvironment in mice.
Marrying immunotherapy with chemotherapy: why say IDO?
Indoleamine 2,3-dioxygenase pathways of pathogenic inflammation and immune escape in cancer.
A highly potent and selective inhibitor RoxyL-WL targeting IDO1 promotes immune response against melanoma.
Inhibiting IDO pathways to treat cancer: lessons from the ECHO-301 trial and beyond.
Targeting indoleamine-2,3-dioxygenase in cancer: Scientific rationale and clinical evidence.
Targeting the indoleamine 2,3-dioxygenase pathway in cancer.
Substrate stereo-specificity in tryptophan dioxygenase and indoleamine 2,3-dioxygenase.
PCC0208009 enhances the anti-tumor effects of temozolomide through direct inhibition and transcriptional regulation of indoleamine 2,3-dioxygenase in glioma models.
Correlation of indoleamine-2,3-dioxygenase 1 inhibitory activity of 4,6-disubstituted indazole derivatives and their heme binding affinity.
Detailed analysis and follow-up studies of a high-throughput screening for indoleamine 2,3-dioxygenase 1 (IDO1) inhibitors.
The indoleamine-2,3-dioxygenase (IDO) inhibitor 1-methyl-D-tryptophan upregulates IDO1 in human cancer cells.
[A new mechanism of tumor resistance to the immune system, based on tryptophan breakdown by indoleamine 2,3-dioxygenase].
Targeting the IDO1/TDO2-KYN-AhR Pathway for Cancer Immunotherapy - Challenges and Opportunities.
Clinical Relevance of Serum Kyn/Trp Ratio and Basal and IFN $\gamma$ -Upregulated IDO1 Expression in Peripheral Monocytes in Early Stage Melanoma.
Discovery of the First Potent IDO1/IDO2 Dual Inhibitors: A Promising Strategy for Cancer Immunotherapy.
New 4-Amino-1,2,3-Triazole Inhibitors of Indoleamine 2,3-Dioxygenase Form a Long-Lived Complex with the Enzyme and Display Exquisite Cellular Potency.
Expression, purification, and kinetic characterization of the human strep-IDO1.
Rational design of 4-aryl-1,2,3-triazoles for indoleamine 2,3-dioxygenase 1 inhibition.
Inhibition of the BET family reduces its new target gene IDO1 expression and the production of L-kynurenine.
A Novel High Throughput Virtual Screening Protocol to Discover New Indoleamine 2,3-Dioxygenase 1 (IDO1) Inhibitors.
Zinc protoporphyrin IX stimulates tumor immunity by disrupting the immunosuppressive enzyme indoleamine 2,3-dioxygenase.
Enhanced tryptophan degradation in patients with ovarian carcinoma correlates with several serum soluble immune activation markers.
Inhibitory effects of flavonoids isolated from <i>Sophora flavescens</i> on indoleamine 2,3-dioxygenase 1 activity.
Is IDO a key enzyme bridging the gap between tumor escape and tolerance induction?
Expression of indoleamine 2,3-dioxygenase in tumor endothelial cells correlates with long-term survival of patients with renal cell carcinoma.
Nanosensor detection of an immunoregulatory tryptophan influx/kynurenine efflux cycle.
Indoleamine 2,3-dioxygenase inhibitory activity of derivatives of marine alkaloid tsitsikammamine A.

Targeting tryptophan availability to tumors: the answer to immune escape?
Identification of selective inhibitors of indoleamine 2,3-dioxygenase 2.
Expression of indoleamine 2,3-dioxygenase and correlation with pathological malignancy in gliomas.
Signal Transducer and Activator of Transcription 1 Plays a Pivotal Role in RET/PTC3 Oncogene-induced Expression of Indoleamine 2,3-Dioxygenase 1.
Endotoxin-Induced Tryptophan Degradation along the Kynurenine Pathway: The Role of Indoleamine 2,3-Dioxygenase and Aryl Hydrocarbon Receptor-Mediated Immunosuppressive Effects in Endotoxin Tolerance and Cancer and Its Implications for Immunoparalysis.
Non-hematopoietic expression of IDO is integrally required for inflammatory tumor promotion.
Inhibition of indoleamine 2,3-dioxygenase, an immunoregulatory target of the cancer suppression gene Bin1, potentiates cancer chemotherapy.
Pathogenetic Interplay Between IL-6 and Tryptophan Metabolism in an Experimental Model of Obesity.
The Kynurenine Pathway and Cancer: Why Keep It Simple When You Can Make It Complicated.
Tumoral Immune Resistance Mediated by Enzymes That Degrade Tryptophan.
Strategic Incorporation of Polarity in Heme-Displacing Inhibitors of Indoleamine-2,3-dioxygenase-1 (IDO1).
Bifunctional Naphthoquinone Aromatic Amide-Oxime Derivatives Exert Combined Immunotherapeutic and Antitumor Effects through Simultaneous Targeting of Indoleamine-2,3-dioxygenase and Signal Transducer and Activator of Transcription 3.
Target exposure and pharmacodynamics study of the indoleamine 2,3-dioxygenase-1 (IDO-1) inhibitor epacadostat in the CT26 mouse tumor model.
Decreased serum tryptophan concentration predicts poor prognosis in malignant melanoma patients.
Canine mast cell tumour cells regulate tryptophan catabolism via the expression of indoleamine 2,3-dioxygenase.
Identification of Human IDO1 Enzyme Activity by Using Genetically Encoded Nitrotyrosine.
IDO-expressing regulatory dendritic cells in cancer and chronic infection.
A highly efficient modality to block the degradation of tryptophan for cancer immunotherapy: locked nucleic acid-modified antisense oligonucleotides to inhibit human indoleamine 2,3-dioxygenase 1/tryptophan 2,3-dioxygenase expression.
A Pt(IV) Pro-drug Preferentially Targets Indoleamine-2,3-dioxygenase, Providing Enhanced Ovarian Cancer Immuno-Chemotherapy.
GBP1 Facilitates Indoleamine 2,3-Dioxygenase Extracellular Secretion to Promote the Malignant Progression of Lung Cancer.
The paradoxical patterns of expression of indoleamine 2,3-dioxygenase in colon cancer.
IDO-Mediated Tryptophan Degradation in the Pathogenesis of Malignant Tumor Disease.
IDO1 and TDO inhibitory evaluation of analogues of the marine pyrroloiminoquinone alkaloids: Wakayin and Tsitsikammamines.
Rational design of indoleamine 2,3-dioxygenase inhibitors.
Nutritional Stress Induced by Tryptophan-Degrading Enzymes Results in ATF4-Dependent Reprogramming of the Amino Acid Transporter Profile in Tumor Cells.
A key in vivo antitumor mechanism of action of natural product-based brassinins is inhibition of indoleamine 2,3-dioxygenase.
Indoleamine 2, 3-Dioxygenase 1 Mediates Survival Signals in Chronic Lymphocytic Leukemia via Kynurenine/Aryl Hydrocarbon Receptor-Mediated MCL1 Modulation.



Preclinical investigations and a first-in-human phase I trial of M4112, the first dual inhibitor of indoleamine 2,3-dioxygenase 1 and tryptophan 2,3-dioxygenase 2, in patients with advanced solid tumors.
Indoleamine 2,3-dioxygenase 1 (IDO1) is up-regulated in thyroid carcinoma and drives the development of an immunosuppressant tumor microenvironment.
Salmonella-mediated therapy targeting indoleamine 2, 3-dioxygenase 1 (IDO) activates innate immunity and mitigates colorectal cancer growth.
Indoleamine 2,3-dioxygenase as a modifier of pathogenic inflammation in cancer and other inflammation-associated diseases.
Discovery of novel hydroxyamidine derivatives as indoleamine 2,3-dioxygenase 1 inhibitors with in vivo anti-tumor efficacy.
Indol-2-yl ethanones as novel indoleamine 2,3-dioxygenase (IDO) inhibitors.
IDO1 involvement in mTOR pathway: a molecular mechanism of resistance to mTOR targeting in medulloblastoma.
Discovery of the first potent proteolysis targeting chimera (PROTAC) degrader of indoleamine 2,3-dioxygenase 1.
Tryptophan catabolism in epithelial ovarian carcinoma.
Localization of indoleamine 2,3-dioxygenase in human esophageal squamous cell carcinomas.
Accumulation of an endogenous tryptophan-derived metabolite in colorectal and breast cancers.
MiR-218 produces anti-tumor effects on cervical cancer cells in vitro.
Immune escape as a fundamental trait of cancer: focus on IDO.
First-in-Human Phase I Study of the Oral Inhibitor of Indoleamine 2,3-Dioxygenase-1 Epacadostat (INCB024360) in Patients with Advanced Solid Malignancies.
Epacadostat Plus Pembrolizumab and Chemotherapy for Advanced Solid Tumors: Results from the Phase I/II ECHO-207/KEYNOTE-723 Study.
On-line screening of indoleamine 2,3-dioxygenase 1 inhibitors by partial filling capillary electrophoresis combined with rapid polarity switching.
The tryptophan-kynurenine pathway in immunomodulation and cancer metastasis.
Role of indoleamine 2,3-dioxygenase in pathology of the gastrointestinal tract.
Inverse correlation between tumoral indoleamine 2,3-dioxygenase expression and tumor-infiltrating lymphocytes in endometrial cancer: its association with disease progression and survival.
1-Methyl-D-tryptophan potentiates TGF- $\beta$ -induced epithelial-mesenchymal transition in T24 human bladder cancer cells.
Serum tryptophan and kynurenine concentrations as parameters for indoleamine 2,3-dioxygenase activity in patients with endometrial, ovarian, and vulvar cancer.
(-)-Epigallocatechin gallate inhibits the expression of indoleamine 2,3-dioxygenase in human colorectal cancer cells.
The indoleamine 2,3-dioxygenase pathway controls complement-dependent enhancement of chemoradiation therapy against murine glioblastoma.
Towards a genetic definition of cancer-associated inflammation: role of the IDO pathway.
Aptamer-conjugated nano-liposome for immunogenic chemotherapy with reversal of immunosuppression.
1-MT inhibits the invasion of CBP-resistant ovarian cancer cells via down-regulating IDO expression and re-activating immune cells function.
Tryptophan metabolites modulate inflammatory bowel disease and colorectal cancer by affecting immune system.
1,2,3-Triazoles as inhibitors of indoleamine 2,3-dioxygenase 2 (IDO2).

Nanoconjugates to enhance PDT-mediated cancer immunotherapy by targeting the indoleamine-2,3-dioxygenase pathway.
Early Carcinogenesis Involves the Establishment of Immune Privilege via Intrinsic and Extrinsic Regulation of Indoleamine 2,3-dioxygenase-1: Translational Implications in Cancer Immunotherapy.
Kynurenines as a Novel Target for the Treatment of Malignancies.
Tryptophan: Its Metabolism along the Kynurenine, Serotonin, and Indole Pathway in Malignant Melanoma.
Calcium-pterin suppresses mitogen-induced tryptophan degradation and neopterin production in peripheral blood mononuclear cells.
Indoleamine-2,3-dioxygenase in an immunotherapy model for Ewing sarcoma.
IDO1(+) Paneth cells promote immune escape of colorectal cancer.
Indoleamine 2,3-dioxygenase, an immunomodulatory protein, is suppressed by (-)-epigallocatechin-3-gallate via blocking of gamma-interferon-induced JAK-PKC-delta-STAT1 signaling in human oral cancer cells.
Not Only Immune Escape-The Confusing Role of the TRP Metabolic Pathway in Carcinogenesis.
Dendritic cell vaccination, immune regulation, and clinical outcomes in ovarian cancer.
The kynurenine pathway in brain tumor pathogenesis.
Synergistic Transcutaneous Immunotherapy Enhances Antitumor Immune Responses through Delivery of Checkpoint Inhibitors.
Kynurenine inhibits melanogenesis in human melanocyte-keratinocyte co-cultures and in a reconstructed 3D skin model.
Targeting the tumor microenvironment by liposomal Epacadostat in combination with liposomal gp100 vaccine.
Eicosapentaenoic acid reduces indoleamine 2,3-dioxygenase 1 expression in tumor cells.
Indoleamine-2,3 dioxygenase: a fate-changer of the tumor microenvironment.
Tryptophan Metabolism: A Versatile Area Providing Multiple Targets for Pharmacological Intervention.
Suppression of TDO-mediated tryptophan catabolism in glioblastoma cells by a steroid-responsive FKBP52-dependent pathway.
Determination of kynurenine and tryptophan, biomarkers of indoleamine 2,3-dioxygenase by LC-MS/MS in plasma and tumor.
Modulation of cancer-specific immune responses by amino acid degrading enzymes.
Recent advances in clinical trials targeting the kynurenine pathway.
Paraptosis Inducer to Effectively Trigger Immunogenic Cell Death for Metastatic Tumor Immunotherapy with IDO Inhibition.
Immunometabolic Network Interactions of the Kynurenine Pathway in Cutaneous Malignant Melanoma.
The extracts of Astragalus membranaceus enhance chemosensitivity and reduce tumor indoleamine 2, 3-dioxygenase expression.
Pt(IV) hybrids containing a TDO inhibitor serve as potential anticancer immunomodulators.
The Kynurenine Pathway Is a Double-Edged Sword in Immune-Privileged Sites and in Cancer: Implications for Immunotherapy.
Novel conjugates with dual suppression of glutathione S-transferases and tryptophan-2,3-dioxygenase activities for improving hepatocellular carcinoma therapy.
Structural Basis of Selective Human Indoleamine-2,3-dioxygenase 1 (hIDO1) Inhibition.
Nicotinamide adenine dinucleotide metabolism as an attractive target for drug discovery.

Discovery of Indoleamine 2,3-Dioxygenase 1 (IDO-1) Inhibitors Based on Ortho-Naphthaquinone-Containing Natural Product.
Investigation of the absolute bioavailability and human mass balance of navoximod, a novel IDO1 inhibitor.
Discovery of Mcl-1 inhibitors through virtual screening, molecular dynamics simulations and in vitro experiments.

Table A2-17, Cluster 16

Cluster 16 focuses on the impact of shedding of MICA/B on tumor cell immune evasion, emphasizing treatments that reduce this shedding and augment the cytotoxic function of Natural Killer cells (171)
Natural killer cell cytotoxicity is suppressed by exposure to the human NKG2D ligand MICA*008 that is shed by tumor cells in exosomes.
Regulation Roles of MICA and NKG2D in Human Renal Cancer Cells.
Shedding of endogenous MHC class I-related chain molecules A and B from different human tumor entities: heterogeneous involvement of the "a disintegrin and metalloproteases" 10 and 17.
Generation of soluble NKG2D ligands: proteolytic cleavage, exosome secretion and functional implications.
MICA immune complex formed with alpha 3 domain-specific antibody activates human NK cells in a Fc-dependent manner.
Development of a screening method to identify regulators of MICA shedding.
Impact on NK cell functions of acute versus chronic exposure to extracellular vesicle-associated MICA: Dual role in cancer immunosurveillance.
BCL11B regulates MICA/B-mediated immune response by acting as a competitive endogenous RNA.
Intracellular retention of the NKG2D ligand MHC class I chain-related gene A in human melanomas confers immune privilege and prevents NK cell-mediated cytotoxicity.
MHC class I chain-related A: Polymorphism, regulation and therapeutic value in cancer.
MICA polymorphism: biology and importance in cancer.
Inhibition of MICA and MICB Shedding Elicits NK-Cell-Mediated Immunity against Tumors Resistant to Cytotoxic T Cells.
Leveraging NKG2D Ligands in Immuno-Oncology.
Altered NKG2D function in NK cells induced by chronic exposure to NKG2D ligand-expressing tumor cells.
Loss of NKG2D in murine NK cells leads to increased perforin production upon long-term stimulation with IL-2.
The role of the NKG2D receptor for tumor immunity.
Molecular Bases for the Regulation of NKG2D Ligands in Cancer.
Natural killer group 2D receptor and its ligands in cancer immune escape.
Estrogen upregulates MICA/B expression in human non-small cell lung cancer through the regulation of ADAM17.
MICA Expression Is Regulated by Cell Adhesion and Contact in a FAK/Src-Dependent Manner.
Downregulation of MICA/B tumor surface expressions and augmented soluble MICA serum levels correlate with disease stage in breast cancer.
Cellular expression, trafficking, and function of two isoforms of human ULBP5/RAET1G.
Loss of expression of MHC class I-related chain A (MICA) is a frequent event and predicts poor survival in patients with hepatocellular carcinoma.

Human tumor-derived exosomes down-modulate NKG2D expression.
Downregulation and/or release of NKG2D ligands as immune evasion strategy of human neuroblastoma.
Reduced immune effector cell NKG2D expression and increased levels of soluble NKG2D ligands in multiple myeloma may not be causally linked.
MICA/B expression is inhibited by unfolded protein response and associated with poor prognosis in human hepatocellular carcinoma.
MMP9 mediates MICA shedding in human osteosarcomas.
Major Histocompatibility Complex Class I-Related Chain A (MICA) Allelic Variants Associate With Susceptibility and Prognosis of Gastric Cancer.
NKG2D Ligands-Critical Targets for Cancer Immune Escape and Therapy.
Mechanisms of Senescence-Related NKG2D Ligands Release and Immune Escape Induced by Chemotherapy in Neuroblastoma Cells.
NKG2D Ligand Shedding in Response to Stress: Role of ADAM10.
Silencing NKG2D ligand-targeting miRNAs enhances natural killer cell-mediated cytotoxicity in breast cancer.
The fatty-acid amide hydrolase inhibitor URB597 inhibits MICA/B shedding.
Soluble ligands for the NKG2D receptor are released during HIV-1 infection and impair NKG2D expression and cytotoxicity of NK cells.
Immune evasion by TGF $\beta$ -induced miR-183 repression of MICA/B expression in human lung tumor cells.
Tumor-associated MICA is shed by ADAM proteases.
Antibody-mediated inhibition of MICA and MICB shedding promotes NK cell-driven tumor immunity.
NKG2D ligands in tumor immunity.
Manipulation of NKG2D ligands by cytomegaloviruses: impact on innate and adaptive immune response.
Soluble MICB in malignant diseases: analysis of diagnostic significance and correlation with soluble MICA.
[Expression and abscission of activated receptors and their ligands on/from NK cells in peripheral blood of patients with acute leukemia].
Tumour-derived soluble MIC ligands impair expression of NKG2D and T-cell activation.
Expression of major histocompatibility complex class I-related chain A/B (MICA/B) in pancreatic carcinoma.
NKG2D and its ligands in cancer.
Radiosensitization effect by HDAC inhibition improves NKG2D-dependent natural killer cytotoxicity in hepatocellular carcinoma.
MICA A5.1 homozygous genotype is associated with a risk for early-onset oral cancer.
Prognostic significance and functional implication of immune activating receptor NKG2D in gastric cancer.
Pseudorabies Virus Infection Causes Downregulation of Ligands for the Activating NK Cell Receptor NKG2D.
KLF4-mediated upregulation of the NKG2D ligand MICA in acute myeloid leukemia: a novel therapeutic target identified by enChIP.
Down-regulation of NKG2D and Nkp80 ligands by Kaposi's sarcoma-associated herpesvirus K5 protects against NK cell cytotoxicity.
Soluble MICA in malignant diseases.

HER2/HER3 signaling regulates NK cell-mediated cytotoxicity via MHC class I chain-related molecule A and B expression in human breast cancer cell lines.
A vaccine targeting resistant tumours by dual T cell plus NK cell attack.
Immune evasion mediated by tumor-derived lactate dehydrogenase induction of NKG2D ligands on myeloid cells in glioblastoma patients.
HBV suppresses expression of MICA/B on hepatoma cells through up-regulation of transcription factors GATA2 and GATA3 to escape from NK cell surveillance.
Effect of NKG2D ligand expression on host immune responses.
Disulphide-isomerase-enabled shedding of tumour-associated NKG2D ligands.
BCR/ABL oncogene directly controls MHC class I chain-related molecule A expression in chronic myelogenous leukemia.
Matrix metalloproteinase 2 (MMP2) mediates MHC class I polypeptide-related sequence A (MICA) shedding in renal cell carcinoma.
TGF-beta and metalloproteinases differentially suppress NKG2D ligand surface expression on malignant glioma cells.
Restoration of antitumor immunity through anti-MICA antibodies elicited with a chimeric protein.
The membrane type matrix metalloproteinase MMP14 mediates constitutive shedding of MHC class I chain-related molecule A independent of A disintegrin and metalloproteinases.
Thermal- and oxidative stress causes enhanced release of NKG2D ligand-bearing immunosuppressive exosomes in leukemia/lymphoma T and B cells.
[The roles of soluble MICA in immune escape of breast tumor].
Functional Characterisation and Analysis of the Soluble NKG2D Ligand Repertoire Detected in Umbilical Cord Blood Plasma.
Decreased NKG2D expression on CD8+ T cell is involved in immune evasion in patients with gastric cancer.
Wiskott-Aldrich syndrome protein (WASp) and N-WASp are involved in the regulation of NK-cell migration upon NKG2D activation.
Evasion from NK cell immunity by MHC class I chain-related molecules expressing colon adenocarcinoma.
Down-regulation of the human major histocompatibility complex class I chain-related gene A (MICA) and its receptor is mediated by microRNA-146b-5p and is a potential mechanism of immunoediting in papillary thyroid carcinoma.
Tumor-derived soluble MICs impair CD3(+)CD56(+) NKT-like cell cytotoxicity in cancer patients.
Ovarian tumor-associated microRNA-20a decreases natural killer cell cytotoxicity by downregulating MICA/B expression.
Lysine acetylation of NKG2D ligand Rae-1 stabilizes the protein and sensitizes tumor cells to NKG2D immune surveillance.
The NKG2D receptor: immunobiology and clinical implications.
Modulation of NKG2D-ligand cell surface expression enhances immune cell therapy of cancer.
A hypoxia-induced decrease of either MICA/B or Hsp70 on the membrane of tumor cells mediates immune escape from NK cells.
ULBP6/RAET1L is an additional human NKG2D ligand.
Tumoral NKG2D alters cell cycle of acute myeloid leukemic cells and reduces NK cell-mediated immune surveillance.
Exploration of the role of NKG2D ligands MICA and MICB in JAK2 V617F-positive myeloproliferative neoplasms.

Soluble MICA-NKG2D interaction upregulates IFN- $\gamma$ production by activated CD3-CD56+ NK cells: potential impact on chronic graft versus host disease.
Corticotropin-releasing factor induces immune escape of cervical cancer cells by downregulation of NKG2D.
Hypoxia-induced shedding of MICA and HIF1A-mediated immune escape of pancreatic cancer cells from NK cells: role of circ_0000977/miR-153 axis.
MUC1-C is a master regulator of MICA/B NKG2D ligand and exosome secretion in human cancer cells.
Secretory pathways generating immunosuppressive NKG2D ligands: New targets for therapeutic intervention.
Tumor-Derived Soluble MICA Obstructs the NKG2D Pathway to Restrain NK Cytotoxicity.
NKG2D engagement on human NK cells leads to DNAM-1 hypo-responsiveness through different converging mechanisms.
Expression of stress ligands of the immunoreceptor NKG2D in melanoma: regulation and clinical significance.
Low-dose valproic acid with low-dose gemcitabine augments MHC class I-related chain A/B expression without inducing the release of soluble MHC class I-related chain A/B.
Adenovirus E3/19K promotes evasion of NK cell recognition by intracellular sequestration of the NKG2D ligands major histocompatibility complex class I chain-related proteins A and B.
Human cytomegalovirus glycoprotein UL16 causes intracellular sequestration of NKG2D ligands, protecting against natural killer cell cytotoxicity.
Human placenta expresses and secretes NKG2D ligands via exosomes that down-modulate the cognate receptor expression: evidence for immunosuppressive function.
Prostate tumor-derived exosomes down-regulate NKG2D expression on natural killer cells and CD8+ T cells: mechanism of immune evasion.
Cutting an NKG2D Ligand Short: Cellular Processing of the Peculiar Human NKG2D Ligand ULBP4.
MicroRNA-mediated down-regulation of NKG2D ligands contributes to glioma immune escape.
A chimeric antigen receptor uniquely recognizing MICA/B stress proteins provides an effective approach to target solid tumors.
Anterior gradient 2-derived peptide upregulates major histocompatibility complex class I-related chains A/B in hepatocellular carcinoma cells.
IL-2-activated haploidentical NK cells restore NKG2D-mediated NK-cell cytotoxicity in neuroblastoma patients by scavenging of plasma MICA.
Differential mechanisms of shedding of the glycosylphosphatidylinositol (GPI)-anchored NKG2D ligands.
Expression of ERp5 and GRP78 on the membrane of chronic lymphocytic leukemia cells: association with soluble MICA shedding.
An acetylated derivative of vitexin halts MDA-MB-231 cellular progression and improves its immunogenic profile through tuning miR-20a-MICA/B axis.
Immune surveillance in glioblastoma: Role of the NKG2D system and novel cell-based therapeutic approaches.
NKG2D ligand MICA is retained in the cis-Golgi apparatus by human cytomegalovirus protein UL142.
Convergent Evolution by Cancer and Viruses in Evading the NKG2D Immune Response.
High glucose promotes pancreatic cancer cells to escape from immune surveillance via AMPK-Bmi1-GATA2-MICA/B pathway.
Increased serum NKG2D-ligands and downregulation of NKG2D in peripheral blood NK cells of patients with major burns.
Tumor cell recognition by the NK cell activating receptor NKG2D.

The RNA binding protein IMP3 facilitates tumor immune escape by downregulating the stress-induced ligands ULBP2 and MICB.
Targeting the NKG2D/NKG2D-L axis in acute myeloid leukemia.
Methylation of NKG2D ligands contributes to immune system evasion in acute myeloid leukemia.
Membrane-bound versus soluble major histocompatibility complex Class I-related chain A and major histocompatibility complex Class I-related chain B differential expression: Mechanisms of tumor eradication versus evasion and current drug development strategies.
Metastamir-mediated immune evasion: miR-10b downregulates the stress-induced molecule MICB, hence avoid recognition by NKG2D receptor.
Heat-killed <i>Helicobacter pylori</i> upregulates NKG2D ligands expression on gastric adenocarcinoma cells via Toll-like receptor 4.
The soluble major histocompatibility complex class I-related chain A protein reduced NKG2D expression on natural killer and T cells from patients with prolactinoma and non-secreting pituitary adenoma.
Discrepant effects of <i>Chlamydia trachomatis</i> infection on MICA expression of HeLa and U373 cells.
New prospects on the NKG2D/NKG2DL system for oncology.
ULBPs, human ligands of the NKG2D receptor, stimulate tumor immunity with enhancement by IL-15.
Manipulating the NKG2D Receptor-Ligand Axis Using CRISPR: Novel Technologies for Improved Host Immunity.
Platelet-mediated shedding of NKG2D ligands impairs NK cell immune-surveillance of tumor cells.
An NKG2D-mediated human lymphoid stress surveillance response with high interindividual variation.
Soluble NKG2D ligand promotes MDSC expansion and skews macrophage to the alternatively activated phenotype.
NKG2D- and T-cell receptor-dependent lysis of malignant glioma cell lines by human $\gamma\delta$ T cells: Modulation by temozolomide and A disintegrin and metalloproteases 10 and 17 inhibitors.
Natural Killer Group 2D Ligand Depletion Reconstitutes Natural Killer Cell Immunosurveillance of Head and Neck Squamous Cell Carcinoma.
Characterization of 5' promoter and exon 1-3 polymorphism of the RAET1E gene.
The alteration of placental-derived soluble MHC class I chain-related protein A and B during pregnancy.
Association of MICA-129 polymorphism with nasopharyngeal cancer risk in a Tunisian population.
Rapamycin downregulates NKG2D ligands in acute myeloid leukemia cells via an activation of the STAT3 pathway: a potential mechanism for rapamycin-induced immune escape in leukemia.
Two Flow Cytometric Approaches of NKG2D Ligand Surface Detection to Distinguish Stem Cells from Bulk Subpopulations in Acute Myeloid Leukemia.
MiR-10b downregulates the stress-induced cell surface molecule MICB, a critical ligand for cancer cell recognition by natural killer cells.
RAET1E2, a soluble isoform of the UL16-binding protein RAET1E produced by tumor cells, inhibits NKG2D-mediated NK cytotoxicity.
Upregulation of Myc promotes the evasion of NK cell-mediated immunity through suppression of NKG2D ligands in K562 cells.
A subgroup of lupus patients with nephritis, innate T cell activation and low vitamin D is identified by the enhancement of circulating MHC class I-related chain A.
CEACAM1 dampens antitumor immunity by down-regulating NKG2D ligand expression on tumor cells.
Soluble NKG2D ligands impair CD8(+) T cell antitumor function dependent of NKG2D downregulation in neuroblastoma.

Placenta-derived soluble MHC class I chain-related molecules down-regulate NKG2D receptor on peripheral blood mononuclear cells during human pregnancy: a possible novel immune escape mechanism for fetal survival.
The murine cytomegalovirus immunoevasin gp40/m152 inhibits NKG2D receptor RAE-1 $\gamma$ by intracellular retention and cell surface masking.
NKG2D ligand tumor expression and association with clinical outcome in early breast cancer patients: an observational study.
LINC01149 variant modulates MICA expression that facilitates hepatitis B virus spontaneous recovery but increases hepatocellular carcinoma risk.
Prevalent expression of the immunostimulatory MHC class I chain-related molecule is counteracted by shedding in prostate cancer.
An six-amino acid motif in the alpha3 domain of MICA is the cancer therapeutic target to inhibit shedding.
Cancer immunoediting of the NK group 2D ligand H60a.
Soluble NKG2D ligands in hepatic autoimmune diseases and in benign diseases involved in marker metabolism.
Cooperative therapeutic anti-tumor effect of IL-15 agonist ALT-803 and co-targeting soluble NKG2D ligand sMIC.
[Mechanism of Anti Apoptosis and Immune Evasion in Drug-Resistant Leukemia Cells Mediated by STAT3].
The Relationship Between Phospho-p38, Matrix Metalloproteinase 9, and Major Histocompatibility Complex Class I Chain-Related Molecule A Expression in Pituitary Adenomas Demonstrates a New Mechanism of Pituitary Adenoma Immune Escape.
NKG2D ligand expression in pediatric brain tumors.
Superior induction and maintenance of protective CD8 T cells in mice infected with mouse cytomegalovirus vector expressing RAE-1 $\gamma$ .
DHA Abolishes the Detrimental Effect of Docetaxel on Downregulation of the MICA via Decreasing the Expression Level of MicroRNA-20a in Gastric Cancer.
Relationship between NF- $\kappa$ B, MMP-9, and MICA expression in pituitary adenomas reveals a new mechanism of pituitary adenomas immune escape.
Increase of CIK cell efficacy by upregulating cell surface MICA and inhibition of NKG2D ligand shedding in multiple myeloma.
HDAC3 represses the expression of NKG2D ligands ULBPs in epithelial tumour cells: potential implications for the immunosurveillance of cancer.
MICA-129Met/Val Polymorphism Is Associated with Early-Onset Breast Cancer Risk.
IDH mutant gliomas escape natural killer cell immune surveillance by downregulation of NKG2D ligand expression.
MALAT-1: Immunomodulatory lncRNA hampering the innate and the adaptive immune arms in triple negative breast cancer.
Soluble MIC is elevated in the serum of patients with pancreatic carcinoma diminishing gammadelta T cell cytotoxicity.
Hypoxia increases tumor cell shedding of MHC class I chain-related molecule: role of nitric oxide.
Recruiting T cells and sensitizing tumors to NKG2D immune surveillance for robust antitumor immune response.
Loss of nonclassical MHC molecules MIC-A/B expression during progression of uveal melanoma.
c-Myc Targets HDAC3 to Suppress NKG2DL Expression and Innate Immune Response in N-Type SCLC through Histone Deacetylation.



Publisher Correction: Absence of NKG2D ligands defines leukaemia stem cells and mediates their immune evasion.
Human $\gamma\delta$ T-Cells: From Surface Receptors to the Therapy of High-Risk Leukemias.
Cannabinoid receptor CB1 regulates STAT3 activity and its expression dictates the responsiveness to SR141716 treatment in human glioma patients' cells.
Cervical Cancer Cells Express Markers Associated with Immunosurveillance.
Hypoxia-mediated immune evasion of pancreatic carcinoma cells.
Potential therapeutic applications of phosphodiesterase inhibition in prostate cancer.
Inhibition of glycolipid shedding rescues recognition of a CD1+ T cell lymphoma by natural killer T (NKT) cells.
SUMOylation inhibitor TAK-981 (subasumstat) synergizes with 5-azacitidine in preclinical models of acute myeloid leukemia.
Bacillus anthracis lethal toxin disrupts TCR signaling in CD1d-restricted NKT cells leading to functional anergy.
Tumour-expressed tissue factor inhibits cellular cytotoxicity.
Expression and modulation of progesterone induced blocking factor (PIBF) and innate immune factors in human leukemia cell lines by progesterone and mifepristone.
Insufficient ex vivo expansion of Valpha24(+) natural killer T cells in malignant lymphoma patients related to the suppressed expression of CD1d molecules on CD14(+) cells.
Investigation of the mechanisms of tissue factor-mediated evasion of tumour cells from cellular cytotoxicity.

Table A2-18, Cluster 17

Cluster 17 focuses on pancreatic cancer/pancreatic ductal adenocarcinoma, and the role of immunotherapies in countering the immune system evasion associated with a tolerogenic tumor microenvironment (199)
Pancreatic Cancer Signaling Pathways, Genetic Alterations, and Tumor Microenvironment: The Barriers Affecting the Method of Treatment.
Immunosuppression, immune escape, and immunotherapy in pancreatic cancer: focused on the tumor microenvironment.
Inflammatory cytokines in human pancreatic cancer.
A subset of epithelial cells mimics regulatory T cells and contributes to immune evasion during development of pancreatic adenocarcinoma.
Is targeting autophagy a promising lead to unveil the cloak of invisibility in pancreatic cancer?
Understanding the immune response and the current landscape of immunotherapy in pancreatic cancer.
Broadening the Impact of Immunotherapy to Pancreatic Cancer: Challenges and Opportunities.
Current immunotherapeutic approaches in pancreatic cancer.
Persistent activation of pancreatic stellate cells creates a microenvironment favorable for the malignant behavior of pancreatic ductal adenocarcinoma.
Gemcitabine inhibits immune escape of pancreatic cancer by down regulating the soluble ULBP2 protein.
Molecular Mechanisms and Potential Therapeutic Reversal of Pancreatic Cancer-Induced Immune Evasion.
Pancreatic cancer and its stroma: a conspiracy theory.

KRAS mutation: The booster of pancreatic ductal adenocarcinoma transformation and progression.
Advances in therapeutic vaccines for pancreatic cancer.
Challenges and Future Perspectives of Immunotherapy in Pancreatic Cancer.
Secretome analysis of multiple pancreatic cancer cell lines reveals perturbations of key functional networks.
Exploiting inflammation for therapeutic gain in pancreatic cancer.
Pancreatic adenocarcinoma exerts systemic effects on the peripheral blood myeloid and plasmacytoid dendritic cells: an indicator of disease severity?
Mirage or long-awaited oasis: reinvigorating T-cell responses in pancreatic cancer.
Fat-Soluble Vitamin Deficiencies and Disruption of the Immune System in Pancreatic Cancer: A Vicious Cycle.
Tumor-Associated Macrophages in Pancreatic Ductal Adenocarcinoma: Origin, Polarization, Function, and Reprogramming.
B7 Family Members in Pancreatic Ductal Adenocarcinoma: Attractive Targets for Cancer Immunotherapy.
[Progressin Tumor-Associated Macrophages in the Treatment of Pancreatic Cancer].
Immunotherapy in pancreatic ductal adenocarcinoma: an emerging entity?
Immune Checkpoint Inhibition for Pancreatic Ductal Adenocarcinoma: Current Limitations and Future Options.
DKK2 Impairs Tumor Immunity Infiltration and Correlates with Poor Prognosis in Pancreatic Ductal Adenocarcinoma.
Uncovering key targets of success for immunotherapy in pancreatic cancer.
CXCL12 in Pancreatic Cancer: Its Function and Potential as a Therapeutic Drug Target.
Challenges and Opportunities Associated With Platelets in Pancreatic Cancer.
Cadherin 11 Promotes Immunosuppression and Extracellular Matrix Deposition to Support Growth of Pancreatic Tumors and Resistance to Gemcitabine in Mice.
miR-128 Regulates Tumor Cell CD47 Expression and Promotes Anti-tumor Immunity in Pancreatic Cancer.
Immunotherapy and Combination Strategies in Pancreatic Cancer: Current Status and Emerging Trends.
Role of immune cells and immune-based therapies in pancreatitis and pancreatic ductal adenocarcinoma.
The diverse roles of circular RNAs in pancreatic cancer.
Stratification of Pancreatic Ductal Adenocarcinoma: Combinatorial Genetic, Stromal, and Immunologic Markers.
Functions and clinical implications of exosomes in pancreatic cancer.
Clinical Scenarios Emerging from Combined Immunophenotypic, Molecular and Morphologic Analysis of Pancreatic Cancer: The Good, the Bad and the Ugly Scenario.
Paracrine production of IL-6 promotes a hypercoagulable state in pancreatic cancer.
An integrated model of N6-methyladenosine regulators to predict tumor aggressiveness and immune evasion in pancreatic cancer.
New thoughts and findings on invasion and metastasis of pancreatic ductal adenocarcinoma (PDAC) from comparative proteomics: multi-target therapy.
Galectins Are Central Mediators of Immune Escape in Pancreatic Ductal Adenocarcinoma.
Major hurdles of immune-checkpoint inhibitors in pancreatic ductal adenocarcinoma.
Apoptosis and anergy of T cell induced by pancreatic stellate cells-derived galectin-1 in pancreatic cancer.

Orchestration of mesenchymal plasticity and immune evasiveness via rewiring of the metabolic program in pancreatic ductal adenocarcinoma.
CA9-Related Acidic Microenvironment Mediates CD8+ T Cell Related Immunosuppression in Pancreatic Cancer.
m(6)A-modified circRNA MYO1C participates in the tumor immune surveillance of pancreatic ductal adenocarcinoma through m(6)A/PD-L1 manner.
Coming in the Air: Hypoxia Meets Epigenetics in Pancreatic Cancer.
HCST Expression Distinguishes Immune-hot and Immune-cold Subtypes in Pancreatic Ductal Adenocarcinoma.
Tracing back to one of the origins of immune evasion in pancreatic cancer.
MicroRNA Expression Profiling of Pancreatic Cancer Cell Line L3.6p1 Following B7-H4 Knockdown.
New therapeutic targets in pancreatic cancer.
Oncolytic virotherapy for pancreatic ductal adenocarcinoma: A glimmer of hope after years of disappointment?
Tumor heterogeneity: An oncogenic driver of PDAC progression and therapy resistance under stress conditions.
Chemotherapy-Derived Inflammatory Responses Accelerate the Formation of Immunosuppressive Myeloid Cells in the Tissue Microenvironment of Human Pancreatic Cancer.
PDAC as an Immune Evasive Disease: Can 3D Model Systems Aid to Tackle This Clinical Problem?
Targeting the Metabolic Rewiring in Pancreatic Cancer and Its Tumor Microenvironment.
Immunotherapy for pancreatic cancer: present and future.
Immune Infiltration of CD8+ T Cells in Patients With Diabetic Pancreatic Cancer Reduces the Malignancy of Cancer Tissues: An In Silico Study.
A GATA6-centred gene regulatory network involving HNFs and $\Delta$ Np63 controls plasticity and immune escape in pancreatic cancer.
Toward stratification of patients with pancreatic cancer: Past lessons from traditional approaches and future applications with physical biomarkers.
VSIG2 promotes malignant progression of pancreatic ductal adenocarcinoma by enhancing LAMTOR2-mediated mTOR activation.
Connecting the Human Microbiome and Pancreatic Cancer.
A comprehensive survey into the role of exosomes in pancreatic cancer; from the origin of cancer to the progress and possibility of diagnosis and treatment.
MUC1 enhances tumor progression and contributes toward immunosuppression in a mouse model of spontaneous pancreatic adenocarcinoma.
Interactions between Cancer-Associated Fibroblasts and T Cells in the Pancreatic Tumor Microenvironment and the Role of Chemokines.
Autophagy is critical for cysteine metabolism in pancreatic cancer through regulation of SLC7A11.
Pancreatic Cancer Small Extracellular Vesicles (Exosomes): A Tale of Short- and Long-Distance Communication.
Pancreatic cancer: role of the immune system in cancer progression and vaccine-based immunotherapy.
Exploiting oxidative phosphorylation to promote the stem and immunoevasive properties of pancreatic cancer stem cells.
Crosstalk Between Peripheral Innervation and Pancreatic Ductal Adenocarcinoma.
A Promising Biomarker and Therapeutic Target in Patients with Advanced PDAC: The Stromal Protein $\beta$ ig-h3.
Regulation and function of autophagy in pancreatic cancer.

KRAS Mutation Dictates the Cancer Immune Environment in Pancreatic Ductal Adenocarcinoma and Other Adenocarcinomas.
The critical roles of activated stellate cells-mediated paracrine signaling, metabolism and onco-immunology in pancreatic ductal adenocarcinoma.
Elucidation of Tumor-Stromal Heterogeneity and the Ligand-Receptor Interactome by Single-Cell Transcriptomics in Real-world Pancreatic Cancer Biopsies.
AXL Inhibitor TP-0903 Reduces Metastasis and Therapy Resistance in Pancreatic Cancer.
Prognostic-related genes for pancreatic cancer typing and immunotherapy response prediction based on single-cell sequencing data and bulk sequencing data.
Urokinase-Type Plasminogen Activator Receptor (uPAR) Cooperates with Mutated KRAS in Regulating Cellular Plasticity and Gemcitabine Response in Pancreatic Adenocarcinomas.
DDX18 drives tumor immune escape through transcription-activated STAT1 expression in pancreatic cancer.
Cellular determinants and therapeutic implications of inflammation in pancreatic cancer.
An integrated model of acinar to ductal metaplasia-related N7-methyladenosine regulators predicts prognosis and immunotherapy in pancreatic carcinoma based on digital spatial profiling.
Expression of galectin-3 in pancreatic ductal adenocarcinoma.
B7-H4 is a prognostic biomarker for poor survival in patients with pancreatic cancer.
Quercetin improves pancreatic cancer chemo-sensitivity by regulating oxidative-inflammatory networks.
Combination treatment of advanced pancreatic cancer using novel vaccine and traditional therapies.
Sequestosome-1/p62-targeted small molecules for pancreatic cancer therapy.
DUSP2 recruits CSNK2A1 to suppress AKT1-mediated apoptosis resistance under hypoxic microenvironment in pancreatic cancer.
Targeting Pin1 renders pancreatic cancer eradicable by synergizing with immunochemotherapy.
Pancreatic stellate cell: Pandora's box for pancreatic disease biology.
Immunosurveillance of pancreatic adenocarcinoma: insights from genetically engineered mouse models of cancer.
New insights in the composition of extracellular vesicles from pancreatic cancer cells: implications for biomarkers and functions.
ENO1 promotes immunosuppression and tumor growth in pancreatic cancer.
Tumor cross-talk networks promote growth and support immune evasion in pancreatic cancer.
FAK suppresses antigen processing and presentation to promote immune evasion in pancreatic cancer.
Pancreatic Ductal Adenocarcinoma Cortical Mechanics and Clinical Implications.
Paracrine and cell autonomous signalling in pancreatic cancer progression and metastasis.
The CD155/TIGIT axis promotes and maintains immune evasion in neoantigen-expressing pancreatic cancer.
Identification of a subset of immunosuppressive P2RX1-negative neutrophils in pancreatic cancer liver metastasis.
Circ_0006790 carried by bone marrow mesenchymal stem cell-derived exosomes regulates S100A11 DNA methylation through binding to CBX7 in pancreatic ductal adenocarcinoma.
A personalized mRNA vaccine has exhibited potential in the treatment of pancreatic cancer.
The Hippo Signaling Pathway in Pancreatic Cancer.
Replication stress identifies novel molecular classification associated with treatment outcomes in pancreatic cancer.
Research on the epigenetic modification of pancreatic cancer vaccine.

PKC $\zeta$ regulates the expression of PDL1 through multiple pathways to modulate immune suppression of pancreatic cancer cells.
Prognostic effect of CD73 in pancreatic ductal adenocarcinoma for disease-free survival after radical surgery.
Targeting autophagy as a therapeutic strategy against pancreatic cancer.
Adenoviral gene therapy for pancreatic cancer: where do we stand?
Prognostic and diagnostic significance of galectins in pancreatic cancer: a systematic review and meta-analysis.
Defining the spatial distribution of extracellular adenosine revealed a myeloid-dependent immunosuppressive microenvironment in pancreatic ductal adenocarcinoma.
Functional reprogramming of peripheral blood monocytes by soluble mediators in patients with pancreatic cancer and intraductal papillary mucinous neoplasms.
CD73, a Promising Therapeutic Target of Diclofenac, Promotes Metastasis of Pancreatic Cancer through a Nucleotidase Independent Mechanism.
Prognostic Stratification Based on HIF-1 Signaling for Evaluating Hypoxic Status and Immune Infiltration in Pancreatic Ductal Adenocarcinomas.
Human Pancreatic Cancer Cells Induce a MyD88-Dependent Stromal Response to Promote a Tumor-Tolerant Immune Microenvironment.
TGF $\beta$ Signaling in the Pancreatic Tumor Microenvironment.
Combining Gemcitabine-Loaded Macrophage-like Nanoparticles and Erlotinib for Pancreatic Cancer Therapy.
CD13(hi) Neutrophil-like myeloid-derived suppressor cells exert immune suppression through Arginase 1 expression in pancreatic ductal adenocarcinoma.
The fatal alliance of cancer and T cells: How pancreatic tumor cells gather immunosuppressive T cells.
Progranulin mediates immune evasion of pancreatic ductal adenocarcinoma through regulation of MHCI expression.
[The Effects of PKC $\zeta$ on Anti-tumor Activity of Cytokine-induced Killer Cells Against Pancreatic Cancer Cells and the Possible Underlying Mechanisms].
Difluoromethylornithine Combined with a Polyamine Transport Inhibitor Is Effective against Gemcitabine Resistant Pancreatic Cancer.
Crosstalk of Sp1 and Stat3 signaling in pancreatic cancer pathogenesis.
The Bromodomain Inhibitor, INCB057643, Targets Both Cancer Cells and the Tumor Microenvironment in Two Preclinical Models of Pancreatic Cancer.
Strong expression of chemokine receptor CXCR4 by pancreatic cancer correlates with advanced disease.
Heterogeneous cancer-associated fibroblasts: A new perspective for understanding immunosuppression in pancreatic cancer.
PTEN/PI3K/mTOR/B7-H1 signaling pathway regulates cell progression and immuno-resistance in pancreatic cancer.
Divulging the Critical Role of HuR in Pancreatic Cancer as a Therapeutic Target and a Means to Overcome Chemoresistance.
STING agonist inflames the pancreatic cancer immune microenvironment and reduces tumor burden in mouse models.
Inhibition of DCLK1 kinase reverses epithelial-mesenchymal transition and restores T-cell activity in pancreatic ductal adenocarcinoma.
Transcription factor Sp1 is upregulated by PKC $\zeta$ to drive the expression of YAP1 during pancreatic carcinogenesis.

MYC- and MIZ1-Dependent Vesicular Transport of Double-Strand RNA Controls Immune Evasion in Pancreatic Ductal Adenocarcinoma.
STAT3-targeting RNA interference inhibits pancreatic cancer angiogenesis in vitro and in vivo.
Propionyl-CoA carboxylase subunit B modulates PIK3CA-regulated immune-surveillance in a pancreatic cancer mouse model.
Pancreatic stellate cells in pancreatic cancer: In focus.
A comprehensive analysis of different gene classes in pancreatic cancer: SIGLEC15 may be a promising immunotherapeutic target.
Aberrant Factors of Fibrinolysis and Coagulation in Pancreatic Cancer.
Epigenetically modified pancreatic carcinoma PANC-1 cells can act as cancer vaccine to enhance antitumor immune response in mice.
B7-H4 enhances oncogenicity and inhibits apoptosis in pancreatic cancer cells.
CD90 highly expressed population harbors a stemness signature and creates an immunosuppressive niche in pancreatic cancer.
Pancreatic Cancers with High Grade Tumor Budding Exhibit Hallmarks of Diminished Anti-Tumor Immunity.
T cell adhesion and cytolysis of pancreatic cancer cells: a role for E-cadherin in immunotherapy?
Extracellular Galectin 4 Drives Immune Evasion and Promotes T-cell Apoptosis in Pancreatic Cancer.
Tumor-associated MUC5AC stimulates in vivo tumorigenicity of human pancreatic cancer.
Downstream mediators of the intratumoral interferon response suppress antitumor immunity, induce gemcitabine resistance and associate with poor survival in human pancreatic cancer.
It Takes Two to Tango: Potential Prognostic Impact of Circulating TGF-Beta and PD-L1 in Pancreatic Cancer.
Identification of LIPH as an unfavorable biomarkers correlated with immune suppression or evasion in pancreatic cancer based on RNA-seq.
[Role of B7-H1 in pancreatic carcinoma immune evasion].
The actin modulator hMENA regulates GAS6-AXL axis and pro-tumor cancer/stromal cell cooperation.
Mass spectrum analysis of membrane proteins reveals that CASK, CD36 and EPB42 are differentially expressed in pancreatic adenocarcinoma.
The CD40-CD154 interaction would correlate with proliferation and immune escape in pancreatic ductal adenocarcinoma.
Chemokine-mucinome interplay in shaping the heterogeneous tumor microenvironment of pancreatic cancer.
Anti-VEGF treatment-resistant pancreatic cancers secrete proinflammatory factors that contribute to malignant progression by inducing an EMT cell phenotype.
Target proteomic profiling of frozen pancreatic CD24+ adenocarcinoma tissues by immuno-laser capture microdissection and nano-LC-MS/MS.
Extracellular vesicles in pancreatic cancer immune escape: Emerging roles and mechanisms.
Ion Channels Orchestrate Pancreatic Ductal Adenocarcinoma Progression and Therapy.
Inhibition of mutant KRAS-driven overexpression of ARF6 and MYC by an eIF4A inhibitor drug improves the effects of anti-PD-1 immunotherapy for pancreatic cancer.
Adipocytes and Neutrophils Give a Helping Hand to Pancreatic Cancers.
Serum miR-338-3p and miR-199b-5p are associated with the absolute neutrophil count in patients with resectable pancreatic cancer.
Systemic and local immunosuppression in pancreatic cancer patients.
Macropinocytosis of Nab-paclitaxel Drives Macrophage Activation in Pancreatic Cancer.
Targeting Galectin-1 in pancreatic cancer: immune surveillance on guard.

Cellular Membrane Localization of Innate Immune Checkpoint Molecule CD47 Is Regulated by Radixin in Human Pancreatic Ductal Adenocarcinoma Cells.
The Role of the Microbiome in Immunologic Development and its Implication For Pancreatic Cancer Immunotherapy.
Clinical significance of B7-H1 and B7-1 expressions in pancreatic carcinoma.
Small extracellular vesicle non-coding RNAs in pancreatic cancer: molecular mechanisms and clinical implications.
Par-4 mediated Smad4 induction in PDAC cells restores canonical TGF- $\beta$ / Smad4 axis driving the cells towards lethal EMT.
ARF6 and AMAP1 are major targets of KRAS and TP53 mutations to promote invasion, PD-L1 dynamics, and immune evasion of pancreatic cancer.
A bioinformatics analysis, pre-clinical and clinical conception of autophagy in pancreatic cancer: Complexity and simplicity in crosstalk.
Transcriptional profiling of peripheral blood mononuclear cells in pancreatic cancer patients identifies novel genes with potential diagnostic utility.
Polarization of Cancer-Associated Macrophages Maneuver Neoplastic Attributes of Pancreatic Ductal Adenocarcinoma.
[High serum levels of Transforming Growth Factor-beta1, Interleukin-10 and Vascular Endothelial Growth Factor in pancreatic adenocarcinoma patients].
Pancreatic adenocarcinoma cell lines show variable susceptibility to TRAIL-mediated cell death.
Circ_0058058 Drives the Malignant Phenotypes and Immune Evasion of Pancreatic Cancer by the MicroRNA-557-Dependent Regulation of PDL1.
Monoclonal Antibody Targeting Sialyl-di-Lewis(a)-Containing Internalizing and Noninternalizing Glycoproteins with Cancer Immunotherapy Development Potential.
Genotyping and expression analysis of IDO2 in human pancreatic cancer: a novel, active target.
The trilogy of P21 activated kinase, autophagy and immune evasion in pancreatic ductal adenocarcinoma.
Pancreatic Cancer Stem-Like Cells With High Calreticulin Expression Associated With Immune Surveillance.
Inflammation Promotes Progression of Pancreatic Cancer Through WNT/ $\beta$ -Catenin Pathway-Dependent Manner.
Expression of the Novel Costimulatory Molecule B7-H5 in Pancreatic Cancer.
The anthelmintic drug niclosamide induces GSK- $\beta$ -mediated $\beta$ -catenin degradation to potentiate gemcitabine activity, reduce immune evasion ability and suppress pancreatic cancer progression.
Mesothelial cell-derived antigen-presenting cancer-associated fibroblasts induce expansion of regulatory T cells in pancreatic cancer.
Immunologic and Metabolic Features of Pancreatic Ductal Adenocarcinoma Define Prognostic Subtypes of Disease.
STAT3 inhibition enhances gemcitabine sensitivity in pancreatic cancer by suppressing EMT, immune escape and inducing oxidative stress damage.
Integrated Analysis of Long Non-Coding RNA and mRNA Expression Profile in Pancreatic Cancer Derived Exosomes Treated Dendritic Cells by Microarray Analysis.
Hypoxia promotes immune escape of pancreatic cancer cells by lncRNA NNT-AS1/METTL3-HuR-mediated ITGB1 m(6)A modification.
B7-H1 up-regulated expression in human pancreatic carcinoma tissue associates with tumor progression.
The Role of Adipokines in Pancreatic Cancer.

CD73 acts as a prognostic biomarker and promotes progression and immune escape in pancreatic cancer.
Exploring the Role of Galectins in Cancer : In Vitro and In Vivo Approaches.
B3GNT3 overexpression promotes tumor progression and inhibits infiltration of CD8(+) T cells in pancreatic cancer.
Pancreatic cancer cell-derived microRNA-155-5p-containing extracellular vesicles promote immune evasion by triggering EHF-dependent activation of Akt/NF- $\kappa$ B signaling pathway.
Genome-wide CRISPR screens define determinants of epithelial-mesenchymal transition mediated immune evasion by pancreatic cancer cells.
Unresolved endoplasmic reticulum stress engenders immune-resistant, latent pancreatic cancer metastases.
Pancreatic cancer escape variants that evade immunogene therapy through loss of sensitivity to IFN $\gamma$ -induced apoptosis.
PDIA6 promotes pancreatic cancer progression and immune escape through CSN5-mediated deubiquitination of $\beta$ -catenin and PD-L1.
Discovery Proteomics Analysis Determines That Driver Oncogenes Suppress Antiviral Defense Pathways Through Reduction in Interferon- $\beta$ Autocrine Stimulation.
Microbiome Dysbiosis and Predominant Bacterial Species as Human Cancer Biomarkers.
KRAS, MYC, and ARF6: inseparable relationships cooperatively promote cancer malignancy and immune evasion.
Localization and upregulation of survivin in cancer health disparities: a clinical perspective.
Immune sculpting of norepinephrine on MHC-I, B7-1, IDO and B7-H1 expression and regulation of proliferation and invasion in pancreatic carcinoma cells.

Table A2-19, Cluster 18

Cluster 18 focuses on 1) the role of exosomes in cancer progression (especially their role in tumor microenvironment therapy resistance), and 2) immunotherapies to reduce the tolerance associated with the immunosuppressive tumor microenvironment (297)
Trends in Research on Exosomes in Cancer Progression and Anticancer Therapy.
Exosomes carrying immune checkpoints, a promising therapeutic approach in cancer treatment.
Focus on exosomes: novel pathogenic components of leukemia.
The Immunomodulation Potential of Exosomes in Tumor Microenvironment.
Exosomes in cancer immunoediting and immunotherapy.
The exosomes in tumor immunity.
Exosomes and the extracellular matrix: a dynamic interplay in cancer progression.
Emerging role of exosomes in cancer progression and tumor microenvironment remodeling.
Exosomes and Their Role in Cancer Progression.
Urinary Exosomes: The Potential for Biomarker Utility, Intercellular Signaling and Therapeutics in Urological Malignancy.
Emerging Function and Clinical Values of Exosomal MicroRNAs in Cancer.
Camouflage strategies for therapeutic exosomes evasion from phagocytosis.
The cancer exosomes: Clinical implications, applications and challenges.
Exosomes in cancer: Small transporters with big functions.
In Situ Capturing and Counting Device for the Specific Depletion and Purification of Cancer-Derived Exosomes.



Role of exosomes in the immune microenvironment of ovarian cancer.
The biological functions and clinical applications of exosomes in lung cancer.
Tumor-derived exosomes in hypoxic microenvironment: release mechanism, biological function and clinical application.
Exosome-derived noncoding RNAs in gastric cancer: functions and clinical applications.
Role of exosomes in immune regulation.
Functions of Cancer-Derived Extracellular Vesicles in Immunosuppression.
Role of exosomes in treatment of hepatocellular carcinoma.
The significance of exosomal RNAs in the development, diagnosis, and treatment of pancreatic cancer.
Role of exosomes in tumour growth, chemoresistance and immunity: state-of-the-art.
Exosomes: an overview of biogenesis, composition and role in ovarian cancer.
Exosomal communication goes viral.
Exosome: From leukemia progression to a novel therapeutic approach in leukemia treatment.
Role of exosomes in tumour and transplant immune regulation.
Exosomes: A Potential Therapeutic Tool Targeting Communications between Tumor Cells and Macrophages.
Dynamic Role of Exosome microRNAs in Cancer Cell Signaling and Their Emerging Role as Noninvasive Biomarkers.
The Crosstalk between Tumor Cells and the Microenvironment in Hepatocellular Carcinoma: The Role of Exosomal microRNAs and their Clinical Implications.
The functional role of exosome in hepatocellular carcinoma.
Diagnostic and Therapeutic Potential of Exosomes in Cancer: The Beginning of a New Tale?
Implications of exosomes as diagnostic and therapeutic strategies in cancer.
Exosomes and organ-specific metastasis.
Exosomes in the Oncobiology, Diagnosis, and Therapy of Hepatic Carcinoma: A New Player of an Old Game.
Cancer-derived exosomal microRNAs shape the immune system within the tumor microenvironment: State of the art.
The role of exosomes in hepatitis, liver cirrhosis and hepatocellular carcinoma.
Exosomes in cancer theranostic: Diamonds in the rough.
Exosomes in gastric cancer: roles, mechanisms, and applications.
[Research Advance of Exosomes in the Immune Microenvironment of Hematological Malignancies--Review].
Role of exosomes and exosomal microRNAs in hepatocellular carcinoma: Potential in diagnosis and antitumour treatments (Review).
Salivary exosomes as potential biomarkers in cancer.
Tumor-derived exosomes in ovarian cancer - liquid biopsies for early detection and real-time monitoring of cancer progression.
Non-coding RNAs shuttled via exosomes reshape the hypoxic tumor microenvironment.
Exosomes: A rising star in breast cancer (Review).
The Potential Roles of Exosomal Non-Coding RNAs in Hepatocellular Carcinoma.
The multifaceted involvement of exosomes in tumor progression: Induction and inhibition.
Tumor-derived exosomes promote tumor progression and T-cell dysfunction through the regulation of enriched exosomal microRNAs in human nasopharyngeal carcinoma.
Exosome-Derived microRNA: Implications in Melanoma Progression, Diagnosis and Treatment.

Tumour-derived exosomes: Tiny envelopes for big stories.
Exosomal miRNAs in tumor microenvironment.
Exosome secretion from hypoxic cancer cells reshapes the tumor microenvironment and mediates drug resistance.
Cancer associated-fibroblast-derived exosomes in cancer progression.
Communication in tiny packages: Exosomes as means of tumor-stroma communication.
The Effect of Hypoxia-Induced Exosomes on Anti-Tumor Immunity and Its Implication for Immunotherapy.
Formation and role of exosomes in cancer.
Mechanism and clinical value of exosomes and exosomal contents in regulating solid tumor radiosensitivity.
Tumor-derived exosomes: the emerging orchestrators in melanoma.
Exosomal circGSE1 promotes immune escape of hepatocellular carcinoma by inducing the expansion of regulatory T cells.
The role of exosomes in lung cancer metastasis and clinical applications: an updated review.
Exosomes in the hypoxic TME: from release, uptake and biofunctions to clinical applications.
Exosomes isolation and characterization in serum is feasible in non-small cell lung cancer patients: critical analysis of evidence and potential role in clinical practice.
Exosomes derived from cancerous and non-cancerous cells regulate the anti-tumor response in the tumor microenvironment.
Exosomes in Immune Regulation.
Exosomes in Cardiovascular Medicine.
Immune Cell-Derived Exosomes in the Cancer-Immunity Cycle.
Human acute myeloid leukemia blast-derived exosomes in patient-derived xenograft mice mediate immune suppression.
Exosomes: New players in cancer (Review).
Role of exosomes in hepatocellular carcinoma and the regulation of traditional Chinese medicine.
MicroRNAs in Tumor Exosomes Drive Immune Escape in Melanoma.
Modulation of tissue tropism and biological activity of exosomes and other extracellular vesicles: New nanotools for cancer treatment.
Hepatocellular carcinoma-derived exosomes in organotropic metastasis, recurrence and early diagnosis application.
Extracellular Vesicles in Oncology: from Immune Suppression to Immunotherapy.
Advances in exosome biomarkers for cervical cancer.
New insights into exosome mediated tumor-immune escape: Clinical perspectives and therapeutic strategies.
Post-translational modifications of exosomal proteins.
Transcriptome reprogramming by cancer exosomes: identification of novel molecular targets in matrix and immune modulation.
Exosomes and the MICA-NKG2D system in cancer.
Tumor exosomes expressing Fas ligand mediate CD8+ T-cell apoptosis.
Exosomes carrying immunoinhibitory proteins and their role in cancer.
Human tumor virus utilizes exosomes for intercellular communication.
Exosome removal as a therapeutic adjuvant in cancer.
Unveiling the Yin-Yang Balance of M1 and M2 Macrophages in Hepatocellular Carcinoma: Role of Exosomes in Tumor Microenvironment and Immune Modulation.

Exosomes: small vesicles participating in intercellular communication.
Emerging Evidence for the Clinical Relevance of Pancreatic Cancer Exosomes.
Exosomes-mediated crosstalk between glioma and immune cells in the tumor microenvironment.
Exosome-Derived microRNA: Efficacy in Cancer.
ExomiRs: A Novel Strategy in Cancer Diagnosis and Therapy.
HCC-Derived Exosomes: Critical Player and Target for Cancer Immune Escape.
Emerging Functions and Clinical Applications of Exosomal ncRNAs in Ovarian Cancer.
Tumor-derived exosomes confer antigen-specific immunosuppression in a murine delayed-type hypersensitivity model.
Exosomal CD47 Plays an Essential Role in Immune Evasion in Ovarian Cancer.
Cancer cell's internal and external warriors: Autophagosomes and exosomes.
Tumour-derived exosomes and their emerging roles in leukaemia (Review).
A Clinician's Guide to Cancer-Derived Exosomes: Immune Interactions and Therapeutic Implications.
Melanoma cell-derived exosomes alter macrophage and dendritic cell functions in vitro.
Rapid and comprehensive 'shotgun' lipidome profiling of colorectal cancer cell derived exosomes.
Regulation of Antitumor Immune Responses by Exosomes Derived from Tumor and Immune Cells.
A Preliminary Proteomic Investigation of Circulating Exosomes and Discovery of Biomarkers Associated with the Progression of Osteosarcoma in a Clinical Model of Spontaneous Disease.
Redox-signaling transmitted in trans to neighboring cells by melanoma-derived TNF-containing exosomes.
Exosomal Cirsaf2 Reshaping Tumor Environment to Promote Renal Cell Carcinoma Progression by Mediating M2 Macrophage Polarization.
The functional roles of exosomes-derived long non-coding RNA in human cancer.
The Double Face of Exosome-Carried MicroRNAs in Cancer Immunomodulation.
Exosomal noncoding RNAs in Glioma: biological functions and potential clinical applications.
The opportunistic effect of exosomes on Non-Hodgkin Lymphoma microenvironment modulation.
Exosomal miRNA in chemoresistance, immune evasion, metastasis and progression of cancer.
Exosomes in Cancer: Another Mechanism of Tumor-Induced Immune Suppression.
Emerging Roles of Exosomal Circular RNAs in Cancer.
The Role of Exo-miRNAs in Cancer: A Focus on Therapeutic and Diagnostic Applications.
Exosomes as intercellular signalosomes and pharmacological effectors.
Plasma-derived exosomal survivin, a plausible biomarker for early detection of prostate cancer.
Exosomal Non-Coding RNAs: Novel Regulators of Macrophage-Linked Intercellular Communication in Lung Cancer and Inflammatory Lung Diseases.
Exosome signaling in mammary gland development and cancer.
The roles of tumor-derived exosomes in altered differentiation, maturation and function of dendritic cells.
Exosomal vesicles enhance immunosuppression in chronic inflammation: Impact in cellular senescence and the aging process.
A Novel Exosome-Relevant Molecular Classification Uncovers Distinct Immune Escape Mechanisms and Genomic Alterations in Gastric Cancer.
Tumor-derived exosomes as mediators of disease and potential diagnostic biomarkers.
Oncogenic Role of Exosomal Circular and Long Noncoding RNAs in Gastrointestinal Cancers.
Exosomal noncoding RNA-mediated spatiotemporal regulation of lipid metabolism: Implications in immune evasion and chronic inflammation.

Exosomal circRNA-001264 promotes AML immunosuppression through induction of M2-like macrophages and PD-L1 overexpression.
Recent advances of exosomal circRNAs in cancer and their potential clinical applications.
The importance of exosomal PDL1 in tumour immune evasion.
Tumor cell cross talk with tumor-associated leukocytes leads to induction of tumor exosomal fibronectin and promotes tumor progression.
Roles of exosome-derived non-coding RNA in tumor micro-environment and its clinical application.
The potential of tumor-derived exosomes for noninvasive cancer monitoring.
Extracellular Vesicles: New Classification and Tumor Immunosuppression.
Exosomal circRNA: emerging insights into cancer progression and clinical application potential.
Exosomes as a tumor immune escape mechanism: possible therapeutic implications.
Exosomes as Efficient Nanocarriers in Osteosarcoma: Biological Functions and Potential Clinical Applications.
Pancreatic cancer-derived exosomes induce apoptosis of T lymphocytes through the p38 MAPK-mediated endoplasmic reticulum stress.
Exosome-derived microRNAs in cancer metabolism: possible implications in cancer diagnostics and therapy.
Extracellular Vesicles and Their Roles in the Tumor Immune Microenvironment.
Extracellular Vesicles: New Players in Lung Immunity.
The impact of hypoxia on extracellular vesicle secretome profile of cancer.
Comprehensive microRNA-sequencing of exosomes derived from head and neck carcinoma cells in vitro reveals common secretion profiles and potential utility as salivary biomarkers.
Contribution of proteomics to understanding the role of tumor-derived exosomes in cancer progression: state of the art and new perspectives.
Glioma-Derived Extracellular Vesicles - Far More Than Local Mediators.
Tumor-derived exosomes (TDEs): How to avoid the sting in the tail.
The Role of Extracellular Vesicles in the Hallmarks of Cancer and Drug Resistance.
MicroRNAs in extracellular vesicles: potential cancer biomarkers.
Role of exosomes in transferring chemoresistance through modulation of cancer glycolytic cell metabolism.
The Uptake of Extracellular Vesicles is Affected by the Differentiation Status of Myeloid Cells.
Effects of exosomes on pre-metastatic niche formation in tumors.
Tumor exosomes block dendritic cells maturation to decrease the T cell immune response.
Systemic T Cells Immunosuppression of Glioma Stem Cell-Derived Exosomes Is Mediated by Monocytic Myeloid-Derived Suppressor Cells.
Pioneer Role of Extracellular Vesicles as Modulators of Cancer Initiation in Progression, Drug Therapy, and Vaccine Prospects.
Pancreatic cancer, stroma, and exosomes.
Extracellular Vesicles for Cancer Therapy: Impact of Host Immune Response.
Exosomal miR-940 maintains SRC-mediated oncogenic activity in cancer cells: a possible role for exosomal disposal of tumor suppressor miRNAs.
Tumor-associated exosomes promote lung cancer metastasis through multiple mechanisms.
Tumor-Derived Extracellular Vesicles: Multifunctional Entities in the Tumor Microenvironment.
Exosomes derived from tumor cells genetically modified to express Mycobacterium tuberculosis antigen: a novel vaccine for cancer therapy.
Exosomes: Mediators in microenvironment of colorectal cancer.

Glycosyl-Phosphatidyl-Inositol (GPI)-Anchors and Metalloproteases: Their Roles in the Regulation of Exosome Composition and NKG2D-Mediated Immune Recognition.
Extracellular Vesicles (EVs) in Tumor Diagnosis and Therapy.
Exosomal B7-H4 from irradiated glioblastoma cells contributes to increase FoxP3 expression of differentiating Th1 cells and promotes tumor growth.
Exosomal cargos-mediated metabolic reprogramming in tumor microenvironment.
Extracellular Vesicles in Cancer Drug Resistance: Roles, Mechanisms, and Implications.
The Role of Extracellular Vesicles in Disease Progression and Detection of Hepatocellular Carcinoma.
Purine Metabolites in Tumor-Derived Exosomes May Facilitate Immune Escape of Head and Neck Squamous Cell Carcinoma.
Comparative Gene Expression Analysis of Lymphocytes Treated with Exosomes Derived from Ovarian Cancer and Ovarian Cysts.
Extracellular Vesicles As miRNA Nano-Shuttles: Dual Role in Tumor Progression.
Blood diffusion and Th1-suppressive effects of galectin-9-containing exosomes released by Epstein-Barr virus-infected nasopharyngeal carcinoma cells.
Extracellular Vesicles Are Important Mediators That Regulate Tumor Lymph Node Metastasis via the Immune System.
Non-coding RNA derived from extracellular vesicles in cancer immune escape: Biological functions and potential clinical applications.
Ovarian-Cancer-Associated Extracellular Vesicles: Microenvironmental Regulation and Potential Clinical Applications.
Investigating the tumor-immune microenvironment through extracellular vesicles from frozen patient biopsies and 3D cultures.
Exosomal evasion of humoral immunotherapy in aggressive B-cell lymphoma modulated by ATP-binding cassette transporter A3.
Signaling by Extracellular Vesicles Advances Cancer Hallmarks.
Extracellular Vesicles in Chemoresistance.
Advances in the Biological Functions of Extracellular Vesicles and their Potential Use in Treating Oral Cancer.
Micromanaging of tumor metastasis by extracellular vesicles.
Cancer-secreted exosomal miR-1468-5p promotes tumor immune escape via the immunosuppressive reprogramming of lymphatic vessels.
Exosomal circRNAs in cancer: Implications for therapy resistance and biomarkers.
Melanoma-Derived Exosomal miR-125b-5p Educates Tumor Associated Macrophages (TAMs) by Targeting Lysosomal Acid Lipase A (LIPA).
Methodological Approaches to Study Extracellular Vesicle miRNAs in Epstein-Barr Virus-Associated Cancers.
Extracellular Vesicles in Renal Cell Carcinoma: Multifaceted Roles and Potential Applications Identified by Experimental and Computational Methods.
Exosomal communication by metastatic osteosarcoma cells modulates alveolar macrophages to an M2 tumor-promoting phenotype and inhibits tumoricidal functions.
Extracellular vesicles in ovarian cancer chemoresistance, metastasis, and immune evasion.
The forces driving cancer extracellular vesicle secretion.
The emergence of extracellular vesicles in urology: fertility, cancer, biomarkers and targeted pharmacotherapy.
Extracellular vesicles in onco-nephrology.

OncomiRs as noncoding RNAs having functions in cancer: Their role in immune suppression and clinical implications.
Extracellular vesicles secreted from cancer cell lines stimulate secretion of MMP-9, IL-6, TGF- $\beta$ 1 and EMMPRIN.
Role of Extracellular Vesicles in Epithelial Ovarian Cancer: A Systematic Review.
The Role and Clinical Interest of Extracellular Vesicles in Pregnancy and Ovarian Cancer.
Exosomes: small vesicles with big roles in hepatocellular carcinoma.
Deciphering the messages carried by extracellular vesicles in hematological malignancies.
Prognostic and immunological role of cancer-associated fibroblasts-derived exosomal protein in esophageal squamous cell carcinoma.
Extracellular vesicles in immunomodulation and tumor progression.
Tumor-derived exosomes modulate T cell function through transfer of RNA.
Extracellular Vesicles and Carried miRNAs in the Progression of Renal Cell Carcinoma.
Extracellular Vesicles-Based Cell-Cell Communication in Melanoma: New Perspectives in Diagnostics and Therapy.
Tumor-derived extracellular vesicle nucleic acids as promising diagnostic biomarkers for prostate cancer.
Bioengineered exosomal-membrane-camouflaged abiotic nanocarriers: neurodegenerative diseases, tissue engineering and regenerative medicine.
Extra-cellular vesicles carry proteome of cancer hallmarks.
The Purification and Characterization of Exosomes from Macrophages.
Exosomal B7-H3 facilitates colorectal cancer angiogenesis and metastasis through AKT1/mTOR/VEGFA pathway.
Urinary Biomarkers in Bladder Cancer: Where Do We Stand and Potential Role of Extracellular Vesicles.
Extracellular Vesicles and Hepatocellular Carcinoma: Opportunities and Challenges.
Alteration of cellular and immune-related properties of bone marrow mesenchymal stem cells and macrophages by K562 chronic myeloid leukemia cell derived exosomes.
Extracellular Vesicle Nanoarchitectonics for Novel Drug Delivery Applications.
TAM-derived extracellular vesicles containing microRNA-29a-3p explain the deterioration of ovarian cancer.
The Role of Extracellular Vesicles in Cancer: Cargo, Function, and Therapeutic Implications.
Analysis of Extracellular Vesicles in Gastric Juice from Gastric Cancer Patients.
Proteomic characterization of circulating extracellular vesicles identifies novel serum myeloma associated markers.
The roles of extracellular vesicles in gastric cancer development, microenvironment, anti-cancer drug resistance, and therapy.
Extracellular Vesicles and Transforming Growth Factor $\beta$ Signaling in Cancer.
Emerging role of exosome-derived non-coding RNAs in tumor-associated angiogenesis of tumor microenvironment.
M2 macrophage-derived exosomal microRNA-155-5p promotes the immune escape of colon cancer by downregulating ZC3H12B.
High-throughput analysis and functional interpretation of extracellular vesicle content in hematological malignancies.
The roles of extracellular vesicles in the development, microenvironment, anticancer drug resistance, and therapy of head and neck squamous cell carcinoma.
The biology and function of extracellular vesicles in nasopharyngeal carcinoma (Review).

EV-Mediated Chemoresistance in the Tumor Microenvironment: Is NF- $\kappa$ B a Player?
Extracellular Vesicle-Mediated Mitochondrial Reprogramming in Cancer.
Targeting nucleic acid sensors in tumor cells to reprogram biogenesis and RNA cargo of extracellular vesicles for T cell-mediated cancer immunotherapy.
Role of exosomal miR-21 in the tumor microenvironment and osteosarcoma tumorigenesis and progression (Review).
Melanoma exosomes enable tumor tolerance in lymph nodes.
Tumor-Associated Macrophage-Derived Exosomes Promote the Progression of Gastric Cancer by Regulating the P38MAPK Signaling Pathway and the Immune Checkpoint PD-L1.
Biological roles and potential applications of immune cell-derived extracellular vesicles.
The Power of Extracellular Vesicles in Myeloproliferative Neoplasms: "Crafting" a Microenvironment That Matters.
Extracellular vesicles as modulators of the cancer microenvironment.
Cancer exosome-derived miR-9 and miR-181a promote the development of early-stage MDSCs via interfering with SOCS3 and PIAS3 respectively in breast cancer.
Exosome-Mediated Metastasis: From Epithelial-Mesenchymal Transition to Escape from Immunosurveillance.
Extracellular Vesicles Enhance Multiple Myeloma Metastatic Dissemination.
Metastatic Dissemination: Role of Tumor-Derived Extracellular Vesicles and Their Use as Clinical Biomarkers.
The "Vesicular Intelligence" Strategy of Blood Cancers.
The biology of extracellular microvesicles.
Anti-HER2 scFv-Directed Extracellular Vesicle-Mediated mRNA-Based Gene Delivery Inhibits Growth of HER2-Positive Human Breast Tumor Xenografts by Prodrug Activation.
Extracellular Vesicles from Ocular Melanoma Have Pro-Fibrotic and Pro-Angiogenic Properties on the Tumor Microenvironment.
[Mannose-capped lipoarabinomannan (ManLAM) binding TLR2 activates mast cells to release exosomes and induces M2 polarization of macrophages].
Treatment Failure in Acute Myeloid Leukemia: Focus on the Role of Extracellular Vesicles.
Role of extracellular vesicles in osteosarcoma.
Urinary exosomes-based Engineered Nanovectors for Homologously Targeted Chemo-Chemodynamic Prostate Cancer Therapy via abrogating EGFR/AKT/NF- $\kappa$ B/I $\kappa$ B signaling.
Proteomic profiling and functional characterization of serum-derived extracellular vesicles in the mucinous and non-mucinous colon adenocarcinoma.
Adenosine Methylation Level of miR-125a-5p Promotes Anti-PD-1 Therapy Escape through the Regulation of IGSF11/VSIG3 Expression.
MicroRNAs, Regulatory Messengers Inside and Outside Cancer Cells.
Malignant ascite-derived extracellular vesicles inhibit T cell activity by upregulating Siglec-10 expression.
[The action mechanism of glioblastoma cell-derived exosome: a review].
MicroRNA-15a Carried by Mesenchymal Stem Cell-Derived Extracellular Vesicles Inhibits the Immune Evasion of Colorectal Cancer Cells by Regulating the KDM4B/HOXC4/PD-L1 Axis.
Comparing extracellular vesicles and cell membranes as biocompatible coatings for gold nanorods: Implications for targeted theranostics.
MicroRNA in lung cancer: role, mechanisms, pathways and therapeutic relevance.
Extracellular Vesicles: New Players in Lymphomas.
miRNAs in the Regulation of Cancer Immune Response: Effect of miRNAs on Cancer Immunotherapy.

Mesenchymal stem cells shuttling miR-503 via extracellular vesicles enhance glioma immune escape.
Exosomal NAMPT from chronic lymphocytic leukemia cells orchestrate monocyte survival and phenotype under endoplasmic reticulum stress.
Non-Coding RNAs Delivery by Small Extracellular Vesicles and Their Applications in Ovarian Cancer.
Modulation of cancer traits by tumor suppressor microRNAs.
Extracellular vesicles-derived microRNA-222 promotes immune escape via interacting with ATF3 to regulate AKT1 transcription in colorectal cancer.
Targeted heart repair by Tβ4-loaded cardiac-resident macrophage-derived extracellular vesicles modified with monocyte membranes.
Recent Advances in Cell-Based Nanotherapy for Cardiovascular Diseases.
Extracellular Vesicle lncRNA Metastasis-Associated Lung Adenocarcinoma Transcript 1 Released From Glioma Stem Cells Modulates the Inflammatory Response of Microglia After Lipopolysaccharide Stimulation Through Regulating miR-129-5p/High Mobility Group Box-1 Protein Axis.
Tumor-derived exosomes encapsulating miR-34a promote apoptosis and inhibit migration and tumor progression of colorectal cancer cells under in vitro condition.
GATA3 Encapsulated by Tumor-Associated Macrophage-Derived Extracellular Vesicles Promotes Immune Escape and Chemotherapy Resistance of Ovarian Cancer Cells by Upregulating the CD24/Siglec-10 Axis.
Circulating mir-320a promotes immunosuppressive macrophages M2 phenotype associated with lung cancer risk.
Small but mighty: microparticles as mediators of tumor progression.
Novel microRNAs modulating ecto-5'-nucleotidase expression.
Noncoding RNAs in Glioblastoma.
M2 macrophage-derived extracellular vesicles augment immune evasion and development of colorectal cancer via a circRNA_CCDC66/microRNA-342-3p/metadherin axis.
Non-Coding RNAs in Gastric Cancer: From Malignant Hallmarks to Clinical Applications.
Cancer Hallmarks and MicroRNAs: The Therapeutic Connection.
Roles of CD133 in microvesicle formation and oncoprotein trafficking in colon cancer.
MicroRNAs in cancer: lessons from melanoma.
Role of microRNAs in programmed cell death in renal diseases: A review.
Clinical significance of microRNAs in chronic and acute human leukemia.
Comparative efficacy and mechanism of action of cardiac progenitor cells after cardiac injury.
Focused Ultrasound Hyperthermia Augments Release of Glioma-derived Extracellular Vesicles with Differential Immunomodulatory Capacity.
Localization of the Epstein-Barr virus protein LMP 1 to exosomes.
MicroRNAs induced in melanoma treated with combination targeted therapy of Temsirolimus and Bevacizumab.
The role of microRNAs and long non-coding RNAs in the pathology, diagnosis, and management of melanoma.
Recent advances of non-coding RNAs in ovarian cancer prognosis and therapeutics.
Non-coding RNAs targeting notch signaling pathway in cancer: From proliferation to cancer therapy resistance.
Interaction of microRNAs with sphingosine kinases, sphingosine-1 phosphate, and sphingosine-1 phosphate receptors in cancer.
Double Insurance for OC: miRNA-Mediated Platinum Resistance and Immune Escape.
Identification of Immune Modulatory miRNAs by miRNA Enrichment via RNA Affinity Purification.
The biology and clinical potential of circulating tumor cells.



The emerging roles of the polycistronic miR-106b~25 cluster in cancer - A comprehensive review.
Current concepts of non-coding RNA regulation of immune checkpoints in cancer.
EBV miRNAs are potent effectors of tumor cell transcriptome remodeling in promoting immune escape.
Regulation of cancer immune escape: The roles of miRNAs in immune checkpoint proteins.
The emerging role of Epstein-Barr virus encoded microRNAs in nasopharyngeal carcinoma.
Modulation of vesicle shedding in 8701 BC human breast carcinoma cells.
Altered expression of miR-181 affects cell fate and targets drug resistance-related mechanisms.
Cytokine Profiling in Low- and High-Density Small Extracellular Vesicles from Epidermoid Carcinoma Cells.
Diverse Roles and Therapeutic Potentials of Circular RNAs in Urological Cancers.
Profiling of microRNAs modulating cytomegalovirus infection in astrocytoma patients.
Erratum: Mechanisms of extracellular vesicle-mediated immune evasion in melanoma.
Novel insights into the roles and therapeutic implications of MUC1 oncoprotein via regulating proteins and non-coding RNAs in cancer.
The Response of microRNAs to Solar UVR in Skin-Resident Melanocytes Differs between Melanoma Patients and Healthy Persons.
Construction of Exosomes that Overexpress CD47 and Evaluation of Their Immune Escape.
Gold nano-particles (AuNPs) carrying anti-EBV-miR-BART7-3p inhibit growth of EBV-positive nasopharyngeal carcinoma.
P53-regulated miR-320a targets PDL1 and is downregulated in malignant mesothelioma.
Circulating nucleic acids: An analysis of their occurrence in malignancies.
Corrigendum: Long Non-Coding RNAs in the Tumor Immune Microenvironment: Biological Properties and Therapeutic Potential.
The Role of Nucleases Cleaving TLR3, TLR7/8 and TLR9 Ligands, Dicer RNase and miRNA/piRNA Proteins in Functional Adaptation to the Immune Escape and Xenophagy of Prostate Cancer Tissue.
A systematic CRISPR screen reveals an IL-20/IL20RA-mediated immune crosstalk to prevent the ovarian cancer metastasis.

Table A2-20, Cluster 19

Cluster 19 focuses on Natural Killer Cell therapies against tumors, emphasizing reduction of immune evasion through enhancing activation and surveillance functions of NK cells (628)
Underlying mechanisms of evasion from NK cells as rationale for improvement of NK cell-based immunotherapies.
Overview of Strategies to Improve Therapy against Tumors Using Natural Killer Cell.
Natural Killer Cell-Based Therapies Targeting Cancer: Possible Strategies to Gain and Sustain Anti-Tumor Activity.
Mechanism of tumor cells escaping from immune surveillance of NK cells.
Pharmacological targeting of natural killer cells for cancer immunotherapy.
Natural Killer Cell Immunomodulation: Targeting Activating, Inhibitory, and Co-stimulatory Receptor Signaling for Cancer Immunotherapy.
Natural killer cell-mediated immunosurveillance of human cancer.
Natural Killer Cell-Based Cancer Immunotherapies: From Immune Evasion to Promising Targeted Cellular Therapies.
NK cell self tolerance, responsiveness and missing self recognition.

Plasticity of NK cells in Cancer.
Viral Evasion of Natural Killer Cell Activation.
NK Cells in the Tumor Microenvironment.
Recent Advances to Augment NK Cell Cancer Immunotherapy Using Nanoparticles.
Mechanisms of NK cell dysfunction in the tumor microenvironment and current clinical approaches to harness NK cell potential for immunotherapy.
Triplebody Mediates Increased Anti-Leukemic Reactivity of IL-2 Activated Donor Natural Killer (NK) Cells and Impairs Viability of Their CD33-Expressing NK Subset.
NK cells to cure cancer.
NK cells in the tumor microenvironment: Prognostic and theranostic impact. Recent advances and trends.
Unraveling the role of natural killer cells in leishmaniasis.
Targeting natural killer cells in cancer immunotherapy.
Natural Killer Cell Dysfunction in Hepatocellular Carcinoma: Pathogenesis and Clinical Implications.
Efficient generation of gene-modified human natural killer cells via alpharetroviral vectors.
NK cell-based cancer immunotherapy: from basic biology to clinical development.
Human breast cancer cells enhance self tolerance by promoting evasion from NK cell antitumor immunity.
Escape of tumor cells from the NK cell cytotoxic activity.
Immune evasion of natural killer cells by viruses.
Understanding of molecular mechanisms in natural killer cell therapy.
Natural killer cell immune escape in acute myeloid leukemia.
Regulation of NK cell activation by stimulatory and inhibitory receptors in tumor escape from innate immunity.
PD-1 mediates functional exhaustion of activated NK cells in patients with Kaposi sarcoma.
Improving efficacy of cancer immunotherapy by genetic modification of natural killer cells.
NK Cell Exhaustion.
NK cells and poxvirus infection.
Insufficient natural killer cell responses against retroviruses: how to improve NK cell killing of retrovirus-infected cells.
Mitochondrial fragmentation limits NK cell-based tumor immunosurveillance.
Tumor-Infiltrating Natural Killer Cells.
Imbalance of NKG2D and its inhibitory counterparts: how does tumor escape from innate immunity?
Natural killer cell epigenetic reprogramming in tumors and potential for cancer immunotherapy.
Current progress in cancer immunotherapy based on natural killer cells.
Boosting Natural Killer Cell-Based Immunotherapy with Anticancer Drugs: a Perspective.
Cancer immunotherapy via stem cell-derived NK cells.
Gastric cancer cells inhibit natural killer cell proliferation and induce apoptosis via prostaglandin E2.
NK Cell-Based Immunotherapy and Therapeutic Perspective in Gliomas.
Strategies to Augment Natural Killer (NK) Cell Activity against Solid Tumors.
Abnormal Expression of c-Myc Oncogene in NK Cells in Patients with Cancer.
NK Cell Dysfunction and Checkpoint Immunotherapy.
Immune Circuits to Shape Natural Killer Cells in Cancer.
Unleashing Natural Killer Cells in the Tumor Microenvironment-The Next Generation of Immunotherapy?

Rescue of impaired NK cell activity in hodgkin lymphoma with bispecific antibodies in vitro and in patients.
Mechanisms of resistance to natural killer cell-mediated cytotoxicity in acute lymphoblastic leukemia.
Immune evasion and therapeutic opportunities based on natural killer cells.
Designing Cancer Immunotherapies That Engage T Cells and NK Cells.
Monocyte/macrophage-elicited natural killer cell dysfunction in hepatocellular carcinoma is mediated by CD48/2B4 interactions.
Next Generation Natural Killer Cells for Cancer Immunotherapy.
NK Cell Memory to Cytomegalovirus: Implications for Vaccine Development.
Soluble ligands for NK cell receptors promote evasion of chronic lymphocytic leukemia cells from NK cell anti-tumor activity.
NK Cell Subset Redistribution during the Course of Viral Infections.
Natural Killer Cell Therapy: A New Treatment Paradigm for Solid Tumors.
Natural killer cells unleashed: Checkpoint receptor blockade and BiKE/TriKE utilization in NK-mediated anti-tumor immunotherapy.
Natural killer cells in cancer biology and therapy.
Viral and Nonviral Engineering of Natural Killer Cells as Emerging Adoptive Cancer Immunotherapies.
The role of NK cell as central communicators in cancer immunity.
Improving NK cell function in multiple myeloma with NKTR-255, a novel polymer-conjugated human IL-15.
Reversal of tumor acidosis by systemic buffering reactivates NK cells to express IFN- $\gamma$ and induces NK cell-dependent lymphoma control without other immunotherapies.
The killing effect of Tanshinol on breast cancer cells: insight into the reversion of TGF- $\beta$ 1-mediated suppression of NK cell functions.
High-dimensional mass cytometry analysis of NK cell alterations in AML identifies a subgroup with adverse clinical outcome.
Cancer-Induced Alterations of NK-Mediated Target Recognition: Current and Investigational Pharmacological Strategies Aiming at Restoring NK-Mediated Anti-Tumor Activity.
Targeting immune checkpoints: how to use natural killer cells for fighting against solid tumors.
Melanoma cells inhibit natural killer cell function by modulating the expression of activating receptors and cytolytic activity.
Genetically Engineered Natural Killer Cells as a Means for Adoptive Tumor Immunotherapy.
NK cells and cancer immunosurveillance.
The Interplay of Exosomes and NK Cells in Cancer Biology.
Targeting NK Cell Inhibitory Receptors for Precision Multiple Myeloma Immunotherapy.
Functional and metabolic targeting of natural killer cells to solid tumors.
Herpesvirus Evasion of Natural Killer Cells.
Altered distribution and function of NK-cell subsets lead to impaired tumor surveillance in JAK2V617F myeloproliferative neoplasms.
Natural killer cells: a review of biology, therapeutic potential and challenges in treatment of solid tumors.
Analysis of the Expression of Surface Receptors on NK Cells and NKG2D on Immunocytes in Peripheral Blood of Patients with Nasopharyngeal Carcinoma.
Natural Killer Cells in Liver Disease and Hepatocellular Carcinoma and the NK Cell-Based Immunotherapy.
Therapeutic Potential of Natural Killer Cells in Gastric Cancer.

NK Cell Regulation in Cervical Cancer and Strategies for Immunotherapy.
Underground Adaptation to a Hostile Environment: Acute Myeloid Leukemia vs. Natural Killer Cells.
Clinical development of natural killer cells expressing chimeric antigen receptors.
Irreversible cancer cell-induced functional anergy and apoptosis in resting and activated NK cells.
Human NK cells in acute myeloid leukaemia patients: analysis of NK cell-activating receptors and their ligands.
Trogocytosis-based generation of suppressive NK cells.
Melanoma cells become resistant to NK-cell-mediated killing when exposed to NK-cell numbers compatible with NK-cell infiltration in the tumor.
Tumor-Associated Monocytes/Macrophages Impair NK-Cell Function via TGFβ1 in Human Gastric Cancer.
Sugar Free: Novel Immunotherapeutic Approaches Targeting Siglecs and Sialic Acids to Enhance Natural Killer Cell Cytotoxicity Against Cancer.
Human NK cells: From surface receptors to clinical applications.
The role of NK cell recognition of nectin and nectin-like proteins in tumor immunosurveillance.
Landscape of natural killer cell activity in head and neck squamous cell carcinoma.
Interdependence of sequential cytotoxic T lymphocyte and natural killer cell cytotoxicity against melanoma cells.
Adoptive NK Cell Therapy: A Promising Treatment Prospect for Metastatic Melanoma.
NK cell receptor imbalance and NK cell dysfunction in HBV infection and hepatocellular carcinoma.
Decreased NKG2D expression on NK cells correlates with impaired NK cell function in patients with gastric cancer.
MicroRNA-29b mediates altered innate immune development in acute leukemia.
NK cells are never alone: crosstalk and communication in tumour microenvironments.
Increased TIGIT expressing NK cells with dysfunctional phenotype in AML patients correlated with poor prognosis.
NK Cells in Chronic Lymphocytic Leukemia and Their Therapeutic Implications.
CHMP2A regulates tumor sensitivity to natural killer cell-mediated cytotoxicity.
Exploiting natural killer cells for therapy of melanoma.
Requirements for control of B-cell lymphoma by NK cells.
Upregulation of thioredoxin-1 in activated human NK cells confers increased tolerance to oxidative stress.
Comprehensive characterization of tumor infiltrating natural killer cells and clinical significance in hepatocellular carcinoma based on gene expression profiles.
Natural killer cell-mediated damage of clinical isolates of mucormycetes.
NK Cell Adoptive Immunotherapy of Cancer: Evaluating Recognition Strategies and Overcoming Limitations.
Suppression of a Natural Killer Cell Response by Simian Immunodeficiency Virus Peptides.
Viral inhibitors of NKG2D ligands for tumor surveillance.
Structural insights into activation of antiviral NK cell responses.
Engineering the TGFβ Receptor to Enhance the Therapeutic Potential of Natural Killer Cells as an Immunotherapy for Neuroblastoma.
Harnessing NK cells for cancer immunotherapy: immune checkpoint receptors and chimeric antigen receptors.
Tailoring Natural Killer cell immunotherapy to the tumour microenvironment.

Hypoxia downregulates the expression of activating receptors involved in NK-cell-mediated target cell killing without affecting ADCC.
Down-regulation of PR/SET domain 10 underlies natural killer cell dysfunction in hepatocellular carcinoma.
Tumor growth impedes natural-killer-cell maturation in the bone marrow.
Selection of tumor-resistant variants following sustained natural killer cell-mediated immune stress.
Split anergized Natural Killer cells halt inflammation by inducing stem cell differentiation, resistance to NK cell cytotoxicity and prevention of cytokine and chemokine secretion.
Defective Localization With Impaired Tumor Cytotoxicity Contributes to the Immune Escape of NK Cells in Pancreatic Cancer Patients.
Natural killer cells and acute myeloid leukemia: promises and challenges.
The cancer-natural killer cell immunity cycle.
Hypoxia-inducible factor-1 alpha expression is induced by IL-2 via the PI3K/mTOR pathway in hypoxic NK cells and supports effector functions in NK cells and ex vivo expanded NK cells.
Pseudorabies Virus US3 Protein Kinase Protects Infected Cells from NK Cell-Mediated Lysis via Increased Binding of the Inhibitory NK Cell Receptor CD300a.
Effect of tumor cells and tumor microenvironment on NK-cell function.
Impact of microRNA-29b on natural killer cells in T-cell acute lymphoblastic leukemia.
Natural killer cells and antifungal host response.
IL-15 Overcomes Hepatocellular Carcinoma-Induced NK Cell Dysfunction.
Elevation of MMP-9 and IDO induced by pancreatic cancer cells mediates natural killer cell dysfunction.
A human NK cell activation/inhibition threshold allows small changes in the target cell surface phenotype to dramatically alter susceptibility to NK cells.
EBV-Upregulated B7-H3 Inhibits NK cell-Mediated Antitumor Function and Contributes to Nasopharyngeal Carcinoma Progression.
Mutually assured destruction: the cold war between viruses and natural killer cells.
Hypoxia Impairs NK Cell Cytotoxicity through SHP-1-Mediated Attenuation of STAT3 and ERK Signaling Pathways.
[Effects of recombinant soluble MICA protein on the biologic activities of NK cells].
Combination of Expanded Allogeneic NK Cells and T Cell-Based Immunotherapy Exert Enhanced Antitumor Effects.
Natural Killer Cell Responses in Hepatocellular Carcinoma: Implications for Novel Immunotherapeutic Approaches.
Secreted Ligands of the NK Cell Receptor NKp30: B7-H6 Is in Contrast to BAG6 Only Marginally Released via Extracellular Vesicles.
Natural Killer Cells Recruitment in Oncolytic Virotherapy: A Mathematical Model.
Characterization and Manipulation of the Crosstalk Between Dendritic and Natural Killer Cells Within the Tumor Microenvironment.
Natural Killer Cells: Tumor Surveillance and Signaling.
Tyrosine kinase pathways modulate tumor susceptibility to natural killer cells.
Human NK Cells Develop an Exhaustion Phenotype During Polar Degranulation at the Aspergillus fumigatus Hyphal Synapse.
Cancer-expanded myeloid-derived suppressor cells induce anergy of NK cells through membrane-bound TGF-beta 1.
CD73 immune checkpoint defines regulatory NK cells within the tumor microenvironment.

CD137 ligand mediates opposite effects in human and mouse NK cells and impairs NK-cell reactivity against human acute myeloid leukemia cells.
Tumor-released Galectin-3, a soluble inhibitory ligand of human Nkp30, plays an important role in tumor escape from NK cell attack.
Natural killer cells as effectors of selection and differentiation of stem cells: role in resolution of inflammation.
HIV-1 Nef-mediated downregulation of CD155 results in viral restriction by KIR2DL5+ NK cells.
HIF-1 $\alpha$ induces immune escape of prostate cancer by regulating NCR1/Nkp46 signaling through miR-224.
Murine cytomegalovirus m157 mutation and variation leads to immune evasion of natural killer cells.
Viral escape from NK-cell-mediated immunosurveillance: A lesson for cancer immunotherapy?
Targeting Checkpoint Receptors and Molecules for Therapeutic Modulation of Natural Killer Cells.
KIR-based inhibitory CARs overcome CAR-NK cell trogocytosis-mediated fratricide and tumor escape.
Natural Killer Cells Suppress T Cell-Associated Tumor Immune Evasion.
Functional expression of CD73 on human natural killer cells.
Human AML activates the aryl hydrocarbon receptor pathway to impair NK cell development and function.
An Uncoupling of Canonical Phenotypic Markers and Functional Potency of Ex Vivo-Expanded Natural Killer Cells.
p53 missense mutant G242A subverts natural killer cells in sheltering mouse breast cancer cells against immune rejection.
IL-2 and Anti-TGF- $\beta$ Promote NK Cell Reconstitution and Anti-tumor Effects after Syngeneic Hematopoietic Stem Cell Transplantation.
T cell receptor engineering of primary NK cells to therapeutically target tumors and tumor immune evasion.
Patient's Natural Killer Cells in the Era of Targeted Therapies: Role for Tumor Killers.
Concentration-Dependent Decitabine Effects on Primary NK Cells Viability, Phenotype, and Function in the Absence of Obvious NK Cells Proliferation-Original Article.
CAR-NK Cell: A New Paradigm in Tumor Immunotherapy.
The Function of NK Cells in Tumor Metastasis and NK Cell-Based Immunotherapy.
Identification of a subset of human natural killer cells expressing high levels of programmed death 1: A phenotypic and functional characterization.
The Multifaceted Role of STAT3 in NK-Cell Tumor Surveillance.
Natural killer cells involved in tumour immune escape of hepatocellular carcinomas.
Studying the Anticancer Effects of Thymoquinone on Breast Cancer Cells through Natural Killer Cell Activity.
Cytomegalovirus immunoevasin reveals the physiological role of "missing self" recognition in natural killer cell dependent virus control in vivo.
Increased sMICA and TGF $\beta$ (1) levels in HNSCC patients impair NKG2D-dependent functionality of activated NK cells.
A high concentration of MMP-2/gelatinase A and MMP-9/gelatinase B reduce NK cell-mediated cytotoxicity against an oral squamous cell carcinoma cell line.
Altered Nkp30, Nkp46, NKG2D, and DNAM-1 Expression on Circulating NK Cells Is Associated with Tumor Progression in Human Gastric Cancer.
Role of NK cells in immunotherapy and virotherapy of solid tumors.
Tim-3 Hampers Tumor Surveillance of Liver-Resident and Conventional NK Cells by Disrupting PI3K Signaling.

Mechanisms of tumor and viral immune escape from natural killer cell-mediated surveillance.
Prostaglandin E2 Secreted by Thyroid Cancer Cells Contributes to Immune Escape Through the Suppression of Natural Killer (NK) Cell Cytotoxicity and NK Cell Differentiation.
The combination of type I IFN, TNF- $\alpha$ , and cell surface receptor engagement with dendritic cells enables NK cells to overcome immune evasion by dengue virus.
Different effects of NK cells and NK-derived soluble factors on cell lines derived from primary or metastatic pancreatic cancers.
Zika Virus Escapes NK Cell Detection by Upregulating Major Histocompatibility Complex Class I Molecules.
Investigating the susceptibility of treatment-resistant oesophageal tumours to natural killer cell-mediated responses.
Decreased levels of circulating regulatory NK cells in patients with head and neck cancer throughout all tumor stages.
Heme oxygenase 1 overexpression induces immune evasion of acute myeloid leukemia against natural killer cells by inhibiting CD48.
Cytokine-driven regulation of NK cell functions in tumor immunity: role of the MICA-NKG2D system.
Binding of excreted and/or secreted products of adult hookworms to human NK cells in <i>Necator americanus</i> -infected individuals from Brazil.
Engineered natural killer cells impede the immunometabolic CD73-adenosine axis in solid tumors.
Attenuation of the glucocorticoid response during Ad5IL-12 adenovirus vector treatment enhances natural killer cell-mediated killing of MHC class I-negative LNCaP prostate tumors.
Control of human herpes virus type 8-associated diseases by NK cells.
Circular EZH2-encoded EZH2-92aa mediates immune evasion in glioblastoma via inhibition of surface NKG2D ligands.
Targeting NKG2D and Nkp30 Ligands Shedding to Improve NK Cell-Based Immunotherapy.
Tumors evade immune cytotoxicity by altering the surface topology of NK cells.
The Oncometabolite 5'-Deoxy-5'-Methylthioadenosine Blocks Multiple Signaling Pathways of NK Cell Activation.
Renal carcinoma cell lines inhibit natural killer activity via the CD94 receptor molecule.
NK cells in pancreatic cancer demonstrate impaired cytotoxicity and a regulatory IL-10 phenotype.
CMV and natural killer cells: shaping the response to vaccination.
NK Cell-Based Immune Checkpoint Inhibition.
Strategies to endow cytotoxic T lymphocytes or natural killer cells with antibody activity against carcinoembryonic antigen.
Interleukin-10 reduces natural killer sensitivity and downregulates MHC class I expression on H-ras-transformed cells.
CD8 T cell help for innate antitumor immunity.
Pleckstrin-2 promotes tumour immune escape from NK cells by activating the MT1-MMP-MICA signalling axis in gastric cancer.
Enhanced interaction between natural killer cells and lung cancer cells: involvement in gefitinib-mediated immunoregulation.
IL-21-mediated reversal of NK cell exhaustion facilitates anti-tumour immunity in MHC class I-deficient tumours.
Adoptive NK Cell Transfer as a Treatment in Colorectal Cancer Patients: Analyses of Tumour Cell Determinants Correlating With Efficacy In Vitro and In Vivo.
Naturally Killing the Silent Killer: NK Cell-Based Immunotherapy for Ovarian Cancer.
PD-L1 Mediates Dysfunction in Activated PD-1(+) NK Cells in Head and Neck Cancer Patients.

Hematological malignancies escape from NK cell innate immune surveillance: mechanisms and therapeutic implications.
NK/DC crosstalk in anti-viral response.
Human breast tumor cells induce self-tolerance mechanisms to avoid NKG2D-mediated and DNAM-mediated NK cell recognition.
Hypoxia Modifies the Transcriptome of Human NK Cells, Modulates Their Immunoregulatory Profile, and Influences NK Cell Subset Migration.
Sensitizing primary acute lymphoblastic leukemia to natural killer cell recognition by induction of NKG2D ligands.
Tanshinone IIA enhances susceptibility of non-small cell lung cancer cells to NK cell-mediated lysis by up-regulating ULBP1 and DR5.
Mathematical model of tumor-immune surveillance.
Effect of vasoactive intestinal peptide (VIP) on NKG2D signal pathway and its contribution to immune escape of MKN45 cells.
Intercellular protein transfer at the NK cell immune synapse: mechanisms and physiological significance.
Monalizumab efficacy correlates with HLA-E surface expression and NK cell activity in head and neck squamous carcinoma cell lines.
Remodeled tumor immune microenvironment (TIME) parade via natural killer cells reprogramming in breast cancer.
Potential of NK cells in multiple Myeloma therapy.
B7-H6-mediated downregulation of Nkp30 in NK cells contributes to ovarian carcinoma immune escape.
HCMV vCXCL1 Binds Several Chemokine Receptors and Preferentially Attracts Neutrophils over NK Cells by Interacting with CXCR2.
Human tumour immune evasion via TGF- $\beta$ blocks NK cell activation but not survival allowing therapeutic restoration of anti-tumour activity.
Head and neck cancer triggers the internalization of TLR3 in natural killer cells.
Defective NK Cells in Acute Myeloid Leukemia Patients at Diagnosis Are Associated with Blast Transcriptional Signatures of Immune Evasion.
Cetuximab Reconstitutes Pro-Inflammatory Cytokine Secretions and Tumor-Infiltrating Capabilities of sMICA-Inhibited NK Cells in HNSCC Tumor Spheroids.
The crosstalk between endometrial stromal cells and macrophages impairs cytotoxicity of NK cells in endometriosis by secreting IL-10 and TGF- $\beta$ .
The BET-bromodomain inhibitor JQ1 renders neuroblastoma cells more resistant to NK cell-mediated recognition and killing by downregulating ligands for NKG2D and DNAM-1 receptors.
Melanoma-associated fibroblasts modulate NK cell phenotype and antitumor cytotoxicity.
Acquired Natural Killer Cell Dysfunction in the Tumor Microenvironment of Classic Hodgkin Lymphoma.
Intrinsic Resistance of Chronic Lymphocytic Leukemia Cells to NK Cell-Mediated Lysis Can Be Overcome In Vitro by Pharmacological Inhibition of Cdc42-Induced Actin Cytoskeleton Remodeling.
Recognition of a virus-encoded ligand by a natural killer cell activation receptor.
Mediastinal lymph node removal modulates natural killer cell exhaustion in patients with non-small cell lung cancer.
Dynamic change in natural killer cell type in the human ocular mucosa in situ as means of immune evasion by adenovirus infection.
Hepatocellular carcinoma-associated fibroblasts trigger NK cell dysfunction via PGE2 and IDO.



NK cell-based therapeutics for lung cancer.
IL-6-induced CD39 expression on tumor-infiltrating NK cells predicts poor prognosis in esophageal squamous cell carcinoma.
Exosomal miRNA Cargo as Mediator of Immune Escape Mechanisms in Neuroblastoma.
Protein kinase C-theta is required for NK cell activation and in vivo control of tumor progression.
Natural killer cells modulation in hematological malignancies.
Influenza Virus Targets Class I MHC-Educated NK Cells for Immunoavoidance.
Increased apoptosis and elevated Fas expression in circulating natural killer cells in gastric cancer patients.
[K562 cells induces apoptosis of activated NK cells in vitro].
Transplantation-induced cancers: Emerging evidence that clonal CMV-specific NK cells are causal immunogenic factors.
Natural Killer Immunotherapy for Minimal Residual Disease Eradication Following Allogeneic Hematopoietic Stem Cell Transplantation in Acute Myeloid Leukemia.
Cancer extracellular vesicles as novel regulators of NK cell response.
MicroRNA analysis of Natural Killer cell-derived exosomes: the microRNA let-7b-5p is enriched in exosomes and participates in their anti-tumor effects against pancreatic cancer cells.
Reduced CD160 Expression Contributes to Impaired NK-cell Function and Poor Clinical Outcomes in Patients with HCC.
T and NK cells: two sides of tumor immunoavoidance.
Inhibiting exosomal MIC-A and MIC-B shedding of cancer cells to overcome immune escape: new insight of approved drugs.
NK cells infiltrating a MHC class I-deficient lung adenocarcinoma display impaired cytotoxic activity toward autologous tumor cells associated with altered NK cell-triggering receptors.
Natural Killer Cell-Derived Vesicular miRNAs: A New Anticancer Approach?
Natural Killer Cells: the Missing Link in Effective Treatment for High-Grade Serous Ovarian Carcinoma.
Compromised activity of natural killer cells in diffuse large b-cell lymphoma is related to lymphoma-induced modification of their surface receptor expression.
Tumor escape mechanisms: potential role of soluble HLA antigens and NK cells activating ligands.
Apigenin Increases Natural Killer Cytotoxicity to Human Hepatocellular Carcinoma Expressing HIF-1 $\alpha$ through High Interaction of CD95/CD95L.
HER-2/neu peptide specificity in the recognition of HLA-A2 by natural killer cells.
Re-targeting of cytotoxic T lymphocytes and/or natural killer cells to CEA-expressing tumor cells with anti-CEA antibody activity.
Targeting the $\alpha$ v integrin/TGF- $\beta$ axis improves natural killer cell function against glioblastoma stem cells.
Peritoneal natural killer cells from epithelial ovarian cancer patients show an altered phenotype and bind to the tumour marker MUC16 (CA125).
Structure-based rational design of a novel chimeric PD1-NKG2D receptor for natural killer cells.
Metabolic Reprogramming of NK Cells by Black Phosphorus Quantum Dots Potentiates Cancer Immunotherapy.
Natural killer cells and malignant haemopathies: a model for the interaction of cancer with innate immunity.
Formyl peptide receptor suppresses melanoma development and promotes NK cell migration.
Overexpression of LLT1 (OCIL, CLEC2D) on prostate cancer cells inhibits NK cell-mediated killing through LLT1-NKRP1A (CD161) interaction.
Varicella zoster virus productively infects human natural killer cells and manipulates phenotype.

Hypoxia-inducible gene 2 promotes the immune escape of hepatocellular carcinoma from nature killer cells through the interleukin-10-STAT3 signaling pathway.
Natural killer cell therapy targeting cancer stem cells: Old wine in a new bottle.
HCMV infection of PDCs deviates the NK cell response into cytokine-producing cells unable to perform cytotoxicity.
Live-cell imaging for analysis of the NK cell immunological synapse.
Genome-Wide CRISPR Screen Reveals Cancer Cell Resistance to NK Cells Induced by NK-Derived IFN- $\gamma$ .
Tumor-derived lactate modifies antitumor immune response: effect on myeloid-derived suppressor cells and NK cells.
Radiation alters PD-L1/NKG2D ligand levels in lung cancer cells and leads to immune escape from NK cell cytotoxicity via IL-6-MEK/Erk signaling pathway.
Altered expression of CD226 and CD96 on natural killer cells in patients with pancreatic cancer.
Involvement of TIGIT in Natural Killer Cell Exhaustion and Immune Escape in Patients and Mouse Model With Liver <i>Echinococcus multilocularis</i> Infection.
MICA-G129R: A bifunctional fusion protein increases PRLR-positive breast cancer cell death in co-culture with natural killer cells.
TM4SF5-mediated liver malignancy involves NK cell exhaustion-like phenotypes.
Metalloprotease-mediated tumor cell shedding of B7-H6, the ligand of the natural killer cell-activating receptor NKp30.
Immune evasion via PD-1/PD-L1 on NK cells and monocyte/macrophages is more prominent in Hodgkin lymphoma than DLBCL.
Myeloid derived suppressor cells inhibit natural killer cells in patients with hepatocellular carcinoma via the NKp30 receptor.
Therapeutic Approaches Targeting the Natural Killer-Myeloid Cell Axis in the Tumor Microenvironment.
Decreased expression of DNAM-1 on NK cells from acute myeloid leukemia patients.
Ganglioside GD3 expression on target cells can modulate NK cell cytotoxicity via siglec-7-dependent and -independent mechanisms.
Defective expression and function of natural killer cell-triggering receptors in patients with acute myeloid leukemia.
Metastatic consequences of immune escape from NK cell cytotoxicity by human breast cancer stem cells.
Natural killer cells, dendritic cells, and the alarmin high-mobility group box 1 protein: a dangerous trio in HIV-1 infection?
Expression of NK cell receptor ligands in primary colorectal cancer tissue in relation to the phenotype of circulating NK- and NKT cells, and clinical outcome.
Critical role of the NKG2D receptor for NK cell-mediated control and immune escape of B-cell lymphoma.
microRNA-889 is downregulated by histone deacetylase inhibitors and confers resistance to natural killer cytotoxicity in hepatocellular carcinoma cells.
Inhibitory Receptors and Checkpoints in Human NK Cells, Implications for the Immunotherapy of Cancer.
Analysis of the mechanisms regulating soluble PD-1 production and function in human NK cells.
Decitabine has a biphasic effect on natural killer cell viability, phenotype, and function under proliferative conditions.
Association of CD47 with natural killer cell-mediated cytotoxicity of head-and-neck squamous cell carcinoma lines.

The TGF- $\beta$ /SMAD pathway is an important mechanism for NK cell immune evasion in childhood B-acute lymphoblastic leukemia.
DNA damage response and evasion from immunosurveillance in CLL: new options for NK cell-based immunotherapies.
UL40-mediated NK evasion during productive infection with human cytomegalovirus.
ELFN1-AS1 promotes GDF15-mediated immune escape of colorectal cancer from NK cells by facilitating GCN5 and SND1 association.
Arsenite suppresses IL-2-dependent tumoricidal activities of natural killer cells.
Glycocalyx engineering reveals a Siglec-based mechanism for NK cell immunoevasion.
Inhibition of natural killer cell-mediated cytotoxicity by Kaposi's sarcoma-associated herpesvirus K5 protein.
Altered-Self MHC Class I Sensing via Functionally Disparate Paired NK Cell Receptors Counters Murine Cytomegalovirus gp34-Mediated Immune Evasion.
CD56(bright)perforin(low) noncytotoxic human NK cells are abundant in both healthy and neoplastic solid tissues and recirculate to secondary lymphoid organs via afferent lymph.
Suppression of Natural Killer cell NKG2D and CD226 anti-tumour cascades by platelet cloaked cancer cells: Implications for the metastatic cascade.
Cell-based immunotherapy in stage IIIA inflammatory breast cancer with declining innate immunity following successive chemotherapies: A case report.
PVR and ICAM-1 on Blast Crisis CML Stem and Progenitor Cells with TKI Resistance Confer Susceptibility to NK Cells.
Critical roles of co-activation receptor DNAX accessory molecule-1 in natural killer cell immunity.
Tumor-Derived PGE2 Gives NK Cells a Headache.
Deficiency of IL-18 Aggravates Esophageal Carcinoma Through Inhibiting IFN- $\gamma$ Production by CD8(+)T Cells and NK Cells.
Iron-induced parafibrin formation in tumors fosters immune evasion.
Taming Tumor Glycolysis and Potential Implications for Immunotherapy.
Hsa_circ_0007456 regulates the natural killer cell-mediated cytotoxicity toward hepatocellular carcinoma via the miR-6852-3p/ICAM-1 axis.
Immune escape mechanisms in Hodgkin's disease.
Advancements in CAR-NK therapy: lessons to be learned from CAR-T therapy.
Immunotherapeutic approaches for hematologic malignancies.
Ligand-Receptor Interactions of Galectin-9 and VISTA Suppress Human T Lymphocyte Cytotoxic Activity.
Malignant glioma cells counteract antitumor immune responses through expression of lectin-like transcript-1.
Increased levels of soluble CD226 in sera accompanied by decreased membrane CD226 expression on peripheral blood mononuclear cells from cancer patients.
Potential contribution of naïve immune effectors to oral tumor resistance: role in synergistic induction of VEGF, IL-6, and IL-8 secretion.
Design, Synthesis, and Biological Evaluation of Small, High-Affinity Siglec-7 Ligands: Toward Novel Inhibitors of Cancer Immune Evasion.
CD73-deficient mice are resistant to carcinogenesis.
Association of killer cell immunoglobulin-like receptors with endemic Burkitt lymphoma in Kenyan children.
Homeobox containing 1 inhibits liver cancer progression by promoting autophagy as well as inhibiting stemness and immune escape.

Opposing consequences of signaling through EGF family members: Escape from CTLs could be a bait for NK cells.
Human melanoma cell secreting human leukocyte antigen-G5 inhibit natural killer cell cytotoxicity by impairing lytic granules polarization toward target cell.
Targeting EpCAM by a Bispecific Trifunctional Antibody Exerts Profound Cytotoxic Efficacy in Germ Cell Tumor Cell Lines.
Molecular and structural basis of TIGIT: Nectin-4 interaction, a recently discovered pathway crucial for cancer immunotherapy.
RNF31 inhibition sensitizes tumors to bystander killing by innate and adaptive immune cells.
Precision glycoalkyl editing as a strategy for cancer immunotherapy.
The Inhibitory NKR-P1B:Clr-b Recognition Axis Facilitates Detection of Oncogenic Transformation and Cancer Immunosurveillance.
High co-expression of immune checkpoint receptors PD-1, CTLA-4, LAG-3, TIM-3, and TIGIT on tumor-infiltrating lymphocytes in early-stage breast cancer.
The chemoattractant chemerin suppresses melanoma by recruiting natural killer cell antitumor defenses.
Metastatic Latency and Immune Evasion through Autocrine Inhibition of WNT.
Adenosinergic signaling as a target for natural killer cell immunotherapy.
Natural Killer Cells from the Subcutaneous Adipose Tissue Underexpress the NKp30 and NKp44 in Obese Persons and Are Less Active against Major Histocompatibility Complex Class I Non-Expressing Neoplastic Cells.
AML1-ETO inhibits acute myeloid leukemia immune escape by CD48.
Expression and purification of soluble and stable ectodomain of natural killer cell receptor LLT1 through high-density transfection of suspension adapted HEK293S GnTI(-) cells.
Immune reaction to breast cancer: for better or for worse?
The presence of IGHG1 in human pancreatic carcinomas is associated with immune evasion mechanisms.
Alterations in target cell membrane phospholipids alter T cell but not NK cell killing.
Differing phenotypes between intraepithelial and stromal lymphocytes in early-stage tongue cancer.
A gastric cancer cell derived extracellular compounds suppresses CD161(+)CD3(-) lymphocytes and aggravates tumor formation in a syngeneic mouse model.
Immune protection function of multipotent mesenchymal stromal cells: role of transforming growth factor- $\beta$ 1.
Aberrant positioning of trophoblast and lymphocytes in the feto-maternal interface with pre-eclampsia.
CD155: A Multi-Functional Molecule in Tumor Progression.
Induced Pluripotent Stem Cell-conditioned Medium Suppressed Melanoma Tumorigenicity Through the Enhancement of Natural-Killer Cellular Immunity.
Adaptive Immunity and the Tumor Microenvironment.
Emerging Canonical and Non-Canonical Roles of Granzyme B in Health and Disease.
Matrix metalloproteinase 9 (MMP-9/gelatinase B) proteolytically cleaves ICAM-1 and participates in tumor cell resistance to natural killer cell-mediated cytotoxicity.
Reduced Expression of Galectin-9 Contributes to a Poor Outcome in Colon Cancer by Inhibiting NK Cell Chemotaxis Partially through the Rho/ROCK1 Signaling Pathway.
miR-4299 inhibits tumor progression in pancreatic cancer through targeting ADAM17.
Reovirus-induced cell-mediated immunity for the treatment of multiple myeloma within the resistant bone marrow niche.

The Tim-3-galectin-9 Secretory Pathway is Involved in the Immune Escape of Human Acute Myeloid Leukemia Cells.
The cooperating mutation or "second hit" determines the immunologic visibility toward MYC-induced murine lymphomas.
Prognostic implication of CD57, CD16, and TGF- $\beta$ expression in oral squamous cell carcinoma.
Evasion of the immune system by adenoviruses.
Immune Landscape of CMV Infection in Cancer Patients: From "Canonical" Diseases Toward Virus-Elicited Oncomodulation.
Alterations in the Transcriptional Programs of Myeloma Cells and the Microenvironment during Extramedullary Progression Affect Proliferation and Immune Evasion.
Mouse Models to Study Natural Killer Cell-Mediated Immunosurveillance and Metastatic Latency.
Dickkopf-1 Can Lead to Immune Evasion in Metastatic Castration-Resistant Prostate Cancer.
Breast cancer stem-like cells can promote metastasis by activating platelets and down-regulating antitumor activity of natural killer cells.
The Role of Natural Killer T Cells in Cancer-A Phenotypical and Functional Approach.
Gene inactivation promotes immune escape in DLBCL.
Plasmonic Nanoparticle-Based Digital Cytometry to Quantify MUC16 Binding on the Surface of Leukocytes in Ovarian Cancer.
The B7 Family Member B7-H6: a New Bane of Tumor.
Good news, bad news for missing-self recognition by NK cells: autoimmune control but viral evasion.
Single-cell analysis reveals the multiple patterns of immune escape in the nasopharyngeal carcinoma microenvironment.
The up side of decidual natural killer cells: new developments in immunology of pregnancy.
Role of innate immunity against human papillomavirus (HPV) infections and effect of adjuvants in promoting specific immune response.
Boronic acid-based arginase inhibitors in cancer immunotherapy.
CD317 mediates immunocytolysis resistance by RICH2/cytoskeleton-dependent membrane protection.
Organoid Co-culture Methods to Capture Cancer Cell-Natural Killer Cell Interactions.
Anti-CD39 and anti-CD73 antibodies A1 and 7G2 improve targeted therapy in ovarian cancer by blocking adenosine-dependent immune evasion.
Lectin-Like Transcript 1 (LLT1) Checkpoint: A Novel Independent Prognostic Factor in HPV-Negative Oropharyngeal Squamous Cell Carcinoma.
AKT inhibition interferes with the expression of immune checkpoint proteins and increases NK-induced killing of HL60-AML cells.
Engineering Human Pluripotent Stem Cell Lines to Evade Xenogeneic Transplantation Barriers.
Role of the ERK pathway for oxidant-induced parthanatos in human lymphocytes.
[Nectins and nectin-like receptors DNAM-1 and CRTAM: new ways for tumor escape].
Inhibition of cell growth and induction of apoptotic cell death by the human tumor-associated antigen RCAS1.
Preclinical Evaluation of CRISPR-Edited CAR-NK-92 Cells for Off-the-Shelf Treatment of AML and B-ALL.
The effect of H-ras expression on tumorigenicity and immunogenicity of Balb/c 3T3 fibroblasts.
Ex vivo detection of primary leukemia cells resistant to granule cytotoxin-induced cell death: a rapid isolation method to study granzyme-B-mediated cell death.
Low concentrations of the soy phytoestrogen genistein induce proteinase inhibitor 9 and block killing of breast cancer cells by immune cells.

Transcriptional heterogeneity of clonal plasma cells and immune evasion in immunoglobulin light chain amyloidosis.
Tumor immunoediting by NKp46.
Lectin-like transcript 1 is a marker of germinal center-derived B-cell non-Hodgkin's lymphomas dampening natural killer cell functions.
Macrophage migration inhibitory factor (MIF) expression in human malignant gliomas contributes to immune escape and tumour progression.
Autocrine WNT Inhibition Drives Immune Evasion and Latency of Metastases.
KIR3DL3 Is an Inhibitory Receptor for HHLA2 that Mediates an Alternative Immunoinhibitory Pathway to PD1.
Scalable Enrichment of Immunomodulatory Human Acute Myeloid Leukemia Cell Line-Derived Extracellular Vesicles.
Deacetylated sialic acids modulates immune mediated cytotoxicity via the sialic acid-Siglec pathway.
Strategies to enhance the graft versus tumour effect after allogeneic haematopoietic stem cell transplantation.
Gnawing Between Cells and Cells in the Immune System: Friend or Foe? A Review of Trogocytosis.
Targeting and amplification of immune killing of tumor cells by pro-Smac.
Suppression of granzyme B activity and caspase-3 activation in leukaemia cells constitutively expressing the protease inhibitor 9.
Invariant Natural Killer T Cells in Immune Regulation of Blood Cancers: Harnessing Their Potential in Immunotherapies.
Donor lymphocyte infusions for multiple myeloma: clinical results and novel perspectives.
To PARP or not to PARP?-Toward sensitizing acute myeloid leukemia stem cells to immunotherapy.
Decreased B7-H3 promotes unexplained recurrent miscarriage via RhoA/ROCK2 signaling pathway and regulates the secretion of decidual NK cells†.
The tumour suppressor gene CEACAM1 is completely but reversibly downregulated in renal cell carcinoma.
Cholangiocarcinoma-derived exosomes inhibit the antitumor activity of cytokine-induced killer cells by down-regulating the secretion of tumor necrosis factor- $\alpha$ and perforin.
Genistein Sensitizes Human Cholangiocarcinoma Cell Lines to Be Susceptible to Natural Killer Cells.
Defective immune function of primary effusion lymphoma cells is associated with distinct KSHV gene expression profiles.
Immunological mechanisms elicited at the tumour site by lymphocyte activation gene-3 (LAG-3) versus IL-12: sharing a common Th1 anti-tumour immune pathway.
Both E6 and E7 oncoproteins of human papillomavirus 16 inhibit IL-18-induced IFN-gamma production in human peripheral blood mononuclear and NK cells.
Intracellular Delivery of Synthetic dsRNA to Leukemic Cells Induces Apoptotic and Necrotic Cell Death.
AML1-ETO targets and suppresses cathepsin G, a serine protease, which is able to degrade AML1-ETO in t(8;21) acute myeloid leukemia.

Table A2-21, Cluster 20

Cluster 20 focuses on herpes simplex virus infections (especially hsv-1), emphasizing immune system evasion by hsv-1, and therapies to reduce this immune evasion (198)
Immunity in latent Herpes simplex virus infection.
Pathogenesis and virulence of herpes simplex virus.

[Immune evasion by herpes simplex viruses].
Autophagy interaction with herpes simplex virus type-1 infection.
Medical application of herpes simplex virus.
Improving immunogenicity and efficacy of vaccines for genital herpes containing herpes simplex virus glycoprotein D.
[Herpes simplex virus vaccines: perspectives].
Activation and evasion of innate antiviral immunity by herpes simplex virus.
Herpes Simplex Vaccines: Prospects of Live-attenuated HSV Vaccines to Combat Genital and Ocular infections.
Herpes Simplex Virus Type 1 and Host Antiviral Immune Responses: An Update.
Dancing with the enemy: the interplay of herpes simplex virus with dendritic cells.
The immune response to ocular herpes simplex virus type 1 infection.
Innate Immune Mechanisms and Herpes Simplex Virus Infection and Disease.
Herpes simplex: insights on pathogenesis and possible vaccines.
The Proteome and Secretome of Cortical Brain Cells Infected With Herpes Simplex Virus.
Herpes simplex virus 1 UL41 protein abrogates the antiviral activity of hZAP by degrading its mRNA.
Induction of apoptosis by herpes simplex virus in Jurkat cells is partly through caspase-3, -8 and -9 activation.
The Us3 Protein of Herpes Simplex Virus 1 Inhibits T Cell Signaling by Confining Linker for Activation of T Cells (LAT) Activation via TRAF6 Protein.
Regulated expression of galectin-1 after in vitro productive infection with herpes simplex virus type 1: implications for T cell apoptosis.
Herpes Simplex Virus: A Versatile Tool for Insights Into Evolution, Gene Delivery, and Tumor Immunotherapy.
Herpes Simplex Virus Evasion of Early Host Antiviral Responses.
Herpes simplex virus $\gamma$ 34.5 interferes with autophagosome maturation and antigen presentation in dendritic cells.
Herpes Simplex Virus 1 $\gamma$ (1)34.5 Protein Inhibits STING Activation That Restricts Viral Replication.
Role of Caspases and Gasdermin A during HSV-1 Infection in Mice.
Herpes Simplex Virus Type 1 Interactions with the Interferon System.
Characteristics of herpes simplex virus infection and pathogenesis suggest a strategy for vaccine development.
Advancing Our Understanding of Corneal Herpes Simplex Virus-1 Immune Evasion Mechanisms and Future Therapeutics.
Investigation of the Anti-HSV Activity of Candidate Antiviral Agents.
Apoptosis and antigen receptor function in T and B cells following exposure to herpes simplex virus.
Molecular characterization of clinical isolates of herpes simplex virus type 1 collected in a tertiary-care hospital in Dublin, Ireland.
The interaction between dendritic cells and herpes simplex virus-1.
The Race between Host Antiviral Innate Immunity and the Immune Evasion Strategies of Herpes Simplex Virus 1.
Role of Microvesicles in the Spread of Herpes Simplex Virus 1 in Oligodendrocytic Cells.
Glycoprotein C of Herpes Simplex Virus 1 Shields Glycoprotein B from Antibody Neutralization.
Herpes simplex virus type 1 renders infected cells resistant to cytotoxic T-lymphocyte-induced apoptosis.
Recent advances in vaccine development for herpes simplex virus types I and II.

Blocking immune evasion as a novel approach for prevention and treatment of herpes simplex virus infection.
Inhibition of dendritic cell maturation by herpes simplex virus.
Infection of mature dendritic cells with herpes simplex virus type 1 dramatically reduces lymphoid chemokine-mediated migration.
Reducing Viral Inhibition of Host Cellular Apoptosis Strengthens the Immunogenicity and Protective Efficacy of an Attenuated HSV-1 Strain.
Novel transcription regulatory sequences and factors of the immune evasion protein ICP47 (US12) of herpes simplex viruses.
Herpes Simplex Virus 1 Specifically Targets Human CD1d Antigen Presentation To Enhance Its Pathogenicity.
Study of interferon- $\beta$ antiviral activity against Herpes simplex virus type 1 in neuron-enriched trigeminal ganglia cultures.
Herpes simplex virus interference with immunity: Focus on dendritic cells.
Suppression of proinflammatory cytokine expression by herpes simplex virus type 1.
Dual impacts of a glycan shield on the envelope glycoprotein B of HSV-1: evasion from human antibodies in vivo and neurovirulence.
Deletion of the virion host shutoff protein (vhs) from herpes simplex virus (HSV) relieves the viral block to dendritic cell activation: potential of vhs- HSV vectors for dendritic cell-mediated immunotherapy.
CD80 Plays a Critical Role in Increased Inflammatory Responses in Herpes Simplex Virus 1-Infected Mouse Corneas.
Herpes Simplex Virus Type 1-Encoded miR-H2-3p Manipulates Cytosolic DNA-Stimulated Antiviral Innate Immune Response by Targeting DDX41.
Using CSF Proteomics to Investigate Herpesvirus Infections of the Central Nervous System.
Evasion of Cytosolic DNA-Stimulated Innate Immune Responses by Herpes Simplex Virus 1.
Non-cytopathic herpes simplex virus type-1 isolated from acyclovir-treated patients with recurrent infections.
Temporal Proteomic Analysis of Herpes Simplex Virus 1 Infection Reveals Cell-Surface Remodeling via pUL56-Mediated GOPC Degradation.
Herpes simplex virus type 2 trivalent protein vaccine containing glycoproteins C, D and E protects guinea pigs against HSV-1 genital infection.
Protection from lethal herpes simplex virus type 1 infection by vaccination with a UL41-deficient recombinant strain.
Decreased T cell stimulatory capacity of monocyte-derived human macrophages following herpes simplex virus type 1 infection.
Vaccines to prevent genital herpes.
Decreased reactivation of a herpes simplex virus type 1 (HSV-1) latency-associated transcript (LAT) mutant using the in vivo mouse UV-B model of induced reactivation.
Implications for herpes simplex virus vaccine strategies based on antibodies produced to herpes simplex virus type 1 glycoprotein gC immune evasion domains.
Herpes Simplex Virus 1 ICP22 Suppresses CD80 Expression by Murine Dendritic Cells.
MicroRNA-649 promotes HSV-1 replication by directly targeting MALT1.
The herpes simplex virus 1 latency-associated transcript promotes functional exhaustion of virus-specific CD8+ T cells in latently infected trigeminal ganglia: a novel immune evasion mechanism.
Herpes simplex virus and varicella-zoster virus: why do these human alphaherpesviruses behave so differently from one another?



Herpes Simplex Virus 1 US3 Phosphorylates Cellular KIF3A To Downregulate CD1d Expression.
Herpes simplex virus protein UL56 inhibits cGAS-Mediated DNA sensing to evade antiviral immunity.
Herpes simplex virus type 1 evades the effects of antibody and complement in vivo.
Altered expression of cytokines in mice infected intranasally with two syncytial variants of Herpes simplex virus type 1.
Quantification of the host response proteome after herpes simplex virus type 1 infection.
Involvement of herpes simplex virus type 1 UL13 protein kinase in induction of SOCS genes, the negative regulators of cytokine signaling.
Short interfering RNA-mediated inhibition of herpes simplex virus type 1 gene expression and function during infection of human keratinocytes.
Vaccines against genital herpes: progress and limitations.
Dynamics of IL-15/IL-15R- $\alpha$ expression in response to HSV-1 infection reveal a novel mode of viral immune evasion counteracted by iNKT cells.
Induction of signaling pathways by herpes simplex virus type 1 through glycoprotein H peptides.
Induction of transforming growth factor-beta 1 production in human cells by herpes simplex virus.
Immune mechanisms induced by an HSV-1 mutant strain: Discrepancy analysis of the immune system gene profile in comparison with a wild-type strain.
Infection of dendritic cells with herpes simplex virus type 1 induces rapid degradation of CYTIP, thereby modulating adhesion and migration.
Novel deletion in glycoprotein G forms a cluster and causes epidemiologic spread of herpes simplex virus type 2 infection.
Blocking herpes simplex virus 2 glycoprotein E immune evasion as an approach to enhance efficacy of a trivalent subunit antigen vaccine for genital herpes.
Species-Specific Deamidation of cGAS by Herpes Simplex Virus UL37 Protein Facilitates Viral Replication.
Herpes simplex virus type 2 recurrent meningitis (Mollaret's meningitis): a consideration for the recurrent pathogenesis.
Direct contact with herpes simplex virus-infected cells results in inhibition of lymphokine-activated killer cells because of cell-to-cell spread of virus.
Innate and adaptive immunity against herpes simplex virus type 2 in the genital mucosa.
In vivo role of complement-interacting domains of herpes simplex virus type 1 glycoprotein gC.
Herpes simplex virus 1 infection induces activation and subsequent inhibition of the IFI16 and NLRP3 inflammasomes.
Immune evasion by herpes simplex virus type 1, strategies for virus survival.
Properties of a herpes simplex virus multiple immediate-early gene-deleted recombinant as a vaccine vector.
Comparison of neuroinvasive abilities of HSV-1 and HSV-2 in mice.
Herpes Simplex Virus 1 Infection of Neuronal and Non-Neuronal Cells Elicits Specific Innate Immune Responses and Immune Evasion Mechanisms.
Herpes simplex virus-1 evasion of CD8+ T cell accumulation contributes to viral encephalitis.
Extracellular Vesicles in Herpes Viral Spread and Immune Evasion.
In vivo immune evasion mediated by the herpes simplex virus type 1 immunoglobulin G Fc receptor.
Herpes simplex virus type 1 induces CD83 degradation in mature dendritic cells with immediate-early kinetics via the cellular proteasome.
Herpes simplex virus 2 (HSV-2) prevents dendritic cell maturation, induces apoptosis, and triggers release of proinflammatory cytokines: potential links to HSV-HIV synergy.

The interplay between human herpes simplex virus infection and the apoptosis and necroptosis cell death pathways.
Are miRNAs critical determinants in herpes simplex virus pathogenesis?
Herpes simplex virus 1 counteracts tetherin restriction via its virion host shutoff activity.
The herpes simplex virus 1 IgG fc receptor blocks antibody-mediated complement activation and antibody-dependent cellular cytotoxicity in vivo.
Herpes simplex virus suppresses necroptosis in human cells.
Infection of murine keratinocytes with herpes simplex virus type 1 induces the expression of interleukin-10, but not interleukin-1 alpha or tumour necrosis factor-alpha.
Herpes simplex virus type 1 glycoprotein D inhibits T-cell proliferation.
Herpes simplex virus: receptors and ligands for cell entry.
A LAT-associated function reduces productive-cycle gene expression during acute infection of murine sensory neurons with herpes simplex virus type 1.
Herpes simplex virus interferes with monocyte accessory cell function.
Toll-like receptors as novel therapeutic targets for herpes simplex virus infection.
The Herpes Simplex Virus Latency-Associated Transcript Gene Is Associated with a Broader Repertoire of Virus-Specific Exhausted CD8+ T Cells Retained within the Trigeminal Ganglia of Latently Infected HLA Transgenic Rabbits.
Herpes simplex virus evades natural killer T cell recognition by suppressing CD1d recycling.
What Goes Around, Comes Around - HSV-1 Replication in Monocyte-Derived Dendritic Cells.
Infection of human dendritic cells with herpes simplex virus type 1 dramatically diminishes the mRNA levels of the prostaglandin E(2) receptors EP2 and EP4.
The herpes simplex virus type 1 latency-associated transcript inhibits phenotypic and functional maturation of dendritic cells.
Kinetic analysis of glycoprotein C of herpes simplex virus types 1 and 2 binding to heparin, heparan sulfate, and complement component C3b.
Optineurin in ocular herpes infection.
Interleukin-1 $\alpha$ released from HSV-1-infected keratinocytes acts as a functional alarmin in the skin.
Herpes simplex virus type 1-induced FasL expression in human monocytic cells and its implications for cell death, viral replication, and immune evasion.
Oncolytic herpes simplex virus and immunotherapy.
HSV-2: in pursuit of a vaccine.
Evolution of rational vaccine designs for genital herpes immunotherapy.
Immune evasion properties of herpes simplex virus type 1 glycoprotein gC.
DNA viruses and genetic modification of dendritic cells.
Inhibition of TCR signaling by herpes simplex virus.
HSV-1 selectively packs the transcription factor Oct-1 into EVs to facilitate its infection.
A Tribute to Professor Steven L. Wechsler (1948-2016): The Man and the Scientist.
Herpes simplex virus US3 tegument protein inhibits Toll-like receptor 2 signaling at or before TRAF6 ubiquitination.
HSV-1-Specific IgG Subclasses Distribution and Serum Neutralizing Activity in Alzheimer's Disease and in Mild Cognitive Impairment.
Double stranded RNA activated EIF2 alpha kinase (EIF2AK2; PKR) is associated with Alzheimer's disease.
VHS, US3 and UL13 viral tegument proteins are required for Herpes Simplex Virus-Induced modification of protein kinase R.

Herpes simplex virus genes Us3, Us5, and Us12 differentially regulate cytotoxic T lymphocyte-induced cytotoxicity.
Alphaherpesvirus-mediated remodeling of the cellular transcriptome results in depletion of m6A-containing transcripts.
Herpes simplex virus downregulates secretory leukocyte protease inhibitor: a novel immune evasion mechanism.
ICP47 mediates viral neuroinvasiveness by induction of TAP protein following intravenous inoculation of herpes simplex virus type 1 in mice.
Oncolytic herpes simplex virus interactions with the host immune system.
Herpes Simplex Virus 1 VP22 Inhibits AIM2-Dependent Inflammasome Activation to Enable Efficient Viral Replication.
Herpes simplex virus 1 infection of T cells causes VP11/12-dependent phosphorylation and degradation of the cellular protein Dok-2.
Corrigendum: A tug of war: DNA-sensing antiviral innate immunity and herpes simplex virus type 1 infection.
Immunization with a vaccine combining herpes simplex virus 2 (HSV-2) glycoprotein C (gC) and gD subunits improves the protection of dorsal root ganglia in mice and reduces the frequency of recurrent vaginal shedding of HSV-2 DNA in guinea pigs compared to immunization with gD alone.
Role of L-Particles during Herpes Simplex Virus Infection.
Immunization strategies to block the herpes simplex virus type 1 immunoglobulin G Fc receptor.
Loss of ICP22 in HSV-1 Elicits Immune Infiltration and Maintains Stromal Keratitis Despite Reduced Primary and Latent Virus Infectivity.
"Non-Essential" Proteins of HSV-1 with Essential Roles In Vivo: A Comprehensive Review.
Modulation of HLA-G and HLA-E expression in human neuronal cells after rabies virus or herpes virus simplex type 1 infections.
Herpes simplex virus blocks Fas-mediated apoptosis independent of viral activation of NF-kappaB in human epithelial HEp-2 cells.
Varicella-zoster virus vaccine, successes and difficulties.
Herpes Simplex Virus 1 Tegument Protein UL41 Counteracts IFIT3 Antiviral Innate Immunity.
Herpes simplex virus type 1 glycoprotein gC mediates immune evasion in vivo.
A Thr/Ser dual residue motif in the cytoplasmic tail of human CD1d is important for the down-regulation of antigen presentation following a herpes simplex virus 1 infection.
[The functions of alpha herpesvirus gene products].
A trivalent gC2/gD2/gE2 vaccine for herpes simplex virus generates antibody responses that block immune evasion domains on gC2 better than natural infection.
A trivalent subunit antigen glycoprotein vaccine as immunotherapy for genital herpes in the guinea pig genital infection model.
HVEM signaling promotes protective antibody-dependent cellular cytotoxicity (ADCC) vaccine responses to herpes simplex viruses.
The Latency-Associated Transcript Inhibits Apoptosis via Downregulation of Components of the Type I Interferon Pathway during Latent Herpes Simplex Virus 1 Ocular Infection.
Herpes simplex virus 1 evades cellular antiviral response by inducing microRNA-24, which attenuates STING synthesis.
A VP22-Null HSV-1 Is Impaired in Inhibiting CD1d-Mediated Antigen Presentation.
Suppression of CD80 Expression by ICP22 Affects Herpes Simplex Virus Type 1 Replication and CD8(+)IFN-γ(+) Infiltrates in the Eyes of Infected Mice but Not Latency Reactivation.
Structure of the C-Terminal Domain of the Multifunctional ICP27 Protein from Herpes Simplex Virus 1.

HSV-1-Specific IgG(3) Titers Correlate with Brain Cortical Thinning in Individuals with Mild Cognitive Impairment and Alzheimer's Disease.
Herpes simplex virus type-1 amplicon vectors for vaccine generation in acute lymphoblastic leukemia.
Interaction of large DNA viruses with dendritic cells.
Nucleoside-modified mRNA encoding HSV-2 glycoproteins C, D, and E prevents clinical and subclinical genital herpes.
Herpes simplex virus type 1 infection of activated cytotoxic T cells: Induction of fratricide as a mechanism of viral immune evasion.
Avoiding the void: cell-to-cell spread of human viruses.
HSV-1 Modulates IL-6 Receptor Expression on Human Dendritic Cells.
Hidden regulation of herpes simplex virus 1 pre-mRNA splicing and polyadenylation by virally encoded immediate early gene ICP27.
Modulation of HLA-G expression in human neural cells after neurotropic viral infections.
Immunobiology and host response.
Differential responses of murine vaginal and uterine epithelial cells prior to and following herpes simplex virus type 2 (HSV-2) infection.
HSV and glycoprotein J inhibit caspase activation and apoptosis induced by granzyme B or Fas.
Cost analysis of treatment strategies for the control of HSV-2 infection in the U.S.: A mathematical modeling-based case study.
CD1 antigen presentation by human dendritic cells as a target for herpes simplex virus immune evasion.
CD8 CTL from genital herpes simplex lesions: recognition of viral tegument and immediate early proteins and lysis of infected cutaneous cells.
Blocking Autophagy in M1 Macrophages Enhances Virus Replication and Eye Disease in Ocularly Infected Transgenic Mice.
Neurodegenerative proteinopathies associated with neuroinfections.
The virion host shutoff RNase plays a key role in blocking the activation of protein kinase R in cells infected with herpes simplex virus 1.
Inhibition of Indoleamine-2,3-dioxygenase (IDO) in Glioblastoma Cells by Oncolytic Herpes Simplex Virus.
HSV-1 ICP27 suppresses NF-kappaB activity by stabilizing IkappaBalpha.
L Particles Transmit Viral Proteins from Herpes Simplex Virus 1-Infected Mature Dendritic Cells to Uninfected Bystander Cells, Inducing CD83 Downmodulation.
Herpes simplex virus US3 protein kinase regulates host responses and determines neurovirulence.
Antiviral Potential of Naphthoquinones Derivatives Encapsulated within Liposomes.
gD-Independent Superinfection Exclusion of Alphaherpesviruses.
Basic amino acids as modulators of an O-linked glycosylation signal of the herpes simplex virus type 1 glycoprotein gC: functional roles in viral infectivity.
Exploiting Mucosal Immunity for Antiviral Vaccines.
Nuclear translocation and activation of the transcription factor NFAT is blocked by herpes simplex virus infection.
Evaluation of HSV-2 gE Binding to IgG-Fc and Application for Vaccine Development.
RIP1/RIP3 binding to HSV-1 ICP6 initiates necroptosis to restrict virus propagation in mice.
Fascinating Molecular and Immune Escape Mechanisms in the Treatment of STIs (Syphilis, Gonorrhea, Chlamydia, and Herpes Simplex).
Host shutoff activity of VHS and SOX-like proteins: role in viral survival and immune evasion.

Herpes simplex virus type 1 ICP0 induces CD83 degradation in mature dendritic cells independent of its E3 ubiquitin ligase function.
Modulation of CD112 by the alpha herpesvirus gD protein suppresses DNAM-1-dependent NK cell-mediated lysis of infected cells.
HSV-1 upregulates the ARE-binding protein tristetraprolin in a STAT1- and p38-dependent manner in mature dendritic cells.
LIGHTing up dendritic cell activation: Immune regulation and viral exploitation.
Virus-Mediated Suppression of the Antigen Presentation Molecule MR1.
A 2.5-kilobase deletion containing a cluster of nine microRNAs in the latency-associated-transcript locus of the pseudorabies virus affects the host response of porcine trigeminal ganglia during established latency.
The herpes virus Fc receptor gE-gI mediates antibody bipolar bridging to clear viral antigens from the cell surface.
HSV-2 glycoprotein gD targets the CC domain of tetherin and promotes tetherin degradation via lysosomal pathway.
Roles of the Different Isoforms of the Pseudorabies Virus Protein Kinase pUS3 in Nuclear Egress.
HSV1 MicroRNA Modulation of GPI Anchoring and Downstream Immune Evasion.
The genome of turkey herpesvirus.
Identification of an important immunological difference between virulent varicella-zoster virus and its avirulent vaccine: viral disruption of dendritic cell instruction.
Immunologic mechanisms of fingolimod and the role of immunosenescence in the risk of cryptococcal infection: A case report and review of literature.
Mapping the Complex Genetic Landscape of Human Neurons.

Table A2-22, Cluster 21

Cluster 21 focuses on SARS-CoV-2, its variants and mutations, and the mechanisms of its immune evasion/immune escape (1260)
The Immune Response to SARS-CoV-2 and Variants of Concern.
[Genomics and epidemiology of SARS-CoV-2 lineage].
Origin and evolution of SARS-CoV-2.
Research progress in spike mutations of SARS-CoV-2 variants and vaccine development.
The spike glycoprotein of SARS-CoV-2: A review of how mutations of spike glycoproteins have driven the emergence of variants with high transmissibility and immune escape.
High-Throughput, Single-Copy Sequencing Reveals SARS-CoV-2 Spike Variants Coincident with Mounting Humoral Immunity during Acute COVID-19.
Pre-Clinical Development of a Potent Neutralizing Antibody MW3321 With Extensive SARS-CoV-2 Variants Coverage.
Understanding the Secret of SARS-CoV-2 Variants of Concern/Interest and Immune Escape.
Facing the wrath of enigmatic mutations: a review on the emergence of severe acute respiratory syndrome coronavirus 2 variants amid coronavirus disease-19 pandemic.
Characterization of SARS-CoV-2 worldwide transmission based on evolutionary dynamics and specific viral mutations in the spike protein.
Immune evasion of SARS-CoV-2 from interferon antiviral system.
Genomic surveillance of severe acute respiratory syndrome coronavirus 2 in Burundi, from May 2021 to January 2022.

The global epidemic of SARS-CoV-2 variants and their mutational immune escape.
Glycosylation is a key in SARS-CoV-2 infection.
All hands on deck: SARS-CoV-2 proteins that block early anti-viral interferon responses.
SARS-CoV-2 epitopes inform future vaccination strategies.
The evolution of SARS-CoV-2.
Cell entry mechanisms of SARS-CoV-2.
Immune evasion of severe acute respiratory syndrome coronavirus-2 (SARS-CoV-2); molecular approaches.
Immune response and potential therapeutic strategies for the SARS-CoV-2 associated with the COVID-19 pandemic.
Structural Comparison and Drug Screening of Spike Proteins of Ten SARS-CoV-2 Variants.
SARS-CoV-2: An Overview of the Genetic Profile and Vaccine Effectiveness of the Five Variants of Concern.
SARS-CoV-2 variants, spike mutations and immune escape.
How do the severe acute respiratory coronavirus 2 (SARS-CoV-2) and its variants escape the host protective immunity and mediate pathogenesis?
Transmissibility and pathogenicity of SARS-CoV-2 variants in animal models.
The mechanisms of immune response and evasion by the main SARS-CoV-2 variants.
Multiple SARS-CoV-2 Variants Exhibit Variable Target Cell Infectivity and Ability to Evade Antibody Neutralization.
Induction of Broadly Cross-Reactive Antibody Responses to SARS-CoV-2 Variants by S1 Nanoparticle Vaccines.
Long-term effects of SARS-CoV-2 infection on human brain and memory.
A recombinant subunit vaccine candidate produced in plants elicits neutralizing antibodies against SARS-CoV-2 variants in macaques.
Variants of Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) and Vaccine Effectiveness.
The Role of the SARS-CoV-2 S-Protein Glycosylation in the Interaction of SARS-CoV-2/ACE2 and Immunological Responses.
When RING finger family proteins meet SARS-CoV-2.
The Evolution and Biology of SARS-CoV-2 Variants.
Evolution of SARS-CoV-2 Variants: Implications on Immune Escape, Vaccination, Therapeutic and Diagnostic Strategies.
Interaction of SARS-CoV-2 with host cells and antibodies: experiment and simulation.
Panorama of Breakthrough Infection Caused by SARS-CoV-2: A Review.
SARS-CoV-2 Mutations and their Viral Variants.
SARS-CoV-2 within-host diversity of human hosts and its implications for viral immune evasion.
The battle between host and SARS-CoV-2: Innate immunity and viral evasion strategies.
Evolution of the SARS-CoV-2 pandemic in India.
A Vaccine with Multiple Receptor-Binding Domain Subunit Mutations Induces Broad-Spectrum Immune Response against SARS-CoV-2 Variants of Concern.
SARS-CoV-2 variants and COVID-19 vaccines: Current challenges and future strategies.
Binding and neutralizing abilities of antibodies towards SARS-CoV-2 S2 domain.
Epidemiology and Characteristics of SARS-CoV-2 Variants of Concern: The Impacts of the Spike Mutations.

Structural Study of SARS-CoV-2 Antibodies Identifies a Broad-Spectrum Antibody That Neutralizes the Omicron Variant by Disassembling the Spike Trimer.
Functional mutations of SARS-CoV-2: implications to viral transmission, pathogenicity and immune escape.
Sensitivity of SARS-CoV-2 Variants to Neutralization by Convalescent Sera and a VH3-30 Monoclonal Antibody.
Towards SARS-CoV-2 serotypes?
Enhanced recombination among Omicron subvariants of SARS-CoV-2 contributes to viral immune escape.
Cross-neutralization of SARS-CoV-2 by a human monoclonal SARS-CoV antibody.
SARS-CoV-2 Vaccines Based on the Spike Glycoprotein and Implications of New Viral Variants.
Manipulation of innate immune signaling pathways by SARS-CoV-2 non-structural proteins.
Target-Specific Drug Discovery of Natural Products against SARS-CoV-2 Life Cycle and Cytokine Storm in COVID-19.
SARS-CoV-2 variants and effectiveness of vaccines: a review of current evidence.
Clinical-Pathological Correlation of the Pathophysiology and Mechanism of Action of COVID-19 - a Primer for Clinicians.
COVID-19 Challenge: A Quest for Effective Vaccine Strategies Against Circulating and Emerging SARS-CoV-2 Variants.
Stable and durable antibody responses in SARS-recovered donors vaccinated with inactivated SARS-CoV-2 vaccine.
Characterization of SARS-CoV-2 Evasion: Interferon Pathway and Therapeutic Options.
Structural and antigenic variations in the spike protein of emerging SARS-CoV-2 variants.
SARS-CoV-2 variant biology: immune escape, transmission and fitness.
Novel receptor, mutation, vaccine, and establishment of coping mode for SARS-CoV-2: current status and future.
The Development of SARS-CoV-2 Variants: The Gene Makes the Disease.
SARS-CoV-2-mediated evasion strategies for antiviral interferon pathways.
Population Immunity to Pre-Omicron and Omicron Severe Acute Respiratory Syndrome Coronavirus 2 Variants in US States and Counties Through 1 December 2021.
Effects of common mutations in the SARS-CoV-2 Spike RBD and its ligand, the human ACE2 receptor on binding affinity and kinetics.
Emerging Variants of SARS-CoV-2 and Novel Therapeutics Against Coronavirus (COVID-19).
Epigenetic Regulator miRNA Pattern Differences Among SARS-CoV, SARS-CoV-2, and SARS-CoV-2 World-Wide Isolates Delineated the Mystery Behind the Epic Pathogenicity and Distinct Clinical Characteristics of Pandemic COVID-19.
Ongoing Positive Selection Drives the Evolution of SARS-CoV-2 Genomes.
SARS-CoV-2 and Emerging Foodborne Pathogens: Intriguing Commonalities and Obvious Differences.
Analysis and comparison of physiochemical properties, mutations and glycosylation patterns between RNA polymerase and membrane protein of SARS-CoV and SARS-CoV-2.
An ACE2-blocking antibody confers broad neutralization and protection against Omicron and other SARS-CoV-2 variants of concern.
The mutational dynamics of the SARS-CoV-2 virus in serial passages in vitro.
Emergence of immune escape at dominant SARS-CoV-2 killer T cell epitope.
Myocardial Damage by SARS-CoV-2: Emerging Mechanisms and Therapies.
Dendritic cells in COVID-19 immunopathogenesis: insights for a possible role in determining disease outcome.

Broadly neutralizing antibodies derived from the earliest COVID-19 convalescents protect mice from SARS-CoV-2 variants challenge.
Antibody resistance of SARS-CoV-2 variants B.1.351 and B.1.1.7.
Updated Insights into the T Cell-Mediated Immune Response against SARS-CoV-2: A Step towards Efficient and Reliable Vaccines.
Understanding the implications of SARS-CoV-2 re-infections on immune response milieu, laboratory tests and control measures against COVID-19.
Analysis of SARS-CoV-2 in Nasopharyngeal Samples from Patients with COVID-19 Illustrates Population Variation and Diverse Phenotypes, Placing the Growth Properties of Variants of Concern in Context with Other Lineages.
Efficacy of Parainfluenza Virus 5 (PIV5)-vectored Intranasal COVID-19 Vaccine as a Single Dose Vaccine and as a Booster against SARS-CoV-2 Variants.
Resilience of Spike-Specific Immunity Induced by COVID-19 Vaccines against SARS-CoV-2 Variants.
The paradigm of immune escape by SARS-CoV-2 variants and strategies for repositioning subverted mAbs against escaped VOCs.
Immunopathology and immunotherapeutic strategies in severe acute respiratory syndrome coronavirus 2 infection.
SARS-CoV-2: Evolution and Emergence of New Viral Variants.
Modeling coronavirus spike protein dynamics: implications for immunogenicity and immune escape.
Antagonism of Type I Interferon by Severe Acute Respiratory Syndrome Coronavirus 2.
[Omicron Variant of SARS-CoV-2 and Its Potential Impact on Dental Practice].
Antibody neutralization to SARS-CoV-2 and variants after 1 year in Wuhan, China.
Epigenetic Lens to Visualize the Severe Acute Respiratory Syndrome Coronavirus-2 (SARS-CoV-2) Infection in COVID-19 Pandemic.
A non-ACE2 competing human single-domain antibody confers broad neutralization against SARS-CoV-2 and circulating variants.
Mitochondria: In the Cross Fire of SARS-CoV-2 and Immunity.
Are the original SARS-CoV-2 novel mutants from in vitro culture able to escape the immune response?
The ins and outs of SARS-CoV-2 variants of concern (VOCs).
Reduced neutralization of SARS-CoV-2 B.1.617 variant by convalescent and vaccinated sera.
Novel SARS-CoV-2 variants: the pandemics within the pandemic.
Antigenic evolution of SARS coronavirus 2.
CD8(+) T-cell immune escape by SARS-CoV-2 variants of concern.
Immunoinformatic analysis of structural and epitope variations in the spike and Orf8 proteins of SARS-CoV-2/B.1.1.7.
Neutralizing and enhancing antibodies against SARS-CoV-2.
The Pathogenic Features of Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2): Possible Mechanisms for Immune Evasion?
Tracking the Genomic Evolution of SARS-CoV-2 for 29 Months in South Korea.
Pathogenesis, Symptomatology, and Transmission of SARS-CoV-2 through Analysis of Viral Genomics and Structure.
Distribution and Functional Analyses of Mutations in Spike Protein and Phylogenic Diversity of SARS-CoV-2 Variants Emerged during the Year 2021 in India.
Differences and similarities between SARS-CoV and SARS-CoV-2: spike receptor-binding domain recognition and host cell infection with support of cellular serine proteases.
CD147 antibody specifically and effectively inhibits infection and cytokine storm of SARS-CoV-2 and its variants delta, alpha, beta, and gamma.



Persistent viral RNA positivity during the recovery period of a patient with SARS-CoV-2 infection.
Evaluation of T cell responses to naturally processed variant SARS-CoV-2 spike antigens in individuals following infection or vaccination.
Protein post-translational modification in SARS-CoV-2 and host interaction.
A Possible Role of Remdesivir and Plasma Therapy in the Selective Sweep and Emergence of New SARS-CoV-2 Variants.
Mutations and Evolution of the SARS-CoV-2 Spike Protein.
SARS-CoV-2 spike S2-specific neutralizing antibodies.
Neurotropism of SARS-CoV-2 and neurological diseases of the central nervous system in COVID-19 patients.
Structural Plasticity and Immune Evasion of SARS-CoV-2 Spike Variants.
Diverse Humoral Immune Responses in Younger and Older Adult COVID-19 Patients.
On the evolution of SARS-CoV-2 and the emergence of variants of concern.
Lack of SARS-CoV-2 Viral RNA Detection among a Convenience Sampling of Ohio Wildlife, Companion, and Agricultural Animals, 2020-2021.
The effect of spike mutations on SARS-CoV-2 neutralization.
CROI 2021: Summary of Basic Science Research in HIV and SARS-CoV-2.
Notable and Emerging Variants of SARS-CoV-2 Virus: A Quick Glance.
Molecular mechanisms involved in pathogenicity of SARS-CoV-2: Immune evasion and implications for therapeutic strategies.
Molecular characteristics, immune evasion, and impact of SARS-CoV-2 variants.
Epidemic history and evolution of an emerging threat of international concern, the severe acute respiratory syndrome coronavirus 2.
The effects of SARS-CoV-2 infection on modulating innate immunity and strategies of combating inflammatory response for COVID-19 therapy.
Molecular switches regulating the potency and immune evasiveness of SARS-CoV-2 spike protein.
HIV-1 and SARS-CoV-2: Patterns in the evolution of two pandemic pathogens.
Effects of SARS-CoV-2 variants on vaccine efficacy and response strategies.
SARS-CoV-2-mediated immune system activation and potential application in immunotherapy.
SARS-CoV-2 Entry Related Viral and Host Genetic Variations: Implications on COVID-19 Severity, Immune Escape, and Infectivity.
Genomic surveillance of genes encoding the SARS-CoV-2 spike protein to monitor for emerging variants on Jeju Island, Republic of Korea.
Dynamical changes of SARS-CoV-2 spike variants in the highly immunogenic regions impact the viral antibodies escaping.
Global Prevalence of Adaptive and Prolonged Infections' Mutations in the Receptor-Binding Domain of the SARS-CoV-2 Spike Protein.
Efficient SARS-CoV-2 Surveillance during the Pandemic-Endemic Transition Using PCR-Based Genotyping Assays.
Analyses of Long-Term Epidemic Trends and Evolution Characteristics of Haplotype Subtypes Reveal the Dynamic Selection on SARS-CoV-2.
SARS-CoV-2 Variants and Clinical Outcomes: A Systematic Review.
PARIS and SPARTA: Finding the Achilles' Heel of SARS-CoV-2.
The Importance of Glycosylation in COVID-19 Infection.
Interferon induction, evasion, and paradoxical roles during SARS-CoV-2 infection.
Mutational hotspots and conserved domains of SARS-CoV-2 genome in African population.

Refocus on Immunogenic Characteristics of Convalescent COVID-19 Challenged by Prototype SARS-CoV-2.
Computational Analysis of Mutations in the Receptor-Binding Domain of SARS-CoV-2 Spike and Their Effects on Antibody Binding.
Structural and physiological changes of the human body upon SARS-CoV-2 infection.
Immune Evasive Effects of SARS-CoV-2 Variants to COVID-19 Emergency Used Vaccines.
Characterization of SARS-CoV-2 Spike mutations important for infection of mice and escape from human immune sera.
Innate Immune Responses to Highly Pathogenic Coronaviruses and Other Significant Respiratory Viral Infections.
Relatively rapid evolution rates of SARS-CoV-2 spike gene at the primary stage of massive vaccination.
Potential transmission chains of variant B.1.1.7 and co-mutations of SARS-CoV-2.
Host cell-intrinsic innate immune recognition of SARS-CoV-2.
After the pandemic: perspectives on the future trajectory of COVID-19.
N-terminal domain mutations of the spike protein are structurally implicated in epitope recognition in emerging SARS-CoV-2 strains.
[Not Available].
Biology and Pathogenesis of SARS-CoV-2: Understandings for Therapeutic Developments against COVID-19.
Spike protein receptor-binding domains from SARS-CoV-2 variants of interest bind human ACE2 more tightly than the prototype spike protein.
Deep phylogenetic-based clustering analysis uncovers new and shared mutations in SARS-CoV-2 variants as a result of directional and convergent evolution.
Vive la Résistance: T-cell Immunity in the Protection against SARS-CoV-2 Infection.
Emergence of Multiple SARS-CoV-2 Antibody Escape Variants in an Immunocompromised Host Undergoing Convalescent Plasma Treatment.
A Methyltransferase-Defective Vesicular Stomatitis Virus-Based SARS-CoV-2 Vaccine Candidate Provides Complete Protection against SARS-CoV-2 Infection in Hamsters.
From SARS to SARS-CoV-2, insights on structure, pathogenicity and immunity aspects of pandemic human coronaviruses.
SARS-CoV-2 Delta variant genomic variation associated with breakthrough infection in Northern California: A retrospective cohort study.
Genomic Surveillance Enables the Identification of Co-infections With Multiple SARS-CoV-2 Lineages in Equatorial Guinea.
Innate and adaptive immunity to SARS-CoV-2 and predisposing factors.
Streptococcus pneumoniae colonization associates with impaired adaptive immune responses against SARS-CoV-2.
Structural Evaluation of the Spike Glycoprotein Variants on SARS-CoV-2 Transmission and Immune Evasion.
Clinical application of SARS-CoV-2 antibody detection and monoclonal antibody therapies against COVID-19.
The immunologic response to severe acute respiratory syndrome coronavirus 2.
Host microRNA interactions with the SARS-CoV-2 viral genome 3'-untranslated region.
SARS-CoV-2 one year on: evidence for ongoing viral adaptation.
TMPRSS2 promotes SARS-CoV-2 evasion from NCOA7-mediated restriction.
A novel mAb broadly neutralizes SARS-CoV-2 VOCs in vitro and in vivo, including the Omicron variants.

Within-host evolution of SARS-CoV-2 in an immunosuppressed COVID-19 patient as a source of immune escape variants.
Longitudinal and proteome-wide analyses of antibodies in COVID-19 patients reveal features of the humoral immune response to SARS-CoV-2.
Structure of SARS-CoV-2 ORF8, a rapidly evolving immune evasion protein.
SARS-CoV-2's Variants of Concern: A Brief Characterization.
SARS-CoV-2 Nsp5 Demonstrates Two Distinct Mechanisms Targeting RIG-I and MAVS To Evade the Innate Immune Response.
Long-Term Evolution of SARS-CoV-2 in an Immunocompromised Patient with Non-Hodgkin Lymphoma.
Neutral evolution test of the spike protein of SARS-CoV-2 and its implications in the binding to ACE2.
Host PDZ-containing proteins targeted by SARS-CoV-2.
Diverse SARS-CoV-2 aptamers overcome variant antigenic shift.
Covid-19: Perspectives on Innate Immune Evasion.
High genetic barrier to SARS-CoV-2 polyclonal neutralizing antibody escape.
SARS-CoV-2 infection activates a subset of intrinsic pathways to inhibit type I interferons in vitro and in vivo.
A novel heterologous receptor-binding domain dodecamer universal mRNA vaccine against SARS-CoV-2 variants.
Strategies and safety considerations of booster vaccination in COVID-19.
SARS-CoV-2 intra-host evolution during prolonged infection in an immunocompromised patient.
Appraisal of SARS-CoV-2 mutations and their impact on vaccination efficacy: an overview.
Activation and evasion of type I interferon responses by SARS-CoV-2.
Emerging roles of SARS-CoV-2 Spike-ACE2 in immune evasion and pathogenesis.
Nanobodies Protecting From Lethal SARS-CoV-2 Infection Target Receptor Binding Epitopes Preserved in Virus Variants Other Than Omicron.
SARS-CoV-2 human T cell epitopes: Adaptive immune response against COVID-19.
Immunoinformatic Analysis of T- and B-Cell Epitopes for SARS-CoV-2 Vaccine Design.
The immune response and immune evasion characteristics in SARS-CoV, MERS-CoV, and SARS-CoV-2: Vaccine design strategies.
SARS-CoV 2; Possible alternative virus receptors and pathophysiological determinants.
How SARS-CoV-2 dodges immune surveillance and facilitates infection: an analytical review.
A Comprehensive Molecular Epidemiological Analysis of SARS-CoV-2 Infection in Cyprus from April 2020 to January 2021: Evidence of a Highly Polyphyletic and Evolving Epidemic.
Discovery and Genomic Characterization of a 382-Nucleotide Deletion in ORF7b and ORF8 during the Early Evolution of SARS-CoV-2.
Brain cross-protection against SARS-CoV-2 variants by a lentiviral vaccine in new transgenic mice.
Cross protection to SARS-CoV-2 variants in hamsters with naturally-acquired immunity.
Recent advances in applying G-quadruplex for SARS-CoV-2 targeting and diagnosis: A review.
Mutations of SARS-CoV-2 spike protein: Implications on immune evasion and vaccine-induced immunity.
Host adaptation of codon usage in SARS-CoV-2 from mammals indicates potential natural selection and viral fitness.
Structural and functional insights into the spike protein mutations of emerging SARS-CoV-2 variants.
SARS-CoV-2 Mutations Responsible for Immune Evasion Leading to Breakthrough Infection.

Severe Acute Respiratory Syndrome Coronavirus 2 Variants of Concern: A Perspective for Emerging More Transmissible and Vaccine-Resistant Strains.
SARS-CoV-2 and the host-immune response.
Nanotechnologies for the Diagnosis and Treatment of SARS-CoV-2 and Its Variants.
Shooting at a Moving Target-Effectiveness and Emerging Challenges for SARS-CoV-2 Vaccine Development.
The ORF8 protein of SARS-CoV-2 mediates immune evasion through down-regulating MHC-I.
SARS-CoV-2 Omicron Variant in Medicinal Chemistry Research.
Conservation and Evolution of Antigenic Determinants of SARS-CoV-2: An Insight for Immune Escape and Vaccine Design.
Of bats and men: Immunomodulatory treatment options for COVID-19 guided by the immunopathology of SARS-CoV-2 infection.
A novel plant lectin, NTL-125, interferes with SARS-CoV-2 interaction with hACE2.
Structure-Function Analyses of New SARS-CoV-2 Variants B.1.1.7, B.1.351 and B.1.1.28.1: Clinical, Diagnostic, Therapeutic and Public Health Implications.
SARS-CoV-2 and Emerging Variants: Unmasking Structure, Function, Infection, and Immune Escape Mechanisms.
Decreased and Heterogeneous Neutralizing Antibody Responses Against RBD of SARS-CoV-2 Variants After mRNA Vaccination.
Novel recombinant SARS-CoV-2 lineage detected through genomic surveillance in Wales, UK.
Structural basis for continued antibody evasion by the SARS-CoV-2 receptor binding domain.
Quantitative Mutation Analysis of Genes and Proteins of Major SARS-CoV-2 Variants of Concern and Interest.
Ocular COVID-19: Eyes as a Reservoir to Conceal and Spread SARS-CoV-2.
Insight into the emerging role of SARS-CoV-2 nonstructural and accessory proteins in modulation of multiple mechanisms of host innate defense.
Dermal Delivery of a SARS-CoV-2 Subunit Vaccine Induces Immunogenicity against Variants of Concern.
SARS-CoV-2 within-host diversity and transmission.
Elicitation of Broadly Neutralizing Antibodies against B.1.1.7, B.1.351, and B.1.617.1 SARS-CoV-2 Variants by Three Prototype Strain-Derived Recombinant Protein Vaccines.
SARS-CoV-2 Non-Structural Proteins and Their Roles in Host Immune Evasion.
The Genetic Variant of SARS-CoV-2: would It Matter for Controlling the Devastating Pandemic?
Applications of nanobodies in the prevention, detection, and treatment of the evolving SARS-CoV-2.
Distinct sensitivities to SARS-CoV-2 variants in vaccinated humans and mice.
SARS-CoV-2 protein ORF8 limits expression levels of Spike antigen and facilitates immune evasion of infected host cells.
Evaluating the effect of SARS-CoV-2 spike mutations with a linear doubly robust learner.
RG203KR Mutations in SARS-CoV-2 Nucleocapsid: Assessing the Impact Using a Virus-Like Particle Model System.
Expanding repertoire of SARS-CoV-2 deletion mutations contributes to evolution of highly transmissible variants.
Discovery of re-purposed drugs that slow SARS-CoV-2 replication in human cells.
Successful treatment of persistent SARS-CoV-2 infection with nirmatrelvir/ritonavir plus sotrovimab in four immunocompromised patients.
Type I and III IFN-mediated antiviral actions counteracted by SARS-CoV-2 proteins and host inherited factors.

The ongoing evolution of variants of concern and interest of SARS-CoV-2 in Brazil revealed by convergent indels in the amino (N)-terminal domain of the spike protein.
Advances in developing ACE2 derivatives against SARS-CoV-2.
The structure of the RBD-E77 Fab complex reveals neutralization and immune escape of SARS-CoV-2.
A comparative review of pathogenesis and host innate immunity evasion strategies among the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), severe acute respiratory syndrome coronavirus (SARS-CoV) and the Middle East respiratory syndrome coronavirus (MERS-CoV).
Long-term passaging of pseudo-typed SARS-CoV-2 reveals the breadth of monoclonal and bispecific antibody cocktails.
Transmission event of SARS-CoV-2 Delta variant reveals multiple vaccine breakthrough infections.
Ongoing global and regional adaptive evolution of SARS-CoV-2.
[Interaction between SARS-CoV-2 and Host Innate Immunity].
An ACE2-Based Decoy Inhibitor Effectively Neutralizes SARS-CoV-2 Omicron BA.5 Variant.
Selection for immune evasion in SARS-CoV-2 revealed by high-resolution epitope mapping and sequence analysis.
Emergence of a recurrent insertion in the N-terminal domain of the SARS-CoV-2 spike glycoprotein.
An update on host immunity correlates and prospects of re-infection in COVID-19.
Adenovirus 5 Vectors Expressing SARS-CoV-2 Proteins.
SARS-CoV-2 Variant Spike and accessory gene mutations alter pathogenesis.
SARS-CoV-2 ORF8: A Rapidly Evolving Immune and Viral Modulator in COVID-19.
Evolutionary Tracking of SARS-CoV-2 Genetic Variants Highlights an Intricate Balance of Stabilizing and Destabilizing Mutations.
Emerging SARS-CoV-2 variants can potentially break set epidemiological barriers in COVID-19.
A comprehensive profile of genomic variations in the SARS-CoV-2 isolates from the state of Telangana, India.
Detecting SARS-CoV-2 neutralizing immunity: highlighting the potential of split nanoluciferase technology.
Persistent SARS-CoV-2 infection and the risk for cancer.
Pathogenesis, Symptomatology, and Transmission of SARS-CoV-2 through Analysis of Viral Genomics and Structure.
SARS-CoV-2 antibody prevalence in a pediatric cohort of unvaccinated children in Mérida, Yucatán, México.
Nanobody repertoire generated against the spike protein of ancestral SARS-CoV-2 remains efficacious against the rapidly evolving virus.
Immunopathology and Immunopathogenesis of COVID-19, what we know and what we should learn.
Modeling of ACE2 and antibodies bound to SARS-CoV-2 provides insights into infectivity and immune evasion.
Emerging Mutations Potentially Related to SARS-CoV-2 Immune Escape: The Case of a Long-Term Patient.
Ten emerging SARS-CoV-2 spike variants exhibit variable infectivity, animal tropism, and antibody neutralization.
Suramin inhibits SARS-CoV-2 nucleocapsid phosphoprotein genome packaging function.
Transmission event of SARS-CoV-2 delta variant reveals multiple vaccine breakthrough infections.
Host genetic diversity and genetic variations of SARS-CoV-2 in COVID-19 pathogenesis and the effectiveness of vaccination.
Bispecific IgG neutralizes SARS-CoV-2 variants and prevents escape in mice.
Shedding of infectious SARS-CoV-2 despite vaccination.

Delta variant (B.1.617.2) of SARS-CoV-2: current understanding of infection, transmission, immune escape, and mutational landscape.
An Update on Innate Immune Responses during SARS-CoV-2 Infection.
Innate immune evasion by SARS-CoV-2: Comparison with SARS-CoV.
The regulation of lncRNAs and miRNAs in SARS-CoV-2 infection.
Genomic analysis of SARS-CoV-2 variants: diagnosis and vaccination challenges.
Type I and III interferon responses in SARS-CoV-2 infection.
Evidence of recurrent selection of mutations commonly found in SARS-CoV-2 variants of concern in viruses infecting immunocompromised patients.
Generation of Novel Severe Acute Respiratory Syndrome Coronavirus 2 Variants on the B.1.1.7 Lineage in 3 Patients With Advanced Human Immunodeficiency Virus-1 Disease.
Understanding Mutations in Human SARS-CoV-2 Spike Glycoprotein: A Systematic Review & Meta-Analysis.
De novo emergence of SARS-CoV-2 spike mutations in immunosuppressed patients.
Neutralization of alpha, gamma, and D614G SARS-CoV-2 variants by CoronaVac vaccine-induced antibodies.
The past, current and future epidemiological dynamic of SARS-CoV-2.
Systematic functional analysis of SARS-CoV-2 proteins uncovers viral innate immune antagonists and remaining vulnerabilities.
SARS-CoV-2 impairs interferon production via NSP2-induced repression of mRNA translation.
Prediction and Evolution of the Molecular Fitness of SARS-CoV-2 Variants: Introducing SpikePro.
Structure of SARS-CoV-2 ORF8, a rapidly evolving coronavirus protein implicated in immune evasion.
Analysis of SARS-CoV-2 Spike Protein Variants with Recombinant Reporter Viruses Created from a Bacmid System.
Site-specific glycan analysis of the SARS-CoV-2 spike.
Advancing Precision Vaccinology by Molecular and Genomic Surveillance of Severe Acute Respiratory Syndrome Coronavirus 2 in Germany, 2021.
SARS-CoV-2 variants: Impact on biological and clinical outcome.
SARS-CoV-2 infects multiple species of North American deer mice and causes clinical disease in the California mouse.
COVID-19, Virology and Geroscience: A Perspective.
A bivalent nanoparticle vaccine exhibits potent cross-protection against the variants of SARS-CoV-2.
SARS-CoV-2 Shedding from Asymptomatic Patients: Contribution of Potential Extrapulmonary Tissue Reservoirs.
In vitro evolution predicts emerging SARS-CoV-2 mutations with high affinity for ACE2 and cross-species binding.
Immuno-epidemiology and pathophysiology of coronavirus disease 2019 (COVID-19).
SARS-CoV-2 variants and vaccination.
Signatures in SARS-CoV-2 spike protein conferring escape to neutralizing antibodies.
The pivotal roles of the host immune response in the fine-tuning the infection and the development of the vaccines for SARS-CoV-2.
Reduced sensitivity of SARS-CoV-2 variant Delta to antibody neutralization.
Dynamic Evolution of SARS-CoV-2 in a Patient on Chemotherapy.
COVID-19 Vaccines-All You Want to Know.
Taxonium, a web-based tool for exploring large phylogenetic trees.
Extremely potent monoclonal antibodies neutralize Omicron and other SARS-CoV-2 variants.

Rise and Fall of SARS-CoV-2 Lineage A.27 in Germany.
The Predictive Value of Mutation Screening for Anticipating COVID-19 Waves.
Allosteric regulation and crystallographic fragment screening of SARS-CoV-2 NSP15 endoribonuclease.
Targeting RdRp of SARS-CoV-2 with De Novo Molecule Generation.
Rapid Increase in SARS-CoV-2 P.1 Lineage Leading to Codominance with B.1.1.7 Lineage, British Columbia, Canada, January-April 2021.
SARS-CoV-2 accessory proteins ORF7a and ORF3a use distinct mechanisms to downregulate MHC-I surface expression.
SARS Coronavirus Fusion Peptide-Derived Sequence Suppresses Collagen-Induced Arthritis in DBA/1J Mice.
Old vaccines for new infections: Exploiting innate immunity to control COVID-19 and prevent future pandemics.
Neutralization of SARS-CoV-2 Variants of Concern in Kidney Transplant Recipients after Standard COVID-19 Vaccination.
Therapeutic effect of CT-P59 against SARS-CoV-2 South African variant.
Transcriptomic study reveals lncRNA-mediated downregulation of innate immune and inflammatory response in the SARS-CoV-2 vaccination breakthrough infections.
Middle East Respiratory Syndrome Coronavirus NS4b Protein Inhibits Host RNase L Activation.
Clinical Presentation of a COVID-19 Delta Variant Patient: Case Report and Literature Review.
Analysis of SARS-CoV-2 mutations associated with resistance to therapeutic monoclonal antibodies that emerge after treatment.
Maturation signatures of conventional dendritic cell subtypes in COVID-19 suggest direct viral sensing.
Cryo-electron Microscopy Structure of the Swine Acute Diarrhea Syndrome Coronavirus Spike Glycoprotein Provides Insights into Evolution of Unique Coronavirus Spike Proteins.
SARS-CoV-2 accessory proteins ORF7a and ORF3a use distinct mechanisms to down-regulate MHC-I surface expression.
Vaccines and variants: Modelling insights into emerging issues in COVID-19 epidemiology.
RdRp inhibitors and COVID-19: Is molnupiravir a good option?
Unraveling the Molecular Mechanism of Recognition of Human Interferon-Stimulated Gene Product 15 by Coronavirus Papain-Like Proteases: A Multiscale Simulation Study.
SARS-CoV-2 immune evasion by the B.1.427/B.1.429 variant of concern.
Vaccines alone will not prevent COVID-19 outbreaks among migrant workers-the example of meat processing plants.
Modulation of the immune response by Middle East respiratory syndrome coronavirus.
The mechanisms of catalysis and ligand binding for the SARS-CoV-2 NSP3 macrodomain from neutron and X-ray diffraction at room temperature.
Immune Responses During Human Coronavirus Infection: Suggestions for Future Studies.
Indirect-Acting Pan-Antivirals vs. Respiratory Viruses: A Fresh Perspective on Computational Multi-Target Drug Discovery.
Host Response to SARS-CoV2 and Emerging Variants in Pre-Existing Liver and Gastrointestinal Diseases.
Characterization of germline antibody libraries from human umbilical cord blood and selection of monoclonal antibodies to viral envelope glycoproteins: Implications for mechanisms of immune evasion and design of vaccine immunogens.
Extensive Positive Selection Drives the Evolution of Nonstructural Proteins in Lineage C Betacoronaviruses.

Recombinant Receptor-Binding Domains of Multiple Middle East Respiratory Syndrome Coronaviruses (MERS-CoVs) Induce Cross-Neutralizing Antibodies against Divergent Human and Camel MERS-CoVs and Antibody Escape Mutants.
Crystal structure of SARS-CoV-2 Orf9b in complex with human TOM70 suggests unusual virus-host interactions.
The emergence, impact, and evolution of human metapneumovirus variants from 2014 to 2021 in Spain.
Use of Whole-Genome Sequencing to Estimate the Contribution of Immune Evasion and Waning Immunity on Decreasing COVID-19 Vaccine Effectiveness.
New Discovery of Myeloid-Derived Suppressor Cell's Tale on Viral Infection and COVID-19.
Uncovering potential host proteins and pathways that may interact with eukaryotic short linear motifs in viral proteins of MERS, SARS and SARS2 coronaviruses that infect humans.
Antibody response and intra-host viral evolution after plasma therapy in COVID-19 patients pre-exposed or not to B-cell-depleting agents.
Viral Nephropathies, Adding SARS-CoV-2 to the List.
Early peripheral blood MCEMP1 and HLA-DRA expression predicts COVID-19 prognosis.
Biochemical and structural insights into the mechanisms of SARS coronavirus RNA ribose 2'-O-methylation by nsp16/nsp10 protein complex.
Structure-Based Discovery of Inhibitors of the SARS-CoV-2 Nsp14 N7-Methyltransferase.
Sustainable policy: Don't get infected and don't infect others.
Pharmacotherapeutic Potential of Natural Products to Target the SARS-CoV-2 PLpro Using Molecular Screening and Simulation Approaches.
Test-trace-isolate-quarantine (TTIQ) intervention strategies after symptomatic COVID-19 case identification.
Room-temperature structural studies of SARS-CoV-2 protein NendoU with an X-ray free-electron laser.
Substantial Impact of Post Vaccination Contacts on Cumulative Infections during Viral Epidemics.
SARS-CoV-2 Lineage Tracking, and Evolving Trends Seen during Three Consecutive Peaks of Infection in Delhi, India: a Clinico-Genomic Study.
Middle East respiratory syndrome coronavirus M protein suppresses type I interferon expression through the inhibition of TBK1-dependent phosphorylation of IRF3.
Exercise, Immune System, Nutrition, Respiratory and Cardiovascular Diseases during COVID-19: A Complex Combination.
Mercapto-pyrimidines are reversible covalent inhibitors of the papain-like protease (PLpro) and inhibit SARS-CoV-2 (SCoV-2) replication.
Bat severe acute respiratory syndrome-like coronavirus ORF3b homologues display different interferon antagonist activities.
Gain-of-function research.
The SARS coronavirus S glycoprotein receptor binding domain: fine mapping and functional characterization.
Local memory CD4 T cell niches in respiratory viral infection.
The human coronavirus HCoV-229E S-protein structure and receptor binding.
Cryo-Electron Microscopy Structure of Porcine Deltacoronavirus Spike Protein in the Prefusion State.
Middle east respiratory syndrome coronavirus 4a protein is a double-stranded RNA-binding protein that suppresses PACT-induced activation of RIG-I and MDA5 in the innate antiviral response.
Artificial intelligence-based HDX (AI-HDX) prediction reveals fundamental characteristics to protein dynamics: Mechanisms on SARS-CoV-2 immune escape.



Glycan shield and epitope masking of a coronavirus spike protein observed by cryo-electron microscopy.
Plausibility of a third wave of COVID-19 in India: A mathematical modelling based analysis.
Do nutrients and other bioactive molecules from foods have anything to say in the treatment against COVID-19?
COVID-19 Pandemic on Fire: Evolved Propensities for Nocturnal Activities as a Liability Against Epidemiological Control.
Ultrastructural study confirms the formation of single and heterotypic syncytial cells in bronchoalveolar fluids of COVID-19 patients.
Flipped over U: structural basis for dsRNA cleavage by the SARS-CoV-2 endoribonuclease.
Secure reversal of immune evasion from refractory NSCLC and highly contagious CoV-2 mutants by using 3D-engineered multifunctional biologics.
Middle East respiratory syndrome coronavirus-encoded ORF8b strongly antagonizes IFN- $\beta$ promoter activation: its implication for vaccine design.
Spike Gene Evolution and Immune Escape Mutations in Patients with Mild or Moderate Forms of COVID-19 and Treated with Monoclonal Antibodies Therapies.
[Advances in the reverse genetics system for RNA viruses].
Resolving the enigma of Iquitos and Manaus: A modeling analysis of multiple COVID-19 epidemic waves in two Amazonian cities.
Enhancing immune protection against MERS-CoV: the synergistic effect of proteolytic cleavage sites and the fusion peptide and RBD domain targeting VLP immunization.
A conserved region of nonstructural protein 1 from alphacoronaviruses inhibits host gene expression and is critical for viral virulence.
Cellular Deubiquitylating Enzyme: A Regulatory Factor of Antiviral Innate Immunity.
Structural insights into the Middle East respiratory syndrome coronavirus 4a protein and its dsRNA binding mechanism.
Pandemics and the English Language: Concepts Critical for Conversing About COVID-19.
SPEAR: Systematic ProtEin AnnotatoR.
Molecular mimicry of ACTH in SARS - implications for corticosteroid treatment and prophylaxis.
Epidemiological impact and cost-effectiveness analysis of COVID-19 vaccination in Kenya.
Risk assessment for long- and short-range airborne transmission of SARS-CoV-2, indoors and outdoors.
Untargeted saliva metabolomics by liquid chromatography-Mass spectrometry reveals markers of COVID-19 severity.
Neutrophil Extracellular Traps and By-Products Play a Key Role in COVID-19: Pathogenesis, Risk Factors, and Therapy.
Porcine Epidemic Diarrhea Virus Deficient in RNA Cap Guanine-N-7 Methylation Is Attenuated and Induces Higher Type I and III Interferon Responses.
The emergence of a virus variant: dynamics of a competition model with cross-immunity time-delay validated by wastewater surveillance data for COVID-19.
Efficient replication of the novel human betacoronavirus EMC on primary human epithelium highlights its zoonotic potential.
Strategies for controlling the innate immune activity of conventional and self-amplifying mRNA therapeutics: Getting the message across.
Strategies for the Emergency Treatment of Pregnant Women with Neurological Symptoms during the COVID-19 Pandemic.

Conformational ensemble of the NSP1 CTD in SARS-CoV-2: Perspectives from the free energy landscape.
Emergent variant modeling of the serological repertoire to norovirus in young children.
Plasma Torquetenovirus (TTV) microRNAs and severity of COVID-19.
A herd immunity approach to the COVID-19 pandemic?
Characterization and comparison of novel adjuvants for a prefusion clamped MERS vaccine.
SARS-CoV-2 Nsp16 activation mechanism and a cryptic pocket with pan-coronavirus antiviral potential.
Engineering ACE2 decoy receptors to combat viral escapability.
Hydrogen-Deuterium Exchange Epitope Mapping of Glycosylated Epitopes Enabled by Online Immobilized Glycosidase.
An Updated Review of Porcine Deltacoronavirus in Terms of Prevalence, Pathogenicity, Pathogenesis and Antiviral Strategy.
The origin and evolution of emerged swine acute diarrhea syndrome coronavirus with zoonotic potential.
Immune Signature of COVID-19: In-Depth Reasons and Consequences of the Cytokine Storm.
Insight into the evolution of nidovirus endoribonuclease based on the finding that nsp15 from porcine Deltacoronavirus functions as a dimer.
Pathogenesis and Preventive Tactics of Immune-Mediated Non-Pulmonary COVID-19 in Children and Beyond.
The age distribution of mortality due to influenza: pandemic and peri-pandemic.
Pulmonary Embolism in COVID-19 Treated with VA-ECLS and Catheter tPA.
Acute respiratory distress syndrome precipitated by granulocyte colony-stimulating factor in undiagnosed <i>Pneumocystis jirovecii</i> pneumonia.

Table A2-23, Cluster 22

Cluster 22 focuses on immune responses to human papillomavirus, especially the evasion of host immune defenses by hpv (135)
Immune responses to human papillomavirus.
HPV-mediated Cervical Cancer: A Systematic Review on Immunological Basis, Molecular Biology, and Immune Evasion Mechanisms.
Immune responses against human papillomavirus (HPV) infection and evasion of host defense in cervical cancer.
HPV innate immunity.
Evasion of host immune defenses by human papillomavirus.
HPV modulation of host immune responses.
The Interplay between Antiviral Signalling and Carcinogenesis in Human Papillomavirus Infections.
Mechanisms used by human papillomaviruses to escape the host immune response.
Current strategies against persistent human papillomavirus infection (Review).
[Toward cervical cancer prevention: strategies employed in the development of HPV vaccines].
Host responses to infection with human papillomavirus.
The immune response to papillomavirus during infection persistence and regression.
The invisible enemy - how human papillomaviruses avoid recognition and clearance by the host immune system.
Genital human papillomavirus infections: current and prospective therapies.

Human papillomavirus can escape immune recognition through Langerhans cell phosphoinositide 3-kinase activation.
Immune evasion mechanisms of human papillomavirus: An update.
An Update on Human Papilloma Virus Vaccines: History, Types, Protection, and Efficacy.
Immune control of human papillomavirus (HPV) associated anogenital disease and potential for vaccination.
Alteration of the IFN-Pathway by Human Papillomavirus Proteins: Antiviral Immune Response Evasion Mechanism.
Human papillomavirus type 16 E6 and NF1-123 mislocalize immune signaling proteins and downregulate immune gene expression in keratinocytes.
Viral Modulation of TLRs and Cytokines and the Related Immunotherapies for HPV-Associated Cancers.
Mechanisms of virus immune evasion lead to development from chronic inflammation to cancer formation associated with human papillomavirus infection.
The role of RNA-binding proteins in the processing of mRNAs produced by carcinogenic papillomaviruses.
Human papillomavirus in the setting of immunodeficiency: Pathogenesis and the emergence of next-generation therapies to reduce the high associated cancer risk.
ADAR1 function affects HPV replication and is associated to recurrent human papillomavirus-induced dysplasia in HIV coinfecting individuals.
[Escape mechanisms to the innate immune response in HPV-associated cervical cancer].
Acquired immune response to oncogenic human papillomavirus associated with prophylactic cervical cancer vaccines.
Suppression of Stromal Interferon Signaling by Human Papillomavirus 16.
Pathogenic Role of Immune Evasion and Integration of Human Papillomavirus in Oropharyngeal Cancer.
[Immune response in cervical cancer. Strategies for the development of therapeutic vaccines].
Suppression of Langerhans cell activation is conserved amongst human papillomavirus $\alpha$ and $\beta$ genotypes, but not a $\mu$ genotype.
Human papillomavirus infection: biology, epidemiology, and prevention.
Biology of HPV Mediated Carcinogenesis and Tumor Progression.
Induction of Interferon Kappa in Human Papillomavirus 16 Infection by Transforming Growth Factor Beta-Induced Promoter Demethylation.
HPV E7 inhibits cell pyroptosis by promoting TRIM21-mediated degradation and ubiquitination of the IFI16 inflammasome.
Human Papillomaviruses; Epithelial Tropisms, and the Development of Neoplasia.
Epithelial cell responses to infection with human papillomavirus.
Engineering the AAV capsid to evade immune responses.
The Route to HPV-Associated Neoplastic Transformation: A Review of the Literature.
A major role for the minor capsid protein of human papillomavirus type 16 in immune escape.
Immune responses to human papilloma viruses.
Host control of human papillomavirus infection and disease.
Association of human papilloma virus with pterygia and ocular-surface squamous neoplasia.
Innate immunity and HPV: friends or foes.
HPV - immune response to infection and vaccination.
The human papillomavirus oncoproteins: a review of the host pathways targeted on the road to transformation.

Langerhans cells from women with cervical precancerous lesions become functionally responsive against human papillomavirus after activation with stabilized Poly-I:C.
Human Papillomavirus 16 E5 Inhibits Interferon Signaling and Supports Episomal Viral Maintenance.
Immune response to human papillomavirus after prophylactic vaccination with AS04-adjuvanted HPV-16/18 vaccine: improving upon nature.
Small But Increasingly Mighty: Latest Advances in AAV Vector Research, Design, and Evolution.
The human papillomavirus E7 oncoprotein as a regulator of transcription.
The human Papillomavirus twilight zone - Latency, immune control and subclinical infection.
Adeno-associated virus as a gene therapy vector: vector development, production and clinical applications.
Suppression of STAT-1 expression by human papillomaviruses is necessary for differentiation-dependent genome amplification and plasmid maintenance.
Adeno-Associated Viruses (AAV) and Host Immunity - A Race Between the Hare and the Hedgehog.
Persistent Human Papillomavirus Infection.
The JAK-STAT transcriptional regulator, STAT-5, activates the ATM DNA damage pathway to induce HPV 31 genome amplification upon epithelial differentiation.
Isolation and functional analysis of five HPV E6 variants with respect to p53 degradation.
Preclinical gene therapy studies for hemophilia using adeno-associated virus (AAV) vectors.
CpG-depleted adeno-associated virus vectors evade immune detection.
Cellular redox, cancer and human papillomavirus.
CD8(+) T-cell responses to adeno-associated virus capsid in humans.
Improving clinical efficacy of adeno associated vectors by rational capsid bioengineering.
Developing immunologically inert adeno-associated virus (AAV) vectors for gene therapy: possibilities and limitations.
The host response to lesions induced by human papillomavirus.
Screening of drugs to counteract human papillomavirus 16 E6 repression of E-cadherin expression.
Directed evolution of AAV mutants for enhanced gene delivery.
Lessons learnt in Japan from adverse reactions to the HPV vaccine: a medical ethics perspective.
Diverse tumorigenic consequences of human papillomavirus integration in primary oropharyngeal cancers.
T-cell response to human papillomavirus type 52 L1, E6, and E7 peptides in women with transient infection, cervical intraepithelial neoplasia, and invasive cancer.
Identification of Broadly Applicable Adeno-Associated Virus Vectors by Systematic Comparison of Commonly Used Capsid Variants In Vitro.
Tumourigenesis driven by the human papillomavirus type 16 Asian-American e6 variant in a three-dimensional keratinocyte model.
Display of Self-Peptide on Adeno-Associated Virus Capsid Decreases Phagocytic Uptake in Vitro.
HPV infection alters vaginal microbiome through down-regulating host mucosal innate peptides used by Lactobacilli as amino acid sources.
Neutralizing Antibody Evasion and Transduction with Purified Extracellular Vesicle-Enveloped Adeno-Associated Virus Vectors.
Expression of different L1 isoforms of Mastomys natalensis papillomavirus as mechanism to circumvent adaptive immunity.
Human papillomavirus virus-like particles do not activate Langerhans cells: a possible immune escape mechanism used by human papillomaviruses.
Can genes transduced by adeno-associated virus vectors elicit or evade an immune response?

AAV as an immunogen.
Modulation of apoptosis by human papillomavirus (HPV) oncoproteins.
IL-18 E42A mutant is resistant to the inhibitory effects of HPV-16 E6 and E7 oncogenes on the IL-18-mediated immune response.
The structure of adeno-associated virus serotype 3B (AAV-3B): insights into receptor binding and immune evasion.
Engineering adeno-associated viral vectors to evade innate immune and inflammatory responses.
Surfactant Protein A Impairs Genital HPV16 Pseudovirus Infection by Innate Immune Cell Activation in A Murine Model.
Structural characterization of an envelope-associated adeno-associated virus type 2 capsid.
Evading and overcoming AAV neutralization in gene therapy.
AAV's anatomy: roadmap for optimizing vectors for translational success.
Host defence and persistent human papillomavirus infection.
CD1d, a sentinel molecule bridging innate and adaptive immunity, is downregulated by the human papillomavirus (HPV) E5 protein: a possible mechanism for immune evasion by HPV.
Human papillomavirus E7 oncoprotein increases production of the anti-inflammatory interleukin-18 binding protein in keratinocytes.
Post-translational control of IL-1 $\beta$ via the human papillomavirus type 16 E6 oncoprotein: a novel mechanism of innate immune escape mediated by the E3-ubiquitin ligase E6-AP and p53.
BRD4S interacts with viral E2 protein to limit human papillomavirus late transcription.
A Single Surface-Exposed Amino Acid Determines Differential Neutralization of AAV1 and AAV6 by Human Alpha-Defensins.
The interferon-related developmental regulator 1 is used by human papillomavirus to suppress NF $\kappa$ B activation.
Enhanced Production of Exosome-Associated AAV by Overexpression of the Tetraspanin CD9.
Manipulation of the innate immune response by human papillomaviruses.
Animal models of papillomavirus pathogenesis.
Human papillomavirus 16 E6 variants are more prevalent in invasive cervical carcinoma than the prototype.
Expression and Purification of Adeno-associated Virus Virus-like Particles in a Baculovirus System and AAVR Ectodomain Constructs in E. coli.
Analysis of Adeno-Associated Virus Serotype 8 (AAV8)-antibody complexes using epitope mapping by molecular imprinting leads to the identification of Fab peptides that potentially evade AAV8 neutralisation.
Downregulation of Toll-Like Receptor 9 Expression by Beta Human Papillomavirus 38 and Implications for Cell Cycle Control.
Deregulation of E-cadherin by human papillomavirus is not confined to high-risk, cancer-causing types.
Evasion of immune responses to introduced human acid alpha-glucosidase by liver-restricted expression in glycogen storage disease type II.
YIP1 family member 4 (YIPF4) is a novel cellular binding partner of the papillomavirus E5 proteins.
Microarray analysis identifies interferon-inducible genes and Stat-1 as major transcriptional targets of human papillomavirus type 31.
Human papillomavirus type 16 antagonizes IRF6 regulation of IL-1 $\beta$ .
Overcoming Immunological Challenges Limiting Capsid-Mediated Gene Therapy With Machine Learning.
Papillomavirus capsid mutation to escape dendritic cell-dependent innate immunity in cervical cancer.

A synthetic E7 gene of human papillomavirus type 16 that yields enhanced expression of the protein in mammalian cells and is useful for DNA immunization studies.
Human papillomavirus-exposed Langerhans cells are activated by stabilized Poly-I:C.
Immune deviation and cervical carcinogenesis.
The immunology of genital human papilloma virus infection.
How not to be seen: immune-evasion strategies in gene therapy.
Mechanisms of persistence by small DNA tumor viruses.
Insidious Insights: Implications of viral vector engineering for pathogen enhancement.
High-risk human papillomavirus targets crossroads in immune signaling.
The Structure of an AAV5-AAVR Complex at 2.5 Å Resolution: Implications for Cellular Entry and Immune Neutralization of AAV Gene Therapy Vectors.
Isoforms of the Papillomavirus Major Capsid Protein Differ in Their Ability to Block Viral Spread and Tumor Formation.
Current strategies and future directions for eluding adenoviral vector immunity.
Improving adenovirus based gene transfer: strategies to accomplish immune evasion.
Current status and future directions of gene and cell therapy for cystic fibrosis.
Evolutionary and functional implications of hypervariable loci within the skin virome.
Contrasting effects of human, canine, and hybrid adenovirus vectors on the phenotypical and functional maturation of human dendritic cells: implications for clinical efficacy.
Vaccines: engineering immune evasion.
Modification of adenovirus gene transfer vectors with synthetic polymers: a scientific review and technical guide.
Regulation of the human papillomavirus type 16 late promoter by transcriptional elongation.
Immune evasion by muscle-specific gene expression in dystrophic muscle.
The E5 proteins.
Extensive cross-reactivity of adenovirus-specific cytotoxic T cells.
Hexon hypervariable region-modified adenovirus type 5 (Ad5) vectors display reduced hepatotoxicity but induce T lymphocyte phenotypes similar to Ad5 vectors.
Human Papillomavirus Downregulates the Expression of IFITM1 and RIPK3 to Escape from IFN $\gamma$ - and TNF $\alpha$ -Mediated Antiproliferative Effects and Necroptosis.
The infectivity of progeny adenovirus in the presence of neutralizing antibody.
Linearized oncolytic adenoviral plasmid DNA delivered by bioreducible polymers.
Application of gene therapy in diabetes care.
The effect of surface modification of adenovirus with an arginine-grafted bioreducible polymer on transduction efficiency and immunogenicity in cancer gene therapy.

Table A2-24, Cluster 23

Cluster 23 focuses on pathogenicity and virulence of Staphylococcus aureus, emphasizing the immune system evasion of this gram-positive bacterium (637)
Pathogenicity and virulence of Staphylococcus aureus.
Messing with the Sentinels-The Interaction of Staphylococcus aureus with Dendritic Cells.
Making the Most of the Host; Targeting the Autophagy Pathway Facilitates Staphylococcus aureus Intracellular Survival in Neutrophils.
Methicillin-Resistant Staphylococcus aureus from Peninsular Malaysian Animal Handlers: Molecular Profile, Antimicrobial Resistance, Immune Evasion Cluster and Genotypic Categorization.

Staphylococcus aureus host interactions and adaptation.
Colonization and infection of the skin by <i>S. aureus</i> : immune system evasion and the response to cationic antimicrobial peptides.
Molecular Assessment of Staphylococcus Aureus Strains in STAT3 Hyper-IgE Syndrome Patients.
Functional Insights of <i>MraZ</i> on the Pathogenicity of Staphylococcus aureus.
Neutrophils in innate host defense against Staphylococcus aureus infections.
Staphylococcus aureus Host Tropism and Its Implications for Murine Infection Models.
Absence of Staphylococcus aureus in Wild Populations of Fish Supports a Spillover Hypothesis.
Laboratory Mice Are Frequently Colonized with Staphylococcus aureus and Mount a Systemic Immune Response-Note of Caution for In vivo Infection Experiments.
High-resolution transcriptomic analysis of the adaptive response of Staphylococcus aureus during acute and chronic phases of osteomyelitis.
Methicillin-resistant Staphylococcus aureus: a pervasive pathogen highlights the need for new antimicrobial development.
Staphylococcus aureus strategies to evade the host acquired immune response.
The Role of Macrophages in Staphylococcus aureus Infection.
Staphylococcus aureus virulence factors in evasion from innate immune defenses in human and animal diseases.
Role of protein A in the evasion of host adaptive immune responses by Staphylococcus aureus.
Complement Inactivation Strategy of Staphylococcus aureus Using Decay-Accelerating Factor and the Response of Infected HaCaT Cells.
Macrophage-driven nutrient delivery to phagosomal Staphylococcus aureus supports bacterial growth.
Murine Models for Staphylococcal Infection.
Clinical Impact of Staphylococcus aureus Skin and Soft Tissue Infections.
Staphylococcus aureus adaptive evolution: Recent insights on how immune evasion, immunometabolic subversion and host genetics impact vaccine development.
Master mechanisms of Staphylococcus aureus: consider its excellent protective mechanisms hindering vaccine development!
Population variation in anti- <i>S. aureus</i> IgG isotypes influences surface protein A mediated immune subversion.
Development of a multicomponent Staphylococcus aureus vaccine designed to counter multiple bacterial virulence factors.
Covering all the Bases: Preclinical Development of an Effective Staphylococcus aureus Vaccine.
Characterisation of Staphylococcus aureus isolates from bovine mastitis in Ningxia, Western China.
Evasion of host defenses by intracellular Staphylococcus aureus.
Molecular mechanisms of Staphylococcus aureus nasopharyngeal colonization.
Staphylococcus aureus protein A promotes immune suppression.
Mouse models for infectious diseases caused by Staphylococcus aureus.
Antimicrobial Resistance and Genetic Lineages of Staphylococcus aureus from Wild Rodents: First Report of <i>mecC</i> -Positive Methicillin-Resistant <i>S. aureus</i> (MRSA) in Portugal.
Host-pathogen interactions between the skin and Staphylococcus aureus.
Inside job: Staphylococcus aureus host-pathogen interactions.
High prevalence of multidrug resistant <i>S. aureus</i> -CC398 and frequent detection of enterotoxin genes among non-CC398 <i>S. aureus</i> from pig-derived food in Spain.
Staphylococcal trafficking and infection-from 'nose to gut' and back.

Immune Evasion by Staphylococcus aureus.
Differential expression and roles of Staphylococcus aureus virulence determinants during colonization and disease.
How methicillin-resistant Staphylococcus aureus evade neutrophil killing.
Genetic and Phenotypic Characteristics of Staphylococcus aureus Isolates from Cystic Fibrosis Patients in Austria.
Staphylococcus aureus induces cell-surface expression of immune stimulatory NKG2D ligands on human monocytes.
Potential Relationship between Phenotypic and Molecular Characteristics in Revealing Livestock-Associated Staphylococcus aureus in Chinese Humans without Occupational Livestock Contact.
The therapeutic effect of chlorogenic acid against Staphylococcus aureus infection through sortase A inhibition.
Epic Immune Battles of History: Neutrophils vs. Staphylococcus aureus.
Association of virulence genes with mecA gene in Staphylococcus aureus isolates from Tertiary Hospitals in Nigeria.
Molecular characterization of Staphylococcus aureus isolated from humans related to a livestock farm in Spain, with detection of MRSA-CC130 carrying mecC gene: A zoonotic case?
Antimicrobial Mechanisms of Macrophages and the Immune Evasion Strategies of Staphylococcus aureus.
Staphylococcus aureus Impairs the Function of and Kills Human Dendritic Cells via the LukAB Toxin.
Wild rodents and shrews are natural hosts of Staphylococcus aureus.
Influence of transcription regulator SAUSA300_1968 on the virulence protein secretion and immune evasion by Staphylococcus aureus.
Staphylococcus aureus uses the ArlRS and MgrA cascade to regulate immune evasion during skin infection.
Staphylococcus aureus: setting its sights on the human innate immune system.
Mechanisms of Immune Evasion and Bone Tissue Colonization That Make Staphylococcus aureus the Primary Pathogen in Osteomyelitis.
Intracellular escape strategies of Staphylococcus aureus in persistent cutaneous infections.
Neutrophils versus Staphylococcus aureus: a biological tug of war.
Methicillin-Resistant Staphylococcus aureus CC398 in Purulent Lesions of Piglets and Fattening Pigs in Portugal.
Genetic diversity, antibiotic resistance, and virulence characteristics of Staphylococcus aureus from raw milk over 10 years in Shanghai.
Clonal complexes and virulence factors of Staphylococcus aureus from several cities in India.
Global spread of mouse-adapted Staphylococcus aureus lineages CC1, CC15, and CC88 among mouse breeding facilities.
Evasion of Immunological Memory by S. aureus Infection: Implications for Vaccine Design.
Whole-Genome Sequencing of Staphylococcus aureus and Staphylococcus haemolyticus Clinical Isolates from Egypt.
Molecular Characterization and Clonal Diversity of Methicillin-Resistant and -Susceptible Staphylococcus aureus Isolates of Milk of Cows with Clinical Mastitis in Tunisia.
Comparative Genomic Reveals Clonal Heterogeneity in Persistent Staphylococcus aureus Infection.
Detection, molecular characterization, and antibiogram of multi-drug resistant and methicillin-resistant Staphylococcus aureus (MRSA) isolated from pets and pet owners in Malaysia.
Community-associated methicillin-resistant Staphylococcus aureus immune evasion and virulence.
Staphylococcus schweitzeri-An Emerging One Health Pathogen?



Unstable chromosome rearrangements in <i>Staphylococcus aureus</i> cause phenotype switching associated with persistent infections.
Genetic lineages, antimicrobial resistance, and virulence in <i>Staphylococcus aureus</i> of meat samples in Spain: analysis of immune evasion cluster (IEC) genes.
Leukocidins and the Nuclease Nuc Prevent Neutrophil-Mediated Killing of <i>Staphylococcus aureus</i> Biofilms.
<i>Staphylococcus aureus</i> Complex in the Straw-Colored Fruit Bat ( <i>Eidolon helvum</i> ) in Nigeria.
Molecular characterisation of methicillin-sensitive <i>Staphylococcus aureus</i> from deep surgical site infections in orthopaedic patients.
Detection of MRSA ST3061-t843-mecC and ST398-t011-mecA in white stork nestlings exposed to human residues.
Recruitment of C4b-binding protein is not a complement evasion strategy employed by <i>Staphylococcus aureus</i> .
Examination of <i>Staphylococcus aureus</i> Prophages Circulating in Egypt.
Neutralizing <i>Staphylococcus aureus</i> Virulence with AZD6389, a Three mAb Combination, Accelerates Closure of a Diabetic Polymicrobial Wound.
A Small Membrane Stabilizing Protein Critical to the Pathogenicity of <i>Staphylococcus aureus</i> .
Diversity and pathogenesis of <i>Staphylococcus aureus</i> from bovine mastitis: current understanding and future perspectives.
Characterization of <i>Staphylococcus aureus</i> strains isolated from faeces of healthy neonates and potential mother-to-infant microbial transmission through breastfeeding.
<i>Staphylococcus aureus</i> chronic and relapsing infections: Evidence of a role for persister cells: An investigation of persister cells, their formation and their role in <i>S. aureus</i> disease.
Persistent <i>Staphylococcus aureus</i> isolates from two independent cases of bacteremia display increased bacterial fitness and novel immune evasion phenotypes.
Determining Impact of Growth Phases on Capacity of <i>Staphylococcus aureus</i> to Adhere to and Invade Host Cells.
Enterotoxin genes and antimicrobial resistance in <i>Staphylococcus aureus</i> isolated from food products in Algeria.
Molecular analysis of human and canine <i>Staphylococcus aureus</i> strains reveals distinct extended-host-spectrum genotypes independent of their methicillin resistance.
Genotypic and Phenotypic Markers of Livestock-Associated Methicillin-Resistant <i>Staphylococcus aureus</i> CC9 in Humans.
Characterisation of <i>Staphylococcus aureus</i> isolates from bloodstream infections, Democratic Republic of the Congo.
Population Analysis of <i>Staphylococcus aureus</i> Reveals a Cryptic, Highly Prevalent Superantigen SEIW That Contributes to the Pathogenesis of Bacteremia.
<i>Staphylococcus aureus</i> infection induces protein A-mediated immune evasion in humans.
Adaptation of <i>Staphylococcus aureus</i> in a Medium Mimicking a Diabetic Foot Environment.
Genomic Characterization of <i>Staphylococcus aureus</i> in Wildlife.
Targeting fundamental pathways to disrupt <i>Staphylococcus aureus</i> survival: clinical implications of recent discoveries.
Antivirulence Strategies for the Treatment of <i>Staphylococcus aureus</i> Infections: A Mini Review.
Adaptive Upregulation of Clumping Factor A (ClfA) by <i>Staphylococcus aureus</i> in the Obese, Type 2 Diabetic Host Mediates Increased Virulence.
Complement regulator C4BP binds to <i>Staphylococcus aureus</i> and decreases opsonization.

Genomic characterization of Staphylococcus aureus isolates causing osteoarticular infections in otherwise healthy children.
In or out: Phagosomal escape of Staphylococcus aureus.
Prevalence and characterization of methicillin-resistant Staphylococcus aureus carrying mecA or mecC and methicillin-susceptible Staphylococcus aureus in dairy sheep farms in central Italy.
Investigation of Virulence Genes and Resistance Genes in a Group of Staphylococcus aureus Isolated in Yuhang, China.
PINK1/parkin-mediated mitophagy alleviates Staphylococcus aureus-induced NLRP3 inflammasome and NF- $\kappa$ B pathway activation in bovine mammary epithelial cells.
Staphylococcus aureus infection dynamics.
Staphylococcus aureus stimulates neutrophil itaconate production that suppresses the oxidative burst.
Pigs as a potential source of emerging livestock-associated Staphylococcus aureus in Africa: a systematic review.
Antimicrobial Susceptibility Profiles and Molecular Characterisation of Staphylococcus aureus from Pigs and Workers at Farms and Abattoirs in Zambia.
Staphylococcal colonization and infection: homeostasis versus disbalance of human (innate) immunity and bacterial virulence.
Immunization of young heifers with staphylococcal immune evasion proteins before natural exposure to Staphylococcus aureus induces a humoral immune response in serum and milk.
Transcriptional adaptations during long-term persistence of Staphylococcus aureus in the airways of a cystic fibrosis patient.
Staphylococcus aureus survival in human blood.
Characterization of staphylococci in urban wastewater treatment plants in Spain, with detection of methicillin resistant Staphylococcus aureus ST398.
Oleate Hydratase (OhyA) Is a Virulence Determinant in Staphylococcus aureus.
Staphylococcus aureus Exploits the Host Apoptotic Pathway To Persist during Infection.
DNA microarray analysis of Staphylococcus aureus causing bloodstream infection: bacterial genes associated with mortality?
Adenosine synthase A contributes to recurrent Staphylococcus aureus infection by dampening protective immunity.
A bacterial outer membrane vesicle-based click vaccine elicits potent immune response against Staphylococcus aureus in mice.
Genetic variation in Staphylococcus aureus surface and immune evasion genes is lineage associated: implications for vaccine design and host-pathogen interactions.
Staphylococcus aureus and CCR5: unveiling commonalities in host-pathogen interactions and potential treatment strategies.
The persistence of intracellular Staphylococcus aureus in the sinuses: a longitudinal study.
Identification of novel targets for host-directed therapeutics against intracellular Staphylococcus aureus.
Staphylococcus aureus innate immune evasion is lineage-specific: a bioinformatics study.
Cathelicidins Mitigate Staphylococcus aureus Mastitis and Reduce Bacterial Invasion in Murine Mammary Epithelium.
[Molecular characterization of Staphylococcus aureus ST6 and ST7 isolates from food-borne illness outbreaks].
A global view of Staphylococcus aureus whole genome expression upon internalization in human epithelial cells.

Shedding of tumor necrosis factor receptor 1 induced by protein A decreases tumor necrosis factor alpha availability and inflammation during systemic Staphylococcus aureus infection.
Genomic Analysis of Bovine Staphylococcus aureus Isolates from Milk To Elucidate Diversity and Determine the Distributions of Antimicrobial and Virulence Genes and Their Association with Mastitis.
A natural human monoclonal antibody targeting Staphylococcus Protein A protects against Staphylococcus aureus bacteremia.
Into the storm: Chasing the opportunistic pathogen Staphylococcus aureus from skin colonisation to life-threatening infections.
Flavone inhibits Staphylococcus aureus virulence via inhibiting the sae two component system.
Staphylococcus aureus Carriage in the Nasotracheal Cavities of White Stork Nestlings (Ciconia ciconia) in Spain: Genetic Diversity, Resistomes and Virulence Factors.
Genomic characterization of Staphylococcus aureus from Canastra Minas Artisanal Cheeses.
Virulence Genes of S. aureus from Dairy Cow Mastitis and Contagiousness Risk.
CCR5 is a receptor for Staphylococcus aureus leukotoxin ED.
Phagocyte subsets and lymphocyte clonal deletion behind ineffective immune response to Staphylococcus aureus.
A privileged intraphagocyte niche is responsible for disseminated infection of Staphylococcus aureus in a zebrafish model.
Characterization of a mouse-adapted Staphylococcus aureus strain.
Pattern characterization of genes involved in non-specific immune response in Staphylococcus aureus isolates from intramammary infections.
Molecular analysis of immune evasion cluster (IEC) genes and intercellular adhesion gene cluster (ICA) among methicillin-resistant and methicillin-sensitive isolates of Staphylococcus aureus.
PSM Peptides From Community-Associated Methicillin-Resistant Staphylococcus aureus Impair the Adaptive Immune Response via Modulation of Dendritic Cell Subsets in vivo.
The TLR2 Antagonist Staphylococcal Superantigen-Like Protein 3 Acts as a Virulence Factor to Promote Bacterial Pathogenicity in vivo.
Antibody-Based Biologics and Their Promise to Combat Staphylococcus aureus Infections.
Host- and microbe determinants that may influence the success of S. aureus colonization.
Immune evasion cluster-positive bacteriophages are highly prevalent among human Staphylococcus aureus strains, but they are not essential in the first stages of nasal colonization.
Detection of methicillin-resistant Staphylococcus aureus (MRSA) carrying the mecC gene in wild small mammals in Spain.
Genomic analysis of Staphylococcus aureus sequential isolates from lungs of patients with cystic fibrosis.
Evaluation of virulence potential of methicillin-sensitive and methicillin-resistant Staphylococcus aureus isolates from a German refugee cohort.
Mechanisms of bacterial virulence in pulmonary infections.
Nasal staphylococci community of healthy pigs and pig-farmers in Aragon (Spain). Predominance and within-host resistome diversity in MRSA-CC398 and MSSA-CC9 lineages.
Use of Flow Cytometry to Evaluate Phagocytosis of Staphylococcus aureus by Human Neutrophils.
Identification of a staphylococcal complement inhibitor with broad host specificity in equid Staphylococcus aureus strains.
Staphylococcus aureus-A Known Opponent against Host Defense Mechanisms and Vaccine Development-Do We Still Have a Chance to Win?
Exploring differentially expressed genes of Staphylococcus aureus exposed to human tonsillar cells using RNA sequencing.

Staphylococcus aureus vaccines: Deviating from the carol.
O-Acetylation of Peptidoglycan Limits Helper T Cell Priming and Permits Staphylococcus aureus Reinfection.
Prolactin regulates H3K9ac and H3K9me2 epigenetic marks and miRNAs expression in bovine mammary epithelial cells challenged with Staphylococcus aureus.
Immunization Against Staphylococcus aureus Infections.
Neutrophil-Mediated Phagocytosis of Staphylococcus aureus.
Occurrence and molecular composition of methicillin-resistant Staphylococcus aureus isolated from ocular surfaces of horses presented with ophthalmologic disease.
Staphylococcal manipulation of host immune responses.
Staphylococcus aureus Induces Shedding of IL-1RII in Monocytes and Neutrophils.
Methicillin Resistant Staphylococcus aureus and public fomites: a review.
Prevalence and characterization of Staphylococcus aureus and Staphylococcus argenteus isolated from rice and flour products in Guangdong, China.
Methicillin-resistant Staphylococcus aureus adaptation to human keratinocytes.
The Nature of Antibacterial Adaptive Immune Responses against Staphylococcus aureus Is Dependent on the Growth Phase and Extracellular Peptidoglycan.
Looking toward basic science for potential drug discovery targets against community-associated MRSA.
The Prevalence of Staphylococcus aureus and the Occurrence of MRSA CC398 in Monkey Feces in a Zoo Park in Eastern China.
Advanced glycation end products reduce macrophage-mediated killing of Staphylococcus aureus by ARL8 upregulation and inhibition of autolysosome formation.
Prevalence and Genomic Structure of Bacteriophage phi3 in Human-Derived Livestock-Associated Methicillin-Resistant Staphylococcus aureus Isolates from 2000 to 2015.
Epigenetic response in mice mastitis: Role of histone H3 acetylation and microRNA(s) in the regulation of host inflammatory gene expression during Staphylococcus aureus infection.
Characterization of Livestock-Associated Methicillin-Resistant Staphylococcus aureus Isolates Obtained From Commercial Rabbitries Located in the Iberian Peninsula.
Prevalence, antibiotic resistance, virulence traits and genetic lineages of Staphylococcus aureus in healthy sheep in Tunisia.
Host chemokines bind to Staphylococcus aureus and stimulate protein A release.
Staphylococcus aureus temperate bacteriophage: carriage and horizontal gene transfer is lineage associated.
Dalbavancin Boosts the Ability of Neutrophils to Fight Methicillin-Resistant Staphylococcus aureus.
Studying Staphylococcal Leukocidins: A Challenging Endeavor.
Attenuation of Staphylococcus aureus-Induced Bacteremia by Human Mini-Antibodies Targeting the Complement Inhibitory Protein Efb.
Immune control of Staphylococcus aureus - regulation and counter-regulation of the adaptive immune response.
Staphylococcus lugdunensis: a Skin Commensal with Invasive Pathogenic Potential.
Adaptive processes of Staphylococcus aureus isolates during the progression from acute to chronic bone and joint infections in patients.
Novel method for detecting complement C3 deposition on Staphylococcus aureus.
Genomic analyses of Staphylococcus aureus isolated from yaks in Ganzi Tibetan Autonomous Prefecture, China.

Horses in Denmark Are a Reservoir of Diverse Clones of Methicillin-Resistant and -Susceptible <i>Staphylococcus aureus</i> .
SpA, ClfA, and FnbA genetic variations lead to Staphaurex test-negative phenotypes in bovine mastitis <i>Staphylococcus aureus</i> isolates.
<i>Staphylococcus aureus</i> Phenol-Soluble Modulins $\alpha 1$ - $\alpha 3$ Act as Novel Toll-Like Receptor (TLR) 4 Antagonists to Inhibit HMGB1/TLR4/NF- $\kappa$ B Signaling Pathway.
A new anti-infective strategy to reduce the spreading of antibiotic resistance by the action on adhesion-mediated virulence factors in <i>Staphylococcus aureus</i> .
The Role of hlb-Converting Bacteriophages in <i>Staphylococcus aureus</i> Host Adaption.
Virulence factor landscape of a <i>Staphylococcus aureus</i> sequence type 45 strain, MCRF184.
Emergence of livestock-associated MRSA isolated from cystic fibrosis patients: Result of a Belgian national survey.
The MpsB protein contributes to both the toxicity and immune evasion capacity of <i>Staphylococcus aureus</i> .
Investigation of <i>Staphylococcus aureus</i> adhesion and invasion of host cells.
Metabolic Adaptations During <i>Staphylococcus aureus</i> and <i>Candida albicans</i> Co-Infection.
First Finding of a Methicillin-Resistant <i>Staphylococcus aureus</i> (MRSA) t304/ST6 from Bovine Clinical Mastitis.
First report of sasX-positive methicillin-resistant <i>Staphylococcus aureus</i> in Japan.
<i>Staphylococcus aureus</i> Infects Osteoclasts and Replicates Intracellularly.
<i>Staphylococcus aureus</i> protects its immune-evasion proteins against degradation by neutrophil serine proteases.
[Aspects of the innate immune response to intramammary <i>Staphylococcus aureus</i> infections in cattle].
In vitro phagocytosis of methicillin resistant and methicillin sensitive <i>Staphylococcus aureus</i> by human polymorphonuclear leucocytes.
Antibiotic resistance and host immune evasion in <i>Staphylococcus aureus</i> mediated by a metabolic adaptation.
Virulence potential of <i>Staphylococcus aureus</i> isolates from Buruli ulcer patients.
High occurrence of enterotoxigenic isolates and low antibiotic resistance rates of <i>Staphylococcus aureus</i> isolated from raw milk from cows and ewes.
A functional menadione biosynthesis pathway is required for capsule production by <i>Staphylococcus aureus</i> .
Molecular Epidemiology of Methicillin-Susceptible and Methicillin-Resistant <i>Staphylococcus aureus</i> in Wild, Captive and Laboratory Rats: Effect of Habitat on the Nasal <i>S. aureus</i> Population.
First report on MRSA CC398 recovered from wild boars in the north of Portugal. Are we facing a problem?
Phenotypic and molecular characteristics of <i>Staphylococcus aureus</i> and methicillin-resistant <i>Staphylococcus aureus</i> in slaughterhouse pig-related workers and control workers in Guangdong Province, China.
Optimization and evaluation of a multiplex PCR assay for detection of <i>Staphylococcus aureus</i> and its major virulence genes for assessing food safety.
Adhesion, invasion and evasion: the many functions of the surface proteins of <i>Staphylococcus aureus</i> .
Bovine mastitis outbreak in Japan caused by methicillin-resistant <i>Staphylococcus aureus</i> New York/Japan clone.
Prophage-encoded immune evasion factors are critical for <i>Staphylococcus aureus</i> host infection, switching, and adaptation.

Staphylococcal superantigen-like protein 10 induces necroptosis through TNFR1 activation of RIPK3-dependent signal pathways.
Staphylococcus aureus sortase A contributes to the Trojan horse mechanism of immune defense evasion with its intrinsic resistance to Cys184 oxidation.
Staphylococcus aureus Manipulates Innate Immunity through Own and Host-Expressed Proteases.
Regulatory adaptation of Staphylococcus aureus during nasal colonization of humans.
Comparative secretome analysis of Staphylococcus aureus strains with different within-herd intramammary infection prevalence.
Livestock origin for a human pandemic clone of community-associated methicillin-resistant Staphylococcus aureus.
Targeting leukocidin-mediated immune evasion protects mice from Staphylococcus aureus bacteremia.
Inhibition of complement activation by a secreted Staphylococcus aureus protein.
Temperate Phages of Staphylococcus aureus.
Characterization of highly virulent community-associated methicillin-resistant Staphylococcus aureus ST9-SCCmec XII causing bloodstream infection in China.
High prevalence of methicillin-resistant Staphylococcus aureus (MRSA) carrying the mecC gene in a semi-extensive red deer (Cervus elaphus hispanicus) farm in Southern Spain.
Disruption of the alternative pathway convertase occurs at the staphylococcal surface via the acquisition of factor H by Staphylococcus aureus.
Livestock-associated methicillin and multidrug resistant S. aureus in humans is associated with occupational pig contact, not pet contact.
Staphylococcus aureus lipoproteins promote abscess formation in mice, shielding bacteria from immune killing.
Staphylococcus aureus small colony variants impair host immunity by activating host cell glycolysis and inducing necroptosis.
Colonization and infection of the human host by staphylococci: adhesion, survival and immune evasion.
Methicillin-resistant and methicillin-susceptible Staphylococcus aureus in dairy sheep and in-contact humans: An intra-farm study.
Capsular polysaccharides are an important immune evasion mechanism for Staphylococcus aureus.
Recurrent infections and immune evasion strategies of Staphylococcus aureus.
Cytoplasmic replication of Staphylococcus aureus upon phagosomal escape triggered by phenol-soluble modulin $\alpha$ .
Equine Methicillin-Resistant Sequence Type 398 Staphylococcus aureus (MRSA) Harbor Mobile Genetic Elements Promoting Host Adaptation.
Staphylococcal Phages Adapt to New Hosts by Extensive Attachment Site Variability.
Simultaneous Nasal Carriage by Methicillin-Resistant and Methicillin Susceptible Staphylococcus aureus of Lineage ST398 in a Live Pig Transporter.
Staphylococci evade the innate immune response by disarming neutrophils and forming biofilms.
Genetic lineages and antimicrobial resistance genotypes in Staphylococcus aureus from children with atopic dermatitis: detection of clonal complexes CC1, CC97 and CC398.
Staphylococcal proteases aid in evasion of the human complement system.
S. aureus Evades Macrophage Killing through NLRP3-Dependent Effects on Mitochondrial Trafficking.
Presence of Immune Evasion Cluster and Molecular Typing of Methicillin-Susceptible Staphylococcus aureus Isolated from Food Handlers.

Identification of secreted and cellular antigens of Staphylococcus aureus causing dairy sheep mastitis and their potential for vaccine development.
Staphylococcus aureus and Neutrophil Extracellular Traps: The Master Manipulator Meets Its Match in Immunothrombosis.
Adaptation of Staphylococcus aureus to the Human Skin Environment Identified Using an ex vivo Tissue Model.
Molecular characterization and clonal diversity of methicillin-susceptible Staphylococcus aureus in milk of cows with mastitis in Brazil.
Within-host evolution of bovine Staphylococcus aureus selects for a SigB-deficient pathotype characterized by reduced virulence but enhanced proteolytic activity and biofilm formation.
Targeting the R domain of coagulase by active vaccination protects mice against lethal Staphylococcus aureus infection.
Secretory proteins in the orchestration of microbial virulence: The curious case of Staphylococcus aureus.
Bacterial immune evasion via an IL-10 mediated host response, a novel pathophysiologic mechanism for chronic rhinosinusitis.
Detection of the biofilm component polysaccharide intercellular adhesin in Staphylococcus aureus infected cow udders.
Staphylococcus aureus leukotoxin GH promotes formation of neutrophil extracellular traps.
Animal and human Staphylococcus aureus associated clonal lineages and high rate of Staphylococcus pseudintermedius novel lineages in Spanish kennel dogs: predominance of S. aureus ST398.
MntC-Dependent Manganese Transport Is Essential for Staphylococcus aureus Oxidative Stress Resistance and Virulence.
Stress Resistance and Pathogenicity of Nonthermal-Plasma-Induced Viable-but-Nonculturable Staphylococcus aureus through Energy Suppression, Oxidative Stress Defense, and Immune-Escape Mechanisms.
Novel Requirement for Staphylococcal Cell Wall-Anchored Protein SasD in Pulmonary Infection.
Immune evasion by staphylococci.
Community-Associated Staphylococcus aureus from Sub-Saharan Africa and Germany: A Cross-Sectional Geographic Correlation Study.
Staphylococcal Protein A Promotes Colonization and Immune Evasion of the Epidemic Healthcare-Associated MRSA ST239.
Staphylococcus aureus: the multi headed hydra resists and controls human complement response in multiple ways.
Prevalence of the immune evasion gene cluster in Staphylococcus aureus CC398.
Production of an attenuated phenol-soluble modulins variant unique to the MRSA clonal complex 30 increases severity of bloodstream infection.
Staphylococcus aureus stimulates neutrophil targeting chemokine expression in keratinocytes through an autocrine IL-1 $\alpha$ signaling loop.
Toll-like receptor 2-dependent endosomal signaling by Staphylococcus aureus in monocytes induces type I interferon and promotes intracellular survival.
Short repeats in the spa gene of Staphylococcus aureus are prone to nonsense mutations: stop codons can be found in strains isolated from patients with generalized infection.
TCA cycle inactivation in Staphylococcus aureus alters nitric oxide production in RAW 264.7 cells.
Adhesive Virulence Factors of Staphylococcus aureus Resist Digestion by Coagulation Proteases Thrombin and Plasmin.

Transcriptional regulation of virulence factors Spa and ClfB by the SpoVG-Rot cascade in <i>Staphylococcus aureus</i> .
Genomic Analysis of <i>Staphylococcus aureus</i> of the Lineage CC130, Including mecC-Carrying MRSA and MSSA Isolates Recovered of Animal, Human, and Environmental Origins.
Characterization of a vraG Mutant in a Genetically Stable <i>Staphylococcus aureus</i> Small-Colony Variant and Preliminary Assessment for Use as a Live-Attenuated Vaccine against Intramammary Infections.
Adaptive immune response to lipoproteins of <i>Staphylococcus aureus</i> in healthy subjects.
<i>Staphylococcus aureus</i> virulence is enhanced by secreted factors that block innate immune defenses.
Extensive horizontal gene transfer during <i>Staphylococcus aureus</i> co-colonization in vivo.
<i>Staphylococcus aureus</i> persistence in non-professional phagocytes.
<i>Staphylococcus aureus</i> determinants for nasal colonization.
The SaeR/S gene regulatory system is essential for innate immune evasion by <i>Staphylococcus aureus</i> .
<i>Staphylococcus aureus</i> elaborates leukocidin AB to mediate escape from within human neutrophils.
A coagulase-negative and non-haemolytic strain of <i>Staphylococcus aureus</i> for investigating the roles of SrtA in a murine model of bloodstream infection.
Regulation of <i>Staphylococcus aureus</i> MntC expression and its role in response to oxidative stress.
A <i>Staphylococcus aureus</i> TIR domain protein virulence factor blocks TLR2-mediated NF- $\kappa$ B signaling.
<i>Staphylococcus aureus</i> -induced clotting of plasma is an immune evasion mechanism for persistence within the fibrin network.
<i>Staphylococcus aureus</i> Depends on Eap Proteins for Preventing Degradation of Its Phenol-Soluble Modulin Toxins by Neutrophil Serine Proteases.
Serological versus molecular typing of surface-associated immune evading polysaccharide antigens-based phenotypes of <i>Staphylococcus aureus</i> .
Comparative Genomic Analysis of the Human Variant of Methicillin-Resistant <i>Staphylococcus aureus</i> CC398 in Japan and Korea.
Methicillin-resistant <i>Staphylococcus aureus</i> alters cell wall glycosylation to evade immunity.
Nanoscaled Discovery of a Shunt Rifamycin from <i>Salinispora arenicola</i> Using a Three-Color GFP-Tagged <i>Staphylococcus aureus</i> Macrophage Infection Assay.
Engineered human antibodies for the opsonization and killing of <i>Staphylococcus aureus</i> .
Effect of filaggrin breakdown products on growth of and protein expression by <i>Staphylococcus aureus</i> .
<i>Staphylococcus aureus</i> secretes a unique class of neutrophil serine protease inhibitors.
Linoleic acid metabolism activation in macrophages promotes the clearing of intracellular <i>Staphylococcus aureus</i> .
<i>Staphylococcus aureus</i> proteins Sbi and Efb recruit human plasmin to degrade complement C3 and C3b.
<i>Staphylococcus aureus</i> intra-mammary infection affects the expression pattern of IL-R8 in goat.
Livestock-Associated Methicillin-Resistant <i>Staphylococcus aureus</i> (MRSA) in Purulent Subcutaneous Lesions of Farm Rabbits.
Identification of a highly transmissible animal-independent <i>Staphylococcus aureus</i> ST398 clone with distinct genomic and cell adhesion properties.
<i>Staphylococcal</i> protein Ecb impairs complement receptor-1 mediated recognition of opsonized bacteria.
Expression and function of protein A in <i>Staphylococcus pseudintermedius</i> .
Genome Sequence of a Highly Virulent pvl-positive Vancomycin-intermediate-resistant <i>Staphylococcus aureus</i> Sequence Type 30.



Different distribution of antimicrobial resistance genes and virulence profiles of <i>Staphylococcus aureus</i> strains isolated from clinical mastitis in six countries.
Comparative genomics of <i>Staphylococcus aureus</i> associated with subclinical and clinical bovine mastitis.
Return of the Trojan horse: intracellular phenotype switching and immune evasion by <i>Staphylococcus aureus</i> .
Molecular Epidemiology of Penicillin-Susceptible <i>Staphylococcus aureus</i> Bacteremia in Australia and Reliability of Diagnostic Phenotypic Susceptibility Methods to Detect Penicillin Susceptibility.
DNA microarray genotyping and virulence and antimicrobial resistance gene profiling of methicillin-resistant <i>Staphylococcus aureus</i> bloodstream isolates from renal patients.
Livestock-Associated Methicillin Resistant and Methicillin Susceptible <i>Staphylococcus aureus</i> Sequence Type (CC)1 in European Farmed Animals: High Genetic Relatedness of Isolates from Italian Cattle Herds and Humans.
The Two-Component System ArlRS and Alterations in Metabolism Enable <i>Staphylococcus aureus</i> to Resist Calprotectin-Induced Manganese Starvation.
Molecular Evolution and Adaptation of Livestock-Associated Methicillin-Resistant <i>Staphylococcus aureus</i> (LA-MRSA) Sequence Type 9.
Gene expression profile of human peripheral blood mononuclear cells induced by <i>Staphylococcus aureus</i> lipoteichoic acid.
<i>Staphylococcus aureus</i> Quorum Regulator SarA Targeted Compound, 2-[(Methylamino)methyl]phenol Inhibits Biofilm and Down-Regulates Virulence Genes.
Bioinformatics study of expression from genomes of epidemiologically related MRSA CC398 isolates from human and wild animal samples.
Interactions between the foreign body reaction and <i>Staphylococcus aureus</i> biomaterial-associated infection. Winning strategies in the derby on biomaterial implant surfaces.
Comparative Prevalence of Immune Evasion Complex Genes Associated with $\beta$ -Hemolysin Converting Bacteriophages in MRSA ST5 Isolates from Swine, Swine Facilities, Humans with Swine Contact, and Humans with No Swine Contact.
Mechanical Strength and Inhibition of the <i>Staphylococcus aureus</i> Collagen-Binding Protein Cna.
<i>Staphylococcal</i> enterotoxin-like X (SEIX) is a unique superantigen with functional features of two major families of <i>staphylococcal</i> virulence factors.
<i>Staphylococcal</i> innate immune evasion.
Key Concepts of Musculoskeletal Infection.
<i>S. aureus</i> IgG-binding proteins SpA and Sbi: host specificity and mechanisms of immune complex formation.
Methionine sulfoxide reductases protect against oxidative stress in <i>Staphylococcus aureus</i> encountering exogenous oxidants and human neutrophils.
Characterization of methicillin-resistant <i>Staphylococcus sciuri</i> isolates from industrially raised pigs, cattle and broiler chickens.
Fibronectin-binding protein B variation in <i>Staphylococcus aureus</i> .
Structure and biological activities of beta toxin from <i>Staphylococcus aureus</i> .
Genomic epidemiology of methicillin-susceptible <i>Staphylococcus aureus</i> across colonisation and skin and soft tissue infection.
Structural insights into recognition of chemokine receptors by <i>Staphylococcus aureus</i> leukotoxins.
The Energy-Coupling Factor Transporter Module EcfAA <sup>T</sup> , a Novel Candidate for the Genetic Basis of Fatty Acid-Auxotrophic Small-Colony Variants of <i>Staphylococcus aureus</i> .

Mucosa-Associated Invariant T Cell Hypersensitivity to Staphylococcus aureus Leukocidin ED and Its Modulation by Activation.
Using Quantitative Spectrometry to Understand the Influence of Genetics and Nutritional Perturbations On the Virulence Potential of Staphylococcus aureus.
Dual RNA regulatory control of a Staphylococcus aureus virulence factor.
Staphylococcal protein a deletes B-1a and marginal zone B lymphocytes expressing human immunoglobulins: an immune evasion mechanism.
Inhibitory effects of lipoteichoic acid from Staphylococcus aureus on platelet function and platelet-monocyte aggregation.
Enzymatic properties of Staphylococcus aureus adenosine synthase (AdsA).
Molecular insights into mechanisms of GPCR hijacking by Staphylococcus aureus.
Inulin-lipid hybrid (ILH) microparticles promote pH-triggered release of rifampicin within infected macrophages.
Staphylococcal Osteomyelitis: Disease Progression, Treatment Challenges, and Future Directions.
Backbone resonance assignments of innate immune evasion protein EapH2 from the S. aureus.
Protein A is released into the Staphylococcus aureus culture supernatant with an unprocessed sorting signal.
Staphylococcal Complement Evasion Protein Sbi Stabilises C3d Dimers by Inducing an N-Terminal Helix Swap.
Staphylococcus aureus PSM peptides induce tolerogenic dendritic cells upon treatment with ligands of extracellular and intracellular TLRs.
Bacteria-Responsive Biomimetic Selenium Nanosystem for Multidrug-Resistant Bacterial Infection Detection and Inhibition.
Identification, Cloning, and Characterization of Staphylococcus pseudintermedius Coagulase.
Identification of peptidic inhibitors of the alternative complement pathway based on Staphylococcus aureus SCIN proteins.
Staphylococcal protein A-formulated immune complexes suppress enterotoxin-induced cellular responses in nasal polyps.
Characterisation of Staphylococcus felis isolated from cats using whole genome sequencing.
Zinc-doped ferric oxyhydroxide nano-layer enhances the bactericidal activity and osseointegration of a magnesium alloy through augmenting the formation of neutrophil extracellular traps.
Staphylococcal Ecb protein and host complement regulator factor H enhance functions of each other in bacterial immune evasion.
Complete non-proline backbone resonance assignments of the S. aureus neutrophil serine protease inhibitor, EapH1.
Fibrinogen Activates the Capture of Human Plasminogen by Staphylococcal Fibronectin-Binding Proteins.
Changes in the spike and nucleocapsid protein of porcine epidemic diarrhea virus strain in Vietnam-a molecular potential for the vaccine development?
3D Cocultures of Osteoblasts and Staphylococcus aureus on Biomimetic Bone Scaffolds as a Tool to Investigate the Host-Pathogen Interface in Osteomyelitis.
Interaction of staphylococci with bone.
Multiscale conformational heterogeneity in staphylococcal protein a: possible determinant of functional plasticity.
Complete genome of a methicillin-resistant Staphylococcus vitulinus from Danish ground beef meat carrying a mecA2 resistance gene and a novel ccr allotype.

WGS analysis of ST9-MRSA-XII isolates from live pigs in China provides insights into transmission among porcine, human and bovine hosts.
The infectious aspects of atopic dermatitis.
Pathogen Receptor Membrane-Coating Facet Structures Boost Nanomaterial Immune Escape and Antibacterial Performance.
Proteomic analysis of adult <i>Schistosoma mekongi</i> somatic and excretory-secretory proteins.
PSM-Mec-A Virulence Determinant that Connects Transcriptional Regulation, Virulence, and Antibiotic Resistance in <i>Staphylococci</i> .
Suppression of conformational heterogeneity at a protein-protein interface.
Context matters: The importance of dimerization-induced conformation of the LukGH leukocidin of <i>Staphylococcus aureus</i> for the generation of neutralizing antibodies.
TLR13 recognizes bacterial 23S rRNA devoid of erythromycin resistance-forming modification.
YycH and YycI Regulate Expression of <i>Staphylococcus aureus</i> Autolysins by Activation of WalRK Phosphorylation.
Structural basis for inhibition of complement C5 by the SSL7 protein from <i>Staphylococcus aureus</i> .
Characterization and virulence of <i>Streptococcus agalactiae</i> deficient in SaeRS of the two-component system.
Staphylococcal superantigen-like 5 activates platelets and supports platelet adhesion under flow conditions, which involves glycoprotein Ib $\alpha$ and $\alpha$ IIb $\beta$ 3.
Transcriptome analysis unveils survival strategies of <i>Streptococcus parauberis</i> against fish serum.
<i>Staphylococcus aureus</i> ST59: Concurrent but Separate Evolution of North American and East Asian Lineages.
Structure-function analysis of the C3 binding region of <i>Staphylococcus aureus</i> immune subversion protein Sbi.
Diversity in the C3b [corrected] contact residues and tertiary structures of the staphylococcal complement inhibitor (SCIN) protein family.
<i>Streptococcus uberis</i> strains isolated from the bovine mammary gland evade immune recognition by mammary epithelial cells, but not of macrophages.
Recombinant lipidated FLIPr effectively enhances mucosal and systemic immune responses for various vaccine types.
In silico assessment of virulence factors in strains of <i>Streptococcus oralis</i> and <i>Streptococcus mitis</i> isolated from patients with Infective Endocarditis.
Immune evasion by a staphylococcal complement inhibitor that acts on C3 convertases.
A competitive mechanism for staphylococcal toxin SSL7 inhibiting the leukocyte IgA receptor, Fc $\alpha$ RI, is revealed by SSL7 binding at the C $\alpha$ 2/C $\alpha$ 3 interface of IgA.
A structurally dynamic N-terminal region drives function of the staphylococcal peroxidase inhibitor (SPIN).
Genomic characterisation of multidrug-resistant <i>Bacillus toyonensis</i> strain 4HC1 isolated from marine plastic in Norway.
[Modulation of Host Immune System by Staphylococcal Superantigen-like (SSL) Proteins].
<i>Streptococcus pneumoniae</i> invades erythrocytes and utilizes them to evade human innate immunity.
Molecular epidemiology of <i>Streptococcus agalactiae</i> isolated from mastitis in Brazilian dairy herds.
Secondary Bacterial Infections Associated with Influenza Pandemics.
Crystal structure of the capsular polysaccharide-synthesis enzyme CapG from <i>Staphylococcus aureus</i> .
The Role of Streptococcal and Staphylococcal Exotoxins and Proteases in Human Necrotizing Soft Tissue Infections.
Resistome and virulome study on pathogenic <i>Streptococcus agalactiae</i> Guangzhou-SAG036.

Staphylococcal Superantigen-Like Protein 1 and 5 (SSL1 & SSL5) Limit Neutrophil Chemotaxis and Migration through MMP-Inhibition.
The MSCRAMM Family of Cell-Wall-Anchored Surface Proteins of Gram-Positive Cocci.
Draft genome sequences of extensively drug resistant and pandrug resistant <i>Acinetobacter baumannii</i> strains isolated from hospital wastewater in South Africa.
Comprehensive Virulence Gene Profiling of Bovine Non-aureus Staphylococci Based on Whole-Genome Sequencing Data.
<i>Staphylococcus aureus</i> SdrE captures complement factor H's C-terminus via a novel 'close, dock, lock and latch' mechanism for complement evasion.
Novel IgG-Degrading Enzymes of the IgdE Protease Family Link Substrate Specificity to Host Tropism of <i>Streptococcus</i> Species.
Local structural plasticity of the <i>Staphylococcus aureus</i> evasion protein EapH1 enables engagement with multiple neutrophil serine proteases.
Convertase inhibitory properties of Staphylococcal extracellular complement-binding protein.
Molecular analysis of the interaction between staphylococcal virulence factor Sbi-IV and complement C3d.
Differential Interaction of the Staphylococcal Toxins Pantone-Valentine Leukocidin and $\gamma$ -Hemolysin CB with Human C5a Receptors.
Characterisation of successive <i>Acinetobacter baumannii</i> isolates from a deceased haemophagocytic lymphohistiocytosis patient.
Simultaneous inhibition of two neutrophil serine proteases by the <i>S. aureus</i> innate immune evasion protein EapH2.
Genomic insights into bla(NDM)-carrying carbapenem-resistant <i>Klebsiella pneumoniae</i> clinical isolates from a university hospital in Thailand.
Comparison of genotypes, antimicrobial resistance and virulence profiles of oral and non oral <i>Enterococcus faecalis</i> from Brazil, Japan and the United Kingdom.
Mathematical Model for MRSA Nasal Carriage.
Evidence for multiple modes of neutrophil serine protease recognition by the EAP family of Staphylococcal innate immune evasion proteins.
Staphylococcal superantigen-like 12 induces the production of interleukin 4 in murine basophils.
Identification of a novel host-specific IgG protease in <i>Streptococcus phocae</i> subsp. <i>phocae</i> .
Mutational analyses reveal that the staphylococcal immune evasion molecule Sbi and complement receptor 2 (CR2) share overlapping contact residues on C3d: implications for the controversy regarding the CR2/C3d cocrystal structure.
A Facile machine learning multi-classification model for <i>Streptococcus agalactiae</i> clonal complexes.
Phylogenomic insights into evolutionary trajectories of multidrug resistant <i>S. pneumoniae</i> CC271 over a period of 14 years in China.
Structural basis for inhibition of TLR2 by staphylococcal superantigen-like protein 3 (SSL3).
<i>Salmonella enterica</i> serovar Typhi plasmid impairs dendritic cell responses to infection.
Biochemical characterization of evasion from peptidoglycan recognition by <i>Staphylococcus aureus</i> D-alanylated wall teichoic acid in insect innate immunity.
Induction of mastitis by cow-to-mouse fecal and milk microbiota transplantation causes microbiome dysbiosis and genomic functional perturbation in mice.
Cell wall-anchored 5'-nucleotidases in Gram-positive cocci.
Current concepts in the pathogenesis and treatment of chronic suppurative otitis media.
Staphylococcal superantigen-like protein 10 binds to phosphatidylserine and apoptotic cells.
Microbiology of fracture related infections.

Clonal diversity and turnover of *Streptococcus mitis* bv. 1 on shedding and nonshedding oral surfaces of human infants during the first year of life.

Table A2-25, Cluster 24

Cluster 24 focuses on Kaposi's sarcoma-associated herpesvirus, emphasizing its immune evasion strategies (233)
Adaptive immune responses to Kaposi's sarcoma-associated herpesvirus.
Looking at Kaposi's Sarcoma-Associated Herpesvirus-Host Interactions from a microRNA Viewpoint.
Modulation of Immune System by Kaposi's Sarcoma-Associated Herpesvirus: Lessons from Viral Evasion Strategies.
KSHV: pathways to tumorigenesis and persistent infection.
Mechanisms of Kaposi's Sarcoma-Associated Herpesvirus Latency and Reactivation.
Immune evasion in Kaposi's sarcoma-associated herpes virus associated oncogenesis.
Immune evasion by Kaposi's sarcoma-associated herpesvirus.
Immune evasion by Kaposi's sarcoma-associated herpesvirus.
Interplay between Kaposi's sarcoma-associated herpesvirus and the innate immune system.
Towards Better Understanding of KSHV Life Cycle: from Transcription and Posttranscriptional Regulations to Pathogenesis.
Immune evasion strategies of Kaposi's sarcoma-associated herpesvirus.
null
Signal Transduction Pathways Associated with KSHV-Related Tumors.
Regulation of KSHV Latency and Lytic Reactivation.
Kaposi's sarcoma-associated herpesvirus ORF54/dUTPase downregulates a ligand for the NK activating receptor NKp44.
Kaposi's Sarcoma-Associated Herpesvirus, the Etiological Agent of All Epidemiological Forms of Kaposi's Sarcoma.
Kaposi's sarcoma-associated herpesvirus immunoevasion and tumorigenesis: two sides of the same coin?
Kaposi's sarcoma-associated herpesvirus and extracellular vesicles.
Major Histocompatibility Complex Class II HLA-DR $\alpha$ Is Downregulated by Kaposi's Sarcoma-Associated Herpesvirus-Encoded Lytic Transactivator RTA and MARCH8.
Multi-step regulation of innate immune signaling by Kaposi's sarcoma-associated herpesvirus.
Cellular and viral oncogenes: the key to unlocking unknowns of Kaposi's sarcoma-associated herpesvirus pathogenesis.
Transcriptional analysis of latent and inducible Kaposi's sarcoma-associated herpesvirus transcripts in the K4 to K7 region.
The viral interferon regulatory factors of KSHV: immunosuppressors or oncogenes?
Immunological and inflammatory features of Kaposi's sarcoma and other Kaposi's sarcoma-associated herpesvirus/human herpesvirus 8-associated neoplasias.
Surface downregulation of major histocompatibility complex class I, PE-CAM, and ICAM-1 following de novo infection of endothelial cells with Kaposi's sarcoma-associated herpesvirus.
Characteristics of circulating KSHV-infected viroblasts during active KSHV+ multicentric Castleman disease.
Noncoding RNAs produced by oncogenic human herpesviruses.
Toll-like receptor 4 mediates innate immunity to Kaposi sarcoma herpesvirus.

Exploitation of the complement system by oncogenic Kaposi's sarcoma-associated herpesvirus for cell survival and persistent infection.
Primary B Lymphocytes Infected with Kaposi's Sarcoma-Associated Herpesvirus Can Be Expanded In Vitro and Are Recognized by LANA-Specific CD4+ T Cells.
Nuclear Innate Immune DNA Sensor IFI16 Is Degraded during Lytic Reactivation of Kaposi's Sarcoma-Associated Herpesvirus (KSHV): Role of IFI16 in Maintenance of KSHV Latency.
Human herpesvirus 8-encoded cytokines.
Kaposi sarcoma-associated herpesvirus latency-associated nuclear antigen: more than a key mediator of viral persistence.
Concurrent Control of the Kaposi's Sarcoma-Associated Herpesvirus Life Cycle through Chromatin Modulation and Host Hedgehog Signaling: a New Prospect for the Therapeutic Potential of Lipoxin A4.
Kaposi's sarcoma-associated herpesvirus immediate early gene activity.
Kaposi's sarcoma associated herpesvirus immune evasion strategies.
Beyond Viral Interferon Regulatory Factors: Immune Evasion Strategies.
Comparative pathobiology of Kaposi sarcoma-associated herpesvirus and related primate rhadinoviruses.
Kaposi's-sarcoma-associated-herpesvirus-activated dendritic cells promote HIV-1 trans-infection and suppress CD4(+) T cell proliferation.
Restoration of immune surface molecules in Kaposi sarcoma-associated herpes virus infected cells by lenalidomide and pomalidomide.
Prospects of a novel vaccination strategy for human gamma-herpesviruses.
Kaposi's sarcoma-associated herpesvirus infection of endothelial cells inhibits neutrophil recruitment through an interleukin-6-dependent mechanism: a new paradigm for viral immune evasion.
Immune mechanisms in murine gammaherpesvirus-68 infection.
Regulation of NF-kappaB inhibitor I kappaBalpha and viral replication by a KSHV microRNA.
Impaired CTL recognition of cells latently infected with Kaposi's sarcoma-associated herpes virus.
Immune control of oncogenic $\gamma$ -herpesviruses.
Kaposi's Sarcoma-Associated Herpesvirus Inhibitor of cGAS (KicGAS), Encoded by ORF52, Is an Abundant Tegument Protein and Is Required for Production of Infectious Progeny Viruses.
Kaposi's sarcoma-associated herpesvirus latent gene vFLIP inhibits viral lytic replication through NF-kappaB-mediated suppression of the AP-1 pathway: a novel mechanism of virus control of latency.
Molecular Virology of KSHV in the Lymphocyte Compartment-Insights From Patient Samples and De Novo Infection Models.
Quantitative Proteomics Analysis of Lytic KSHV Infection in Human Endothelial Cells Reveals Targets of Viral Immune Modulation.
Lack of CD8(+) T-cell co-localization with Kaposi's sarcoma-associated herpesvirus infected cells in Kaposi's sarcoma tumors.
Regulation of the Macroautophagic Machinery, Cellular Differentiation, and Immune Responses by Human Oncogenic $\gamma$ -Herpesviruses.
The oncogenic gamma herpesviruses Epstein-Barr virus (EBV) and Kaposi's sarcoma-associated herpesvirus (KSHV) hijack retinoic acid-inducible gene I (RIG-I) facilitating both viral and tumour immune evasion.
Intracellular-activated Notch1 can reactivate Kaposi's sarcoma-associated herpesvirus from latency.
Guanylate-Binding Protein 1 Inhibits Nuclear Delivery of Kaposi's Sarcoma-Associated Herpesvirus Virions by Disrupting Formation of Actin Filament.
Interference with the Autophagic Process as a Viral Strategy to Escape from the Immune Control: Lesson from Gamma Herpesviruses.

Proteomic screening of human targets of viral microRNAs reveals functions associated with immune evasion and angiogenesis.
Functions of Kaposi's sarcoma-associated herpesvirus microRNAs.
The viral interferon regulatory factors of kaposi's sarcoma-associated herpesvirus differ in their inhibition of interferon activation mediated by toll-like receptor 3.
Immune escape of $\gamma$ -herpesviruses from adaptive immunity.
Subversion of autophagy by Kaposi's sarcoma-associated herpesvirus impairs oncogene-induced senescence.
Inhibition of RIG-I-mediated signaling by Kaposi's sarcoma-associated herpesvirus-encoded deubiquitinase ORF64.
Ago HITS-CLIP expands understanding of Kaposi's sarcoma-associated herpesvirus miRNA function in primary effusion lymphomas.
Kaposi's sarcoma-associated herpesvirus viral interferon regulatory factor-2 inhibits type 1 interferon signalling by targeting interferon-stimulated gene factor-3.
Phase-dependent immune evasion of herpesviruses.
Immunobiology and host response to KSHV infection.
Evasion of oncogene-induced senescence by gammaherpesviruses.
Immune evasion strategies of the human gamma-herpesviruses: implications for viral tumorigenesis.
Concurrent expression of latent and a limited number of lytic genes with immune modulation and antiapoptotic function by Kaposi's sarcoma-associated herpesvirus early during infection of primary endothelial and fibroblast cells and subsequent decline of lytic gene expression.
The product of Kaposi's sarcoma-associated herpesvirus immediate early gene K4.2 regulates immunoglobulin secretion and calcium homeostasis by interacting with and inhibiting pERP1.
ORF45, a multifunctional immediate early and tegument protein of KSHV.
Disarming Cellular Alarm Systems-Manipulation of Stress-Induced NKG2D Ligands by Human Herpesviruses.
Kaposi's sarcoma-associated herpesvirus-encoded replication and transcription activator impairs innate immunity via ubiquitin-mediated degradation of myeloid differentiation factor 88.
Regulation of herpesvirus lifecycle by viral microRNAs.
Kaposi sarcoma herpesvirus pathogenesis.
The chromatin modification by SUMO-2/3 but not SUMO-1 prevents the epigenetic activation of key immune-related genes during Kaposi's sarcoma associated herpesvirus reactivation.
KSHV strategies for host dsDNA sensing machinery.
Kaposi's Sarcoma-Associated Herpesvirus Interleukin-6 Modulates Endothelial Cell Movement by Upregulating Cellular Genes Involved in Migration.
Viral subversion of autophagy impairs oncogene-induced senescence.
Kaposi's sarcoma-associated herpesvirus-encoded viral IRF3 modulates major histocompatibility complex class II (MHC-II) antigen presentation through MHC-II transactivator-dependent and -independent mechanisms: implications for oncogenesis.
Dissecting the regions of virion-associated Kaposi's sarcoma-associated herpesvirus complement control protein required for complement regulation and cell binding.
Discovery of a viral NLR homolog that inhibits the inflammasome.
Deletion of immune evasion genes provides an effective vaccine design for tumor-associated herpesviruses.
Comparative analysis of the viral interferon regulatory factors of KSHV for their requisite for virus production and inhibition of the type I interferon pathway.
Management of herpesvirus infections.

Kaposi's Sarcoma-Associated Herpesvirus Reduces Cellular Myeloid Differentiation Primary-Response Gene 88 (MyD88) Expression via Modulation of Its RNA.
Anti-viral and pro-inflammatory functions of Toll-like receptors during gamma-herpesvirus infections.
Kaposi sarcoma-associated herpesvirus latency-associated nuclear antigen inhibits interferon (IFN) beta expression by competing with IFN regulatory factor-3 for binding to IFNB promoter.
Human Oncogenic Herpesvirus and Post-translational Modifications - Phosphorylation and SUMOylation.
Modulation of interferon regulatory factor 5 activities by the Kaposi sarcoma-associated herpesvirus-encoded viral interferon regulatory factor 3 contributes to immune evasion and lytic induction.
Kaposi sarcoma herpesvirus-encoded vFLIP and vIRF1 regulate antigen presentation in lymphatic endothelial cells.
A human herpesvirus miRNA attenuates interferon signaling and contributes to maintenance of viral latency by targeting IKKε.
The Heme Metabolite Carbon Monoxide Facilitates KSHV Infection by Inhibiting TLR4 Signaling in Endothelial Cells.
Genome-Wide Mapping of the Binding Sites and Structural Analysis of Kaposi's Sarcoma-Associated Herpesvirus Viral Interferon Regulatory Factor 2 Reveal that It Is a DNA-Binding Transcription Factor.
Herpesviruses hijack host exosomes for viral pathogenesis.
Kaposi's sarcoma-associated herpesvirus viral interferon regulatory factor 3 inhibits gamma interferon and major histocompatibility complex class II expression.
Diverse herpesvirus microRNAs target the stress-induced immune ligand MICB to escape recognition by natural killer cells.
Productive herpesvirus lytic replication in primary effusion lymphoma cells requires S-phase entry.
Intracellular Kaposi's sarcoma-associated herpesvirus load determines early loss of immune synapse components.
Kaposi sarcoma-associated herpesvirus degrades cellular Toll-interleukin-1 receptor domain-containing adaptor-inducing beta-interferon (TRIF).
Computational analysis predicts the Kaposi's sarcoma-associated herpesvirus tegument protein ORF63 to be alpha helical.
HMGB1, a potential regulator of tumor microenvironment in KSHV-infected endothelial cells.
Human herpesvirus-8: Kaposi sarcoma, multicentric Castlemann disease, and primary effusion lymphoma.
Kaposi's sarcoma-associated herpesvirus microRNAs target IRAK1 and MYD88, two components of the toll-like receptor/interleukin-1R signaling cascade, to reduce inflammatory-cytokine expression.
A seek-and-hide game between Cd1-restricted T cells and herpesviruses.
Rodent herpesvirus Peru encodes a secreted chemokine decoy receptor.
The association between human herpesviruses and periodontal disease: Part 1. Herpesviruses immune evasion. A review.
The Kaposi's Sarcoma-associated Herpesvirus-encoded vIRF-3 Inhibits Cellular IRF-5.
Kaposi's sarcoma-associated herpesvirus K3 and K5 ubiquitin E3 ligases have stage-specific immune evasion roles during lytic replication.
Kaposi's sarcoma-associated herpesvirus-encoded latency-associated nuclear antigen reduces interleukin-8 expression in endothelial cells and impairs neutrophil chemotaxis by degrading nuclear p65.
KSHV ORF K9 (vIRF) is an oncogene which inhibits the interferon signaling pathway.
Immune evasion by gamma-herpesviruses.



Multi-targeted therapy of everolimus in Kaposi's sarcoma associated herpes virus infected primary effusion lymphoma.
Kaposi's sarcoma-associated herpesvirus latency-associated nuclear antigen 1 mimics Epstein-Barr virus EBNA1 immune evasion through central repeat domain effects on protein processing.
Immune evasion by gammaherpesvirus genome maintenance proteins.
Human gammaherpesvirus immune evasion strategies.
Herpesviral microRNAs in Cellular Metabolism and Immune Responses.
Identification of AIDS-Associated Kaposi Sarcoma: A Functional Genomics Approach.
KRAB-ZFP Repressors Enforce Quiescence of Oncogenic Human Herpesviruses.
Kaposi's sarcoma-associated herpesvirus noncoding polyadenylated nuclear RNA interacts with virus- and host cell-encoded proteins and suppresses expression of genes involved in immune modulation.
Plasmacytoid dendritic cells in skin lesions of classic Kaposi's sarcoma.
Kaposi's sarcoma-associated herpesvirus ubiquitin ligases downregulate cell surface expression of I-selectin.
Immune evasion strategies of the herpesviruses.
Getting the message direct manipulation of host mRNA accumulation during gammaherpesvirus lytic infection.
Herpesvirus latency confers symbiotic protection from bacterial infection.
Conquering the Host: Determinants of Pathogenesis Learned from Murine Gammaherpesvirus 68.
Long Non-Coding RNAs: Novel Players in Regulation of Immune Response Upon Herpesvirus Infection.
The inflammatory component of Kaposi sarcoma.
A gamma-herpesvirus deficient in replication establishes chronic infection in vivo and is impervious to restriction by adaptive immune cells.
MicroRNAs in large herpesvirus DNA genomes: recent advances.
Murine herpesvirus pathogenesis: a model for the analysis of molecular mechanisms of human gamma herpesvirus infections.
Chemokine Subversion by Human Herpesviruses.
Natural history of murine gamma-herpesvirus infection.
Selective inhibition of miRNA processing by a herpesvirus-encoded miRNA.
The DNase of gammaherpesviruses impairs recognition by virus-specific CD8+ T cells through an additional host shutoff function.
Constitutive activation of T cells by $\gamma$ 2-herpesviral GPCR through the interaction with cellular CXCR4.
Evasion of innate cytosolic DNA sensing by a gammaherpesvirus facilitates establishment of latent infection.
Small RNAs growing tall: miRNAs as drug targets in herpesvirus infections.
Kaposi' s sarcoma in HIV-positive patients: the state of art in the HAART-era.
Murine gammaherpesvirus (MHV) MK3 gene sequence diversity among 72, 4556, and 68 strains.
Identification of functional domains in kaposica, the complement control protein homolog of Kaposi's sarcoma-associated herpesvirus (human herpesvirus 8).
Gene expression and transcription factor profiling reveal inhibition of transcription factor cAMP-response element-binding protein by gamma-herpesvirus replication and transcription activator.
Transcription pattern of human herpesvirus 8 open reading frame K3 in primary effusion lymphoma and Kaposi's sarcoma.
Autocrine and paracrine promotion of cell survival and virus replication by human herpesvirus 8 chemokines.

Molecular mechanism of BST2/tetherin downregulation by K5/MIR2 of Kaposi's sarcoma-associated herpesvirus.
Solution structure of the Kaposi's sarcoma-associated herpesvirus K3 N-terminal domain reveals a Novel E2-binding C4HC3-type RING domain.
Immune control of mammalian gamma-herpesviruses: lessons from murid herpesvirus-4.
Regulation of CD1d expression and function by a herpesvirus infection.
Deletion of Murid Herpesvirus 4 ORF63 Affects the Trafficking of Incoming Capsids toward the Nucleus.
Cooperative DNA binding mediated by KicGAS/ORF52 oligomerization allows inhibition of DNA-induced phase separation and activation of cGAS.
NK cell activity controls human herpesvirus 8 latent infection and is restored upon highly active antiretroviral therapy in AIDS patients with regressing Kaposi's sarcoma.
Masters of deception: a review of herpesvirus immune evasion strategies.
The Role of microRNAs in the Pathogenesis of Herpesvirus Infection.
Evolutionary effects of the AID/APOBEC family of mutagenic enzymes on human gamma-herpesviruses.
Immune regulation of human herpesviruses and its implications for human transplantation.
Human cytomegalovirus miR-UL112-1 promotes the down-regulation of viral immediate early-gene expression during latency to prevent T-cell recognition of latently infected cells.
Kaposi's sarcoma-associated herpesvirus ORF57 promotes escape of viral and human interleukin-6 from microRNA-mediated suppression.
Chemokines encoded by herpesviruses.
Structural Proteomics of Herpesviruses.
Manipulation of ubiquitin/SUMO pathways in human herpesviruses infection.
Virally encoded 7TM receptors.
Autophagy in herpesvirus immune control and immune escape.
The intertransmembrane region of Kaposi's sarcoma-associated herpesvirus modulator of immune recognition 2 contributes to B7-2 downregulation.
Chemokine-binding proteins encoded by herpesviruses.
Kaposi sarcoma.
The central repeat domain 1 of Kaposi's sarcoma-associated herpesvirus (KSHV) latency associated-nuclear antigen 1 (LANA1) prevents cis MHC class I peptide presentation.
Human herpesviridae methods of natural killer cell evasion.
Messenger RNA turnover and its regulation in herpesviral infection.
Immune escape of equine herpesvirus 1 and other herpesviruses of veterinary importance.
Aberrant herpesvirus-induced polyadenylation correlates with cellular messenger RNA destruction.
Illumination of murine gammaherpesvirus-68 cycle reveals a sexual transmission route from females to males in laboratory mice.
The different activities of RNA G-quadruplex structures are controlled by flanking sequences.
Herpesviruses induce aggregation and selective autophagy of host signalling proteins NEMO and RIPK1 as an immune-evasion mechanism.
Repression of interferon- $\alpha$ stimulated genes expression by Kaposi's sarcoma-associated herpesvirus K-bZIP protein.
A basic charge cluster near the C-terminus of the cytoplasmic tail contributes to the molecular stability of human herpesvirus 8 E3 ubiquitin ligases.
An internal ribosome entry site directs translation of the murine gammaherpesvirus 68 MK3 open reading frame.

Modulation of cellular signaling by herpesvirus-encoded G protein-coupled receptors.
Regulation of telomerase and telomeres: human tumor viruses take control.
Marek's disease: Genetic regulation of gallid herpesvirus 2 infection and latency.
Intrinsic and Innate Defenses of Neurons: Détente with the Herpesviruses.
MHV-68 producing mIFN $\alpha$ 1 is severely attenuated in vivo and effectively protects mice against challenge with wt MHV-68.
The Critical Role of Genome Maintenance Proteins in Immune Evasion During Gammaherpesvirus Latency.
Critical role of complement and viral evasion of complement in acute, persistent, and latent gamma-herpesvirus infection.
Emerging roles of chicken and viral microRNAs in avian disease.
Virally encoded chemokine binding proteins.
In cis inhibition of antigen processing by the latency-associated nuclear antigen I of Kaposi sarcoma herpes virus.
A viral conspiracy: hijacking the chemokine system through virally encoded pirated chemokine receptors.
Structure, function and physiological consequences of virally encoded chemokine seven transmembrane receptors.
Identification and sequencing of a novel rodent gammaherpesvirus that establishes acute and latent infection in laboratory mice.
Identification of a gammaherpesvirus selective chemokine binding protein that inhibits chemokine action.
The HHV-6A Proteins U20 and U21 Target NKG2D Ligands to Escape Immune Recognition.
Herpesvirus microRNAs for use in gene therapy immune-evasion strategies.
Disruption of CCL21-induced chemotaxis in vitro and in vivo by M3, a chemokine-binding protein encoded by murine gammaherpesvirus 68.
Virus-encoded 7 transmembrane receptors.
LANA and hnRNP A1 Regulate the Translation of LANA mRNA through G-Quadruplexes.
[Research advances in herpesviruses glycoprotein N gene and its encoded protein].
Immune modulation by virus-encoded secreted chemokine binding proteins.
T-cell responses to the M3 immune evasion protein of murid gammaherpesvirus 68 are partially protective and induced with lytic antigen kinetics.
Human Herpesvirus 6B Downregulates Expression of Activating Ligands during Lytic Infection To Escape Elimination by Natural Killer Cells.
Viral Encoded miRNAs in Tumorigenesis: Theranostic Opportunities in Precision Oncology.
The DNA sequence of human herpesvirus-6: structure, coding content, and genome evolution.
Kinetics of murine gammaherpesvirus 68 gene expression following infection of murine cells in culture and in mice.
Viral miRNAs: tools for immune evasion.
Partial genome sequence of murine gammaherpesvirus 72 and its analysis.
Biology and host response to Cyprinid herpesvirus 3 infection in common carp.
A Murine Herpesvirus Closely Related to Ubiquitous Human Herpesviruses Causes T-Cell Depletion.
Reduction in RNA levels rather than retardation of translation is responsible for the inhibition of major histocompatibility complex class I antigen presentation by the glutamic acid-rich repeat of herpesvirus saimiri open reading frame 73.
Viral chemokine-modulatory proteins: tools and targets.

Evasion of adaptive and innate immune response mechanisms by $\gamma$ -herpesviruses.
The m4 gene of murine gammaherpesvirus modulates productive and latent infection in vivo.
Viral semaphorin inhibits dendritic cell phagocytosis and migration but is not essential for gammaherpesvirus-induced lymphoproliferation in malignant catarrhal fever.
Inhibition of interleukin-2 gene expression by human herpesvirus 6B U54 tegument protein.
Koala and Wombat Gammaherpesviruses Encode the First Known Viral NTPDase Homologs and Are Phylogenetically Divergent from All Known Gammaherpesviruses.
Possible role of human herpesvirus 6 as a trigger of autoimmune disease.
Partial genome analysis of murine gammaherpesvirus 4556.
Differential HHV-6A gene expression in T cells and primary human astrocytes based on multi-virus array analysis.
Blockade of chemokine activity by a soluble chemokine binding protein from vaccinia virus.
vCCL2/vMIP-II, the viral master KEYmokine.
B cells latently infected with murine gammaherpesvirus 68 (MHV-68) are present in the mouse thymus-A step toward immune evasion?
The chemokine decoy receptor M3 blocks CC chemokine ligand 2 and CXC chemokine ligand 13 function in vivo.
Interactions between herpesvirus entry mediator (TNFRSF14) and latency-associated transcript during herpes simplex virus 1 latency.
Identification and characterization of U83A viral chemokine, a broad and potent beta-chemokine agonist for human CCRs with unique selectivity and inhibition by spliced isoform.
Macavirus latency-associated protein evades immune detection through regulation of protein synthesis in cis depending upon its glycin/glutamate-rich domain.
A novel highly potent therapeutic antibody neutralizes multiple human chemokines and mimics viral immune modulation.
HHV-8 reduces dendritic cell migration through down-regulation of cell-surface CCR6 and CCR7 and cytoskeleton reorganization.
Human and viral membrane-associated E3 ubiquitin ligases MARCH1 and MIR2 recognize different features of CD86 to downregulate surface expression.
Mechanism of action of the viral chemokine-binding protein E163 from ectromelia virus.
Immunomodulation by herpesvirus U51A chemokine receptor via CCL5 and FOG-2 down-regulation plus XCR1 and CCR7 mimicry in human leukocytes.
Selective recruitment of Th2-type cells and evasion from a cytotoxic immune response mediated by viral macrophage inhibitory protein-II.
MicroRNA based immunoevasion mechanism of human polyomaviruses.
Establishment of HSV1 latency in immunodeficient mice facilitates efficient in vivo reactivation.
Molecular characterization of Cyprinid herpesvirus 3 encoded viral interleukin10.
Lentiviral latency in peripheral CD4+ T cells isolated from feline immunodeficiency virus-infected cats during the asymptomatic phase is not associated with hypermethylation of the proviral promoter.

Table A2-26, Cluster 25

Cluster 25 focuses on the role of immunosuppressive regulatory T cells in cancer immunology (which augment tumor development and progression by inhibiting antitumor immunity), emphasizing strategies to deplete Treg cells and control Treg cell functions to increase antitumor immune responses (472)
---

Regulatory T cells in gastrointestinal tumors.
T regulatory cells in cancer: recent advances and therapeutic potential.
The role of regulatory T cells in cancer immunology.
Regulatory T Cells in Ovarian Carcinogenesis and Future Therapeutic Opportunities.
Regulatory T cells as adjuvant target for enhancing the viral disease vaccine efficacy.
FOXP3+ regulatory T cells: Current controversies and future perspectives.
Regulatory T cells and potential immunotherapeutic targets in lung cancer.
Regulatory T cells in breast cancer as a potent anti-cancer therapeutic target.
Monitoring regulatory T cells in clinical samples: consensus on an essential marker set and gating strategy for regulatory T cell analysis by flow cytometry.
The role of regulatory T cells in malignant glioma.
Regulatory T Cells in Autoimmunity and Cancer: A Duplicitous Lifestyle.
Immunotherapy approaches targeting regulatory T-cells.
FOXP3+ Tregs: heterogeneous phenotypes and conflicting impacts on survival outcomes in patients with colorectal cancer.
Role of heterogeneous regulatory T cells in the tumor microenvironment.
Increased intratumoral regulatory T cells are related to intratumoral macrophages and poor prognosis in hepatocellular carcinoma patients.
Immunosuppressive effect of regulatory T lymphocytes in lung cancer, with special reference to their effects on the induction of autologous tumor-specific cytotoxic T lymphocytes.
Immunotherapy of Cancer by Targeting Regulatory T cells.
Different subpopulations of regulatory T cells in human autoimmune disease, transplantation, and tumor immunity.
Molecular Signatures of Human Regulatory T Cells in Colorectal Cancer and Polyps.
Intranuclear interactomic inhibition of FoxP3 suppresses functions of Treg cells.
Roles of regulatory T cells in cancer immunity.
Targeting CD4+CD25+FoxP3+ regulatory T-cells for the augmentation of cancer immunotherapy.
Regulatory T Cells in Gynecologic Cancer.
Requirement of Treg-intrinsic CTLA4/PKC $\eta$ signaling pathway for suppressing tumor immunity.
The Dual Role of Treg in Cancer.
Modulating Treg stability to improve cancer immunotherapy.
Regulatory T Cells in an Endogenous Mouse Lymphoma Recognize Specific Antigen Peptides and Contribute to Immune Escape.
Treg Fragility: A Prerequisite for Effective Antitumor Immunity?
Regulatory T cells in the immunotherapy of melanoma.
Local and systemic induction of CD4+CD25+ regulatory T-cell population by non-Hodgkin lymphoma.
TNFR2-expressing CD4(+)Foxp3(+) regulatory T cells in cancer immunology and immunotherapy.
Tumor-infiltrating T-regulatory cells adapt to altered metabolism to promote tumor-immune escape.
Unlocking the Complexities of Tumor-Associated Regulatory T Cells.
Intratumoral regulatory T cells: markers, subsets and their impact on anti-tumor immunity.
Regulatory T cells as therapeutic targets and mediators.
Neoadjuvant chemo-immunotherapy modifies CD4(+)CD25(+) regulatory T cells (Treg) in non-small cell lung cancer (NSCLC) patients.
CD4+CD25hiCD127low regulatory T cells are increased in oral squamous cell carcinoma patients.
Crosstalk of Microorganisms and Immune Responses in Autoimmune Neuroinflammation: A Focus on Regulatory T Cells.

Induced regulatory T cells in inhibitory microenvironments created by cancer.
T-regulatory cells: key players in tumor immune escape and angiogenesis.
Regulatory T cell depletion enhances tumor specific CD8 T-cell responses, elicited by tumor antigen NY-ESO-1b in hepatocellular carcinoma patients, in vitro.
Regulatory T cells in melanoma: the final hurdle towards effective immunotherapy?
Hyperfunction of CD4 CD25 regulatory T cells in de novo acute myeloid leukemia.
Transcriptional regulation of FOXP3 requires integrated activation of both promoter and CNS regions in tumor-induced CD8(+) Treg cells.
Metabolic Regulation of Tregs in Cancer: Opportunities for Immunotherapy.
Treg(s) in Cancer: Friends or Foe?
PD-L1 Is Expressed and Promotes the Expansion of Regulatory T Cells in Acute Myeloid Leukemia.
Presence of regulatory T-cells in endometrial cancer predicts poorer overall survival and promotes progression of tumor cells.
Mechanisms of regulatory T cell infiltration in tumors: implications for innovative immune precision therapies.
Antibody-based cancer immunotherapy by targeting regulatory T cells.
Intratumoral regulatory T cells with higher prevalence and more suppressive activity in hepatocellular carcinoma patients.
Distepharinamide, a novel dimeric proaporphine alkaloid from <i>Diplocisia glaucescens</i> , inhibits the differentiation and proliferative expansion of CD4(+)Foxp3(+) regulatory T cells.
Role of Regulatory T Cells (Treg) and the Treg Effector Molecule Fibrinogen-like Protein 2 in Alloimmunity and Autoimmunity.
Isolation and Analysis of Tumor-Infiltrating Treg.
Regulation of regulatory T cells in cancer.
Heat Shock Protein 60 in Eggs Specifically Induces Tregs and Reduces Liver Immunopathology in Mice with Schistosomiasis Japonica.
Regulatory T cell: a protection for tumour cells.
Treg programming and therapeutic reprogramming in cancer.
Colorectal Cancer-Infiltrating Regulatory T Cells: Functional Heterogeneity, Metabolic Adaptation, and Therapeutic Targeting.
Searching for Peptide Inhibitors of T Regulatory Cell Activity by Targeting Specific Domains of FOXP3 Transcription Factor.
T Regulatory Cells and Priming the Suppressive Tumor Microenvironment.
Tumor-infiltrating FoxP3(+) Tregs predict favorable outcome in colorectal cancer patients: A meta-analysis.
Immune checkpoint inhibitors in cancer therapy: a focus on T-regulatory cells.
Manipulation of regulatory T cells and antigen-specific cytotoxic T lymphocyte-based tumour immunotherapy.
Modulation of regulatory T cells by natural products in cancer.
CD4+FoxP3+ regulatory T cells gradually accumulate in gliomas during tumor growth and efficiently suppress antiglioma immune responses in vivo.
Regulatory T cells in cancer: an overview and perspectives on cyclooxygenase-2 and Foxp3 DNA methylation.
Targeting of Cdc42 GTPase in regulatory T cells unleashes antitumor T-cell immunity.
Lkb1 aggravates diffuse large B-cell lymphoma by promoting the function of Treg cells and immune escape.
Targeting regulatory T cells for anticancer therapy.

Activated tumor-infiltrating CD4+ regulatory T cells restrain antitumor immunity in patients with primary or metastatic liver cancer.
Hepatitis C Virus Improves Human Tregs Suppressive Function and Promotes Their Recruitment to the Liver.
[Correlation analysis of the Epstein-Barr virus recruited regulatory T cells in nasopharyngeal carcinoma of immune escape].
IL-8 in bone marrow and peripheral blood of patients with B-cell acute lymphoblastic leukemia is associated with high regulatory T cell counts, degree of tumor infiltration and expression of CXCR1 in blasts.
[Anti-CCR4 mAb and regulatory T cells].
Cyclophosphamide inhibits the generation and function of CD8(+) regulatory T cells.
Regulatory T cells in cancer: where are we now?
Disruption of CCR5-dependent homing of regulatory T cells inhibits tumor growth in a murine model of pancreatic cancer.
Fetal alloantigen is responsible for the expansion of the CD4(+)CD25(+) regulatory T cell pool during pregnancy.
FOXP3+ regulatory T cells and the immune escape in solid tumours.
Increased ectonucleotidase expression and activity in regulatory T cells of patients with head and neck cancer.
Permanent up-regulation of regulatory T-lymphocytes in patients with head and neck cancer.
Inducers, Attractors and Modulators of CD4(+) Treg Cells in Non-Small-Cell Lung Cancer.
The frequency and suppressor function of CD4+CD25highFoxp3+ T cells in the circulation of patients with squamous cell carcinoma of the head and neck.
Treg-specific demethylated region activity in isolated regulatory t lymphocytes is a surrogate for disease severity in hepatocellular carcinoma.
Remodeling of the tumor microenvironment via disrupting Blimp1(+) effector Treg activity augments response to anti-PD-1 blockade.
Deciphering the Role of Regulatory CD4 T Cells in Oral and Oropharyngeal Cancer: A Systematic Review.
Protein-bound polysaccharide-K reduces the proportion of regulatory T cells in vitro and in vivo.
Up-regulation of proliferative and migratory genes in regulatory T cells from patients with metastatic castration-resistant prostate cancer.
Clinical significance of the frequency of regulatory T cells in regional lymph node lymphocytes as a prognostic factor for non-small-cell lung cancer.
Regulatory T cell-targeted hybrid nanoparticles combined with immuno-checkpoint blockage for cancer immunotherapy.
Regulatory T cells (Tregs) in lymphoid malignancies and the impact of novel therapies.
Targeting regulatory T cells for improving cancer therapy: Challenges and prospects.
CD27 expression on Treg cells limits immune responses against tumors.
Phenotypic alterations, clinical impact and therapeutic potential of regulatory T cells in cancer.
Treg Cells Promote the SREBP1-Dependent Metabolic Fitness of Tumor-Promoting Macrophages via Repression of CD8(+) T Cell-Derived Interferon-γ.
The regulatory T-cell response during acute retroviral infection is locally defined and controls the magnitude and duration of the virus-specific cytotoxic T-cell response.
The Antitumor Effect of Xihuang Pill on Treg Cells Decreased in Tumor Microenvironment of 4T1 Breast Tumor-Bearing Mice by PI3K/AKT~AP-1 Signaling Pathway.

Conversion of intratumoral regulatory T cells by human gastric cancer cells is dependent on transforming growth factor- $\beta$ 1.
Selective inhibition of regulatory T cells by targeting the PI3K-Akt pathway.
<i>Strongyloides ratti</i> infection induces expansion of Foxp3+ regulatory T cells that interfere with immune response and parasite clearance in BALB/c mice.
Precise Spatiotemporal Interruption of Regulatory T-cell-Mediated CD8(+) T-cell Suppression Leads to Tumor Immunity.
Cutaneous and oral squamous cell carcinoma-dual immunosuppression via recruitment of FOXP3+ regulatory T cells and endogenous tumour FOXP3 expression?
Tumor-derived microvesicles induce, expand and up-regulate biological activities of human regulatory T cells (Treg).
The immunology of pregnancy: regulatory T cells control maternal immune tolerance toward the fetus.
Inhibitory Receptors and Pathways of Lymphocytes: The Role of PD-1 in Treg Development and Their Involvement in Autoimmunity Onset and Cancer Progression.
Activated CTLA-4-independent immunosuppression of Treg cells disturbs CTLA-4 blockade-mediated antitumor immunity.
Interaction of indoleamine-2,3-dioxygenase and CD4+CD25+ regulatory T cells in tumor immune escape.
Regulatory T cells and plasmacytoid dendritic cells contribute to the immune escape of papillary thyroid cancer coexisting with multinodular non-toxic goiter.
4-1BBL-containing leukemic extracellular vesicles promote immunosuppressive effector regulatory T cells.
HIF-1 $\alpha$ Is a Metabolic Switch between Glycolytic-Driven Migration and Oxidative Phosphorylation-Driven Immunosuppression of Tregs in Glioblastoma.
Expression of indoleamine 2, 3-dioxygenase and the recruitment of Foxp3-expressing regulatory T cells in the development and progression of uterine cervical cancer.
Providence of the CD25(+) KIR(+) CD127(-) FOXP3(-) CD8(+) T-cell subset determines the dynamics of tumor immune surveillance.
Single-Cell Sequencing Reveals the Transcriptome and TCR Characteristics of pTregs and in vitro Expanded iTregs.
Helios serves as a suppression marker to reduce regulatory T cell function in pancreatic cancer patients.
Regulatory T cells in oral squamous cell carcinoma.
Regulatory T-cell expansion during chronic viral infection is dependent on endogenous retroviral superantigens.
Pathogen-specific Treg cells expand early during mycobacterium tuberculosis infection but are later eliminated in response to Interleukin-12.
Antitumor T cell responses in bladder cancer are directed against a limited set of antigens and are modulated by regulatory T cells and routine treatment approaches.
IL-21 Is an Accomplice of PD-L1 in the Induction of PD-1-Dependent Treg Generation in Head and Neck Cancer.
Overcoming immunoescape mechanisms of BCL1 leukemia and induction of CD8+ T-cell-mediated BCL1-specific resistance in mice cured by targeted polymer-bound doxorubicin.
Tumor-infiltrating regulatory T cells inhibit endogenous cytotoxic T cell responses to lung adenocarcinoma.



Adenosine Generated by Regulatory T Cells Induces CD8(+) T Cell Exhaustion in Gastric Cancer through A2aR Pathway.
Trafficking of FoxP3+ regulatory T cells: myths and facts.
Renal cell carcinoma may evade the immune system by converting CD4+Foxp3- T cells into CD4+CD25+Foxp3+ regulatory T cells: Role of tumor COX-2-derived PGE2.
Altering regulatory T cell function in cancer immunotherapy: a novel means to boost the efficacy of cancer vaccines.
Radiation enhances regulatory T cell representation.
The immune checkpoint B7x expands tumor-infiltrating Tregs and promotes resistance to anti-CTLA-4 therapy.
Behavior of circulating CD4+CD25+Foxp3+ regulatory T cells in colon cancer patients undergoing surgery.
Graded RhoA GTPase Expression in Treg Cells Distinguishes Tumor Immunity From Autoimmunity.
Immune modulation by regulatory T cells in Helicobacter pylori-associated diseases.
Targeting regulatory T cells.
Analysis of FOXP3+ regulatory T cells that display apparent viral antigen specificity during chronic hepatitis C virus infection.
Blockade of CCL1 inhibits T regulatory cell suppressive function enhancing tumor immunity without affecting T effector responses.
CCL20-CCR6 Cytokine Network Facilitate Treg Activity in Advanced Grades and Metastatic Variants of Hepatocellular Carcinoma.
Deregulated Mucosal Immune Surveillance through Gut-Associated Regulatory T Cells and PD-1(+) T Cells in Human Colorectal Cancer.
Metronomic cyclophosphamide treatment in metastasized breast cancer patients: immunological effects and clinical outcome.
T-cell immunoglobulin and ITIM domain in cancer immunotherapy: A focus on tumor-infiltrating regulatory T cells.
Tumor cells induced-M2 macrophage favors accumulation of Treg in nasopharyngeal carcinoma.
Frequency of CD4+CD25+Foxp3+ cells in peripheral blood in relation to urinary bladder cancer malignancy indicators before and after surgical removal.
INFα-2b inhibitory effects on CD4(+)CD25(+)FOXP3(+) regulatory T cells in the tumor microenvironment of C57BL/6 J mice with melanoma xenografts.
Adenocarcinoma contains more immune tolerance regulatory t-cell lymphocytes (versus squamous carcinoma) in non-small-cell lung cancer.
Cancer immunotherapy with check point inhibitor can cause autoimmune adverse events due to loss of Treg homeostasis.
RA8, a human anti-CD25 antibody against human Treg cells.
Treg-mediated acquired resistance to immune checkpoint inhibitors.
Expression of TNFR2 by regulatory T cells in peripheral blood is correlated with clinical pathology of lung cancer patients.
CCR8 blockade primes anti-tumor immunity through intratumoral regulatory T cells destabilization in muscle-invasive bladder cancer.
Tumor-secreted miR-214 induces regulatory T cells: a major link between immune evasion and tumor growth.
Analysis of CD8+ Treg cells in patients with ovarian cancer: a possible mechanism for immune impairment.

Systems biology analysis identifies TNFRSF9 as a functional marker of tumor-infiltrating regulatory T-cell enabling clinical outcome prediction in lung cancer.
Induction of stable human FOXP3(+) Tregs by a parasite-derived TGF- $\beta$ mimic.
Acute Myeloid Leukemia Cells Express ICOS Ligand to Promote the Expansion of Regulatory T Cells.
The role of FoxP3+ regulatory T cells and IDO+ immune and tumor cells in malignant melanoma - an immunohistochemical study.
Cutting edge: expression of TNFR2 defines a maximally suppressive subset of mouse CD4+CD25+FoxP3+ T regulatory cells: applicability to tumor-infiltrating T regulatory cells.
Natural regulatory (CD4+CD25+FOXP+) T cells control the production of pro-inflammatory cytokines during Plasmodium chabaudi adami infection and do not contribute to immune evasion.
Prognostic value of regulatory T cells in newly diagnosed chronic myeloid leukemia patients.
A TNFR2 antibody by countering immunosuppression cooperates with HMGN1 and R848 immune stimulants to inhibit murine colon cancer.
Foxp3 expression in melanoma cells as a possible mechanism of resistance to immune destruction.
Solanine Inhibits Immune Escape Mediated by Hepatoma Treg Cells via the TGF $\beta$ /Smad Signaling Pathway.
FOXP3+ Cells Recruited by CCL22 into Breast Cancer Correlates with Less Tumor Nodal Infiltration.
SA-4-1BBL costimulation inhibits conversion of conventional CD4+ T cells into CD4+ FoxP3+ T regulatory cells by production of IFN- $\gamma$ .
Roles of Tregs in development of hepatocellular carcinoma: a meta-analysis.
Follicular lymphoma B cells induce the conversion of conventional CD4+ T cells to T-regulatory cells.
Myeloid-derived suppressor cells from tumor-bearing mice impair TGF- $\beta$ -induced differentiation of CD4+CD25+FoxP3+ Tregs from CD4+CD25-FoxP3- T cells.
Steroid receptor coactivator 3 (SRC-3/AIB1) is enriched and functional in mouse and human Tregs.
After Treatment Decrease of Bone Marrow Tregs and Outcome in Younger Patients with Newly Diagnosed Acute Myeloid Leukemia.
Radiofrequency ablation does not induce the significant increase of CD4(+) CD25(+) Foxp3(+) regulatory T cells compared with surgical resection in Hepal-6 tumor model.
T regulatory cells control T-cell proliferation partly by the release of soluble CD25 in patients with B-cell malignancies.
Localization of FOXP3-positive cells in renal cell carcinoma.
IL-16 processing in sentinel node regulatory T cells is a factor in bladder cancer immunity.
Increased prevalence of regulatory T cells (Treg) is induced by pancreas adenocarcinoma.
"High Treg" Inflammations Promote (Most) Non-Hematologic Cancers While "Low Treg" Inflammations Promote Lymphoid Cancers.
Expression of indoleamine 2,3-dioxygenase predicts shorter survival in patients with vulvar squamous cell carcinoma (vSCC) not influencing on the recruitment of FOXP3-expressing regulatory T cells in cancer nests.
Regulatory T cells recruited through CCL22/CCR4 are selectively activated in lymphoid infiltrates surrounding primary breast tumors and lead to an adverse clinical outcome.
CD4+CD25+ T regulatory cells dominate multiple immune evasion mechanisms in early but not late phases of tumor development in a B cell lymphoma model.
Changes of regulatory T cells and FoxP3 gene expression in the aging process and its relationship with lung tumors in humans and mice.
The glutathione peroxidase Gpx4 prevents lipid peroxidation and ferroptosis to sustain Treg cell activation and suppression of antitumor immunity.

Induction of CD4(+)CD25(+)FOXP3(+) regulatory T cells during human hookworm infection modulates antigen-mediated lymphocyte proliferation.
Myeloid-derived suppressor cells promote cross-tolerance in B-cell lymphoma by expanding regulatory T cells.
Oncogenic BRAF(V600E) Governs Regulatory T-cell Recruitment during Melanoma Tumorigenesis.
Intratumor hypoxia promotes immune tolerance by inducing regulatory T cells via TGF- $\beta$ 1 in gastric cancer.
Analysis of Treg cell population alterations in the peripheral blood of patients treated surgically for ovarian cancer - a preliminary report.
Increased expression of CD4+CD25 +FOXP3+ regulatory T cells correlates with Epstein-Barr virus and has no impact on survival in patients with classical Hodgkin lymphoma in Brazil.
Suppression of intratumoral CCL22 by type I interferon inhibits migration of regulatory T cells and blocks cancer progression.
Depletion of Foxp3+ regulatory T cells but not the absence of CD19+IL-10+ regulatory B cells hinders tumor growth in a para-orthotopic neuroblastoma mouse model.
Toward understanding the genetics of regulatory T cells in ovarian cancer.
Secretory High-Mobility Group Box 1 Protein Affects Regulatory T Cell Differentiation in Neuroblastoma Microenvironment In Vitro.
Patients with oral squamous cell carcinoma are characterized by increased frequency of suppressive regulatory T cells in the blood and tumor microenvironment.
Immune responses regulation following antitumor dendritic cell-based prophylactic, concurrent, and therapeutic vaccination.
Tregs and Platelets Play Synergistic Roles in Tumor Immune Escape and Inflammatory Diseases.
Superantigen-induced proliferation of human CD4+CD25- T cells is followed by a switch to a functional regulatory phenotype.
Forkhead box M1 recruits FoxP3(+) Treg cells to induce immune escape in hilar cholangiocarcinoma.
Sphingosine 1 phosphate receptor-1 (S1P1) promotes tumor-associated regulatory T cell expansion: leading to poor survival in bladder cancer.
Effector/memory but not naive regulatory T cells are responsible for the loss of concomitant tumor immunity.
CD4+CD25+FOXP3+ T regulatory cells reconstitute and accumulate in the bone marrow of patients with multiple myeloma following allogeneic stem cell transplantation.
Increased CTLA-4 and FOXP3 transcripts in peripheral blood mononuclear cells of patients with breast cancer.
HMGB1 is overexpressed in tumor cells and promotes activity of regulatory T cells in patients with head and neck cancer.
Upregulated expression of indoleamine 2, 3-dioxygenase in CHO cells induces apoptosis of competent T cells and increases proportion of Treg cells.
Dynamic Treg interactions with intratumoral APCs promote local CTL dysfunction.
Increased frequency and suppressive activity of CD127(low/-) regulatory T cells in the peripheral circulation of patients with head and neck squamous cell carcinoma are associated with advanced stage and nodal involvement.
The expression of the regulatory T cell-specific forkhead box transcription factor FoxP3 is associated with poor prognosis in ovarian cancer.
microRNA-128-3p inhibits CD4+ regulatory T cells enrichment by targeting interleukin 16 in gastric cancer.
Hydrogen Sulfide Creates a Favorable Immune Microenvironment for Colon Cancer.

Staphylococcus aureus Protein A Induces Human Regulatory T Cells Through Interaction With Antigen-Presenting Cells.
T Regulatory Cell Subpopulations Associated with Recent Ultraviolet Radiation Exposure in a Skin Cancer Screening Cohort.
3-Methylcholanthrene-induced transforming growth factor-beta-producing carcinomas, but not sarcomas, are refractory to regulatory T cell-depletion therapy.
FOXP3+ Tregs and B7-H1+/PD-1+ T lymphocytes co-infiltrate the tumor tissues of high-risk breast cancer patients: Implication for immunotherapy.
Despite increased CD4+Foxp3+ cells within the infection site, BALB/c IL-4 receptor-deficient mice reveal CD4+Foxp3-negative T cells as a source of IL-10 in Leishmania major susceptibility.
Effect of nasopharyngeal carcinoma-derived exosomes on human regulatory T cells.
VEGFA-VEGFR pathway blockade inhibits tumor-induced regulatory T-cell proliferation in colorectal cancer.
CXCR3+ T regulatory cells selectively accumulate in human ovarian carcinomas to limit type I immunity.
The interplay of effector and regulatory T cells in cancer.
Hypoxia Induces Overexpression of CCL28 to Recruit Treg Cells to Enhance Angiogenesis in Lung Adenocarcinoma.
CD39 is highly involved in mediating the suppression activity of tumor-infiltrating CD8+ T regulatory lymphocytes.
Surgical trauma-induced CCL18 promotes recruitment of regulatory T cells and colon cancer progression.
A CCR4 antagonist combined with vaccines induces antigen-specific CD8+ T cells and tumor immunity against self antigens.
Phosphoinositide 3-kinase $\delta$ is a regulatory T-cell target in cancer immunotherapy.
Aspergillus fumigatus Influences Gasdermin-D-Dependent Pyroptosis of the Lung via Regulating Toll-Like Receptor 2-Mediated Regulatory T Cell Differentiation.
Dendritic cell generation and CD4+ CD25 <sup>high</sup> FOXP3+ regulatory t cells in human head and neck carcinoma during radio-chemotherapy.
Epigenetic quantification of tumor-infiltrating T-lymphocytes.
miR141-CXCL1-CXCR2 signaling-induced Treg recruitment regulates metastases and survival of non-small cell lung cancer.
Melanoma cells express ICOS ligand to promote the activation and expansion of T-regulatory cells.
Adoptive cellular therapy enhances the helper T cell response and reduces the number of regulatory T cells.
Taming regulatory T cells by autologous T cell immunization: a potential new strategy for cancer immune therapy.
MEK inhibition prevents tumour-shed transforming growth factor- $\beta$ -induced T-regulatory cell augmentation in tumour milieu.
Role of plasmacytoid dendritic cells and inducible costimulator-positive regulatory T cells in the immunosuppression microenvironment of gastric cancer.
Functional heterogeneity of circulating T regulatory cell subsets in breast cancer patients.
Anti-TNFR2 enhanced the antitumor activity of a new HMGN1/3M-052 stimulated dendritic cell vaccine in a mouse model of colon cancer.
The role and significance of VEGFR2(+) regulatory T cells in tumor immunity.
MiR-125b-5p modulates the function of regulatory T cells in tumor microenvironment by targeting TNFR2.

The biological role of Treg cells in ectopic endometrium homeostasis.
Increased liver-infiltrating CD8+FoxP3+ regulatory T cells are associated with tumor stage in hepatocellular carcinoma patients.
Enrichment of Foxp3+ CD4 regulatory T cells in migrated T cells to IL-6- and IL-8-expressing tumors through predominant induction of CXCR1 by IL-6.
[Effects of Chinese drugs for activating blood and Chinese drugs for nourishing qi and activating blood on the metastasis of Lewis lung carcinoma in different stages].
Regulatory T cells promote cancer immune-escape through integrin $\alpha\text{v}\beta 8$ -mediated TGF- $\beta$ activation.
Foxp3 expression in pancreatic carcinoma cells as a novel mechanism of immune evasion in cancer.
Helios, CD73 and CD39 Induction in Regulatory T Cells Exposed to Adipose Derived Mesenchymal Stem Cells.
Aiduqing formula inhibits breast cancer metastasis by suppressing TAM/CXCL1-induced Treg differentiation and infiltration.
Regulatory T cells-derived IL-35 promotes the growth of adult acute myeloid leukemia blasts.
Targeted inhibition of IL-10-secreting CD25- Treg via p38 MAPK suppression in cancer immunotherapy.
Computational study for suppression of CD25/IL-2 interaction.
Foxp3 Reprograms T Cell Metabolism to Function in Low-Glucose, High-Lactate Environments.
Identification of CD8+CD25+Foxp3+ suppressive T cells in colorectal cancer tissue.
Classical Hodgkin Lymphoma with Positive Epstein-Barr Virus Status is Associated with More FOXP3 Regulatory T Cells.
Regulatory T cell subsets in human cancer: are they regulating for or against tumor progression?
CCR2 Influences T Regulatory Cell Migration to Tumors and Serves as a Biomarker of Cyclophosphamide Sensitivity.
CD19+CD24 <sup>hi</sup> CD38 <sup>hi</sup> Bregs involved in downregulate helper T cells and upregulate regulatory T cells in gastric cancer.
The use of FoxP3 as a biomarker and prognostic factor for malignant human tumors.
Cyclic adenosine monophosphate involvement in low-dose cyclophosphamide-reversed immune evasion in a mouse lymphoma model.
Histidine decarboxylase (HDC)-expressing granulocytic myeloid cells induce and recruit Foxp3(+) regulatory T cells in murine colon cancer.
Clinical Application of Anti-CCR4 Monoclonal Antibody.
CD39/ENTPD1 expression by CD4+Foxp3+ regulatory T cells promotes hepatic metastatic tumor growth in mice.
The presence of tumor-infiltrating FOXP3+ lymphocytes correlates with intratumoral angiogenesis in endometrial cancer.
TGF- $\beta$ 1 contributes to CD8+ Treg induction through p38 MAPK signaling in ovarian cancer microenvironment.
A low proportion of regulatory T cells before chemoradiotherapy predicts better overall survival in esophageal cancer.
EBV-EBNA1 constructs an immunosuppressive microenvironment for nasopharyngeal carcinoma by promoting the chemoattraction of Treg cells.
A comprehensive review on the role of co-signaling receptors and Treg homeostasis in autoimmunity and tumor immunity.
Circulating regulatory T cell subsets predict overall survival of patients with unresectable pancreatic cancer.

Peripheral blood immune markers in breast cancer: Differences in regulatory T cell abundance are related to clinical parameters.
Role of tumor endothelium in CD4+ CD25+ regulatory T cell infiltration of human pancreatic carcinoma.
Knockdown of HMGB1 in tumor cells attenuates their ability to induce regulatory T cells and uncovers naturally acquired CD8 T cell-dependent antitumor immunity.
Regulatory T cells promote the stemness of leukemia stem cells through IL10 cytokine-related signaling pathway.
Non-platelet-derived CXCL4 differentially regulates cytotoxic and regulatory T cells through CXCR3 to suppress the immune response to colon cancer.
Dopamine inhibits human CD8+ Treg function through D(1)-like dopaminergic receptors.
Circulating FoxP3+ Regulatory T and Interleukin17-Producing Th17 Cells Actively Influence HBV Clearance in De Novo Hepatitis B Virus Infected Patients after Orthotopic Liver Transplantation.
Salidroside regulates tumor microenvironment of non-small cell lung cancer via Hsp70/Stub1/Foxp3 pathway in Tregs.
Gastric Cancer Stem Cells Effect on Th17/Treg Balance; A Bench to Beside Perspective.
Mycobacteria-responsive sonic hedgehog signaling mediates programmed death-ligand 1- and prostaglandin E2-induced regulatory T cell expansion.
CXCL12 mediates immunosuppression in the lymphoma microenvironment after allogeneic transplantation of hematopoietic cells.
The dark side of cyclophosphamide: cyclophosphamide-mediated ablation of regulatory T cells.
Regulatory T cells: potential target in anticancer immunotherapy.
LncRNA SNHG1 regulates the differentiation of Treg cells and affects the immune escape of breast cancer via regulating miR-448/IDO.
CD69+ CD4+ CD25- T cells, a new subset of regulatory T cells, suppress T cell proliferation through membrane-bound TGF-beta 1.
Expression of the Epstein-Barr virus-encoded Epstein-Barr virus nuclear antigen 1 in Hodgkin's lymphoma cells mediates Up-regulation of CCL20 and the migration of regulatory T cells.
Local accumulation of FOXP3+ regulatory T cells: evidence for an immune evasion mechanism in patients with large condylomata acuminata.
The dendritic cell-regulatory T lymphocyte crosstalk contributes to tumor-induced tolerance.
Active site-targeted covalent irreversible inhibitors of USP7 impair the functions of Foxp3+ T-regulatory cells by promoting ubiquitination of Tip60.
B7-H1 and B7-H4 expression in colorectal carcinoma: correlation with tumor FOXP3(+) regulatory T-cell infiltration.
Tnfrsf4-expressing regulatory T cells promote immune escape of chronic myeloid leukemia stem cells.
Activin A Promotes Regulatory T-cell-Mediated Immunosuppression in Irradiated Breast Cancer.
Tumor microenvironmental modification by the current target therapy for head and neck squamous cell carcinoma.
ICOS-ligand expression on plasmacytoid dendritic cells supports breast cancer progression by promoting the accumulation of immunosuppressive CD4+ T cells.
Lenalidomide-based maintenance therapy reduces TNF receptor 2 on CD4 T cells and enhances immune effector function in acute myeloid leukemia patients.
Cancer cell-derived IL-1 $\alpha$ induces CCL22 and the recruitment of regulatory T cells.
Foxp3 expression in human cancer cells.
Human prostate tumor antigen-specific CD8+ regulatory T cells are inhibited by CTLA-4 or IL-35 blockade.

Tumor sialylation impedes T cell mediated anti-tumor responses while promoting tumor associated-regulatory T cells.
FKBP51s signature in peripheral blood mononuclear cells of melanoma patients as a possible predictive factor for immunotherapy.
Pembrolizumab Interferes with the Differentiation of Human FOXP3(+)-Induced T Regulatory Cells, but Not with FOXP3 Stability, through Activation of mTOR.
TNFR2: Role in Cancer Immunology and Immunotherapy.
Effector memory regulatory T-cell expansion marks a pivotal point of immune escape in myelodysplastic syndromes.
Characterization of tumor-infiltrating immune cells in relation to microbiota in colorectal cancers.
Regulatory T cells are redirected to kill glioblastoma by an EGFRvIII-targeted bispecific antibody.
Expression of B7-H3, a potential factor of tumor immune evasion in combination with the number of regulatory T cells, affects against recurrence-free survival in breast cancer patients.
Arsenic trioxide is an immune adjuvant in liver cancer treatment.
Targeting cytokines secreted by CD4(+) CD25(high) CD127(low) regulatory T cells inhibits ovarian cancer progression.
CCR 20th Anniversary Commentary: From Regulatory T Cells to Checkpoint Monoclonal Antibodies--Immuno-oncology Advances Clinical Cancer Research.
Tumor antigen specific iTreg accumulate in the tumor microenvironment and suppress therapeutic vaccination.
Comparison of immunosuppressive and immunomodulatory cells in keratoacanthoma and cutaneous squamous cell carcinoma.
Activated Galectin-9/Tim3 promotes Treg and suppresses Th1 effector function in chronic lymphocytic leukemia.
Increased expression of the regulatory T cell-associated marker CTLA-4 in bovine leukemia virus infection.
An immune-related gene prognostic index for acute myeloid leukemia associated with regulatory T cells infiltration.
Complex Immune Evasion Strategies in Classical Hodgkin Lymphoma.
Investigating the Role of Indoleamine 2,3-Dioxygenase in Acute Myeloid Leukemia: A Systematic Review.
Chemoresistance in mesenchymal lung cancer cells is correlated to high regulatory T cell presence in the tumor microenvironment.
GDF15 induces immunosuppression via CD48 on regulatory T cells in hepatocellular carcinoma.
Matrix metalloproteinase 12 expression is associated with tumor FOXP3(+) regulatory T cell infiltration and poor prognosis in hepatocellular carcinoma.
Expression of Forkhead box P3 in tumour cells causes immunoregulatory function of signet ring cell carcinoma of the stomach.
Inhibition of EphB4-Ephrin-B2 Signaling Reprograms the Tumor Immune Microenvironment in Head and Neck Cancers.
Ganoderic acid Me induces the apoptosis of competent T cells and increases the proportion of Treg cells through enhancing the expression and activation of indoleamine 2,3-dioxygenase in mouse lewis lung cancer cells.
Accumulation and suppressive function of regulatory T cells in malignant ascites: Reducing their suppressive function using arsenic trioxide in vitro.
Regulatory T cells do not suppress rather activate human basophils by IL-3 and STAT5-dependent mechanisms.

Inverse correlation between NKG2D expression on CD8+ T cells and the frequency of CD4+CD25+ regulatory T cells in patients with esophageal cancer.
ST2 and regulatory T cells in the colorectal adenoma/carcinoma microenvironment: implications for diseases progression and prognosis.
Neoadjuvant Chemotherapy Reinforces Antitumour T cell Response in Urothelial Urinary Bladder Cancer.
Foxp3 overexpression decreases sensitivity to chemotherapy in mouse Lewis lung cancer cells.
Targeting regulatory T cells in cytokine-induced killer cell cultures (Review).
Analysis of Treg cell population in patients with breast cancer with respect to progesterone receptor status.
PD-L1 is a critical mediator of regulatory B cells and T cells in invasive breast cancer.
Regulatory T Cells and Plasmacytoid Dendritic Cells Within the Tumor Microenvironment in Gastric Cancer Are Correlated With Gastric Microbiota Dysbiosis: A Preliminary Study.
A link between PDL1 and T regulatory cells in fetomaternal tolerance.
HDAC5 controls the functions of Foxp3(+) T-regulatory and CD8(+) T cells.
MicroRNA-15a/16-1 Prevents Hepatocellular Carcinoma by Disrupting the Communication Between Kupffer Cells and Regulatory T Cells.
Galectin-1 fosters an immunosuppressive microenvironment in colorectal cancer by reprogramming CD8(+) regulatory T cells.
Advances in Human Immune System Mouse Models for Personalized Treg-Based Immunotherapies.
Increase in tumour-infiltrating lymphocytes with regulatory T cell immunophenotypes and reduced zeta-chain expression in nasopharyngeal carcinoma patients.
Standardized approach for evaluating tumor infiltrating lymphocytes in canine mammary carcinoma: Spatial distribution and score as relevant features of tumor malignancy.
Tumor infiltrating regulatory T cells: tractable targets for immunotherapy.
TNFR2 expression predicts the responses to immune checkpoint inhibitor treatments.
Lysosomal Acid Lipase Deficiency Controls T- and B-Regulatory Cell Homeostasis in the Lymph Nodes of Mice with Human Cancer Xenotransplants.
Nodal metastasis in cervical cancer occurs in clearly delineated fields of immune suppression in the pelvic lymph catchment area.
Impact of Allium sativum ethanol extract on immuno-regulatory T cells and anti-inflammatory cytokine profile in murine schistosomiasis.
The role of cytokine signaling in the pathogenesis of cutaneous T-cell lymphoma.
Clinical implications and characteristics of factor forkhead box protein 3 in gastric cancer.
Effect of cytotoxic T-lymphocyte-associated protein 4 on CD4(+)CD25(+) regulatory T cells in murine schistosomiasis japonica.
Monocyte-derived dendritic cells from breast cancer patients are biased to induce CD4+CD25+Foxp3+ regulatory T cells.
TIGIT enhances CD4(+) regulatory T-cell response and mediates immune suppression in a murine ovarian cancer model.
B7.1/NHS76: a new costimulator fusion protein for the immunotherapy of solid tumors.
Overexpression of the transcription factor FOXP3 in lung adenocarcinoma sustains malignant character by promoting G1/S transition gene CCND1.
The immunosuppressive enzyme IL4I1 promotes FoxP3(+) regulatory T lymphocyte differentiation.
Unique roles of Schistosoma japonicum protein Sj16 to induce IFN-γ and IL-10 producing CD4(+)CD25(+) regulatory T cells in vitro and in vivo.
Immuno-regulatory malignant B cells contribute to Chronic Lymphocytic Leukemia progression.



Kindlin-1 regulates IL-6 secretion and modulates the immune environment in breast cancer models.
Regulatory T-cells and micrometastasis in lymph nodes of stage I NSCLC.
Enhanced autoimmunity associated with induction of tumor immunity in thyroiditis-susceptible mice.
Kras(G12D) mutation contributes to regulatory T cell conversion through activation of the MEK/ERK pathway in pancreatic cancer.
Immunopathogenesis of lymphoma: focus on CCR4.
Induction of liver fibrosis in a murine hepatoma model by thioacetamide is associated with enhanced tumor growth and suppressed antitumor immunity.
miR-22 promotes immunosuppression via activating JAK/STAT3 signaling in cutaneous squamous cell carcinoma.
Downregulation of miR-4772-3p promotes enhanced regulatory T cell capacity in malignant pleural effusion by elevating Helios levels.
A key role of GARP in the immune suppressive tumor microenvironment.
Astragaloside IV inhibits progression of lung cancer by mediating immune function of Tregs and CTLs by interfering with IDO.
Up-regulated S100 calcium binding protein A8 in Plasmodium-infected patients correlates with CD4(+)CD25(+)Foxp3 regulatory T cell generation.
Blockade of $\beta$ -Catenin-Induced CCL28 Suppresses Gastric Cancer Progression via Inhibition of Treg Cell Infiltration.
Discovery of secondary sulphonamides as IDO1 inhibitors with potent antitumour effects in vivo.
Targeting MARCO and IL37R on Immunosuppressive Macrophages in Lung Cancer Blocks Regulatory T Cells and Supports Cytotoxic Lymphocyte Function.
Hypothermic microenvironment plays a key role in tumor immune subversion.
Nocardia brasiliensis induces an immunosuppressive microenvironment that favors chronic infection in BALB/c mice.
Autocrine Adenosine Regulates Tumor Polyfunctional CD73(+)CD4(+) Effector T Cells Devoid of Immune Checkpoints.
KDM3A promotes inhibitory cytokines secretion by participating in TLR4 regulation of Foxp3 transcription in lung adenocarcinoma cells.
Long Non-coding RNA FENDRR Acts as a miR-423-5p Sponge to Suppress the Treg-Mediated Immune Escape of Hepatocellular Carcinoma Cells.
Circulating regulatory T cells from breast cancer patients in response to neoadjuvant chemotherapy.
TGF- $\beta$ -miR-34a-CCL22 signaling-induced Treg cell recruitment promotes venous metastases of HBV-positive hepatocellular carcinoma.
Cross talk between follicular Th cells and tumor cells in human follicular lymphoma promotes immune evasion in the tumor microenvironment.
CCR4 as a novel molecular target for immunotherapy of cancer.
Human epithelial ovarian carcinoma cell-derived cytokines cooperatively induce activated CD4+CD25-CD45RA+ naïve T cells to express forkhead box protein 3 and exhibit suppressive ability in vitro.
A structurally distinct TGF- $\beta$ mimic from an intestinal helminth parasite potently induces regulatory T cells.
Regulation of hematopoietic and leukemia stem cells by regulatory T cells.
Depletion of endogenous tumor-associated regulatory T cells improves the efficacy of adoptive cytotoxic T-cell immunotherapy in murine acute myeloid leukemia.
Expression of costimulatory molecules B7-H1, B7-H4 and Foxp3+ Tregs in gastric cancer and its clinical significance.

Humanization of an anti-CCR4 antibody that kills cutaneous T-cell lymphoma cells and abrogates suppression by T-regulatory cells.
[Paclitaxel blocks immunologic escape through up-regulating TAP-1, TAP-2 and eliminatiing Treg cells in 3LL-bearing mice].
Mass cytometry of Hodgkin lymphoma reveals a CD4(+) regulatory T-cell-rich and exhausted T-effector microenvironment.
Peripheral Tr1 and Foxp3(+) Treg as Markers of Recurrent Malignancies in Patients with Hepato-Biliary Pancreatic Cancers.
A Brief Communication on Circulating PD-1-positive T-Regulatory Lymphocytes in Melanoma Patients Undergoing Adjuvant Immunotherapy With Nivolumab.
Loss of histone macroH2A1 in hepatocellular carcinoma cells promotes paracrine-mediated chemoresistance and CD4(+)CD25(+)FoxP3(+) regulatory T cells activation.
Downregulation of FOXP3 inhibits invasion and immune escape in cholangiocarcinoma.
The transcription factor Forkhead box P3 (FoxP3) is expressed in glioma cells and associated with increased apoptosis.
Adenovirus-mediated intratumoral expression of immunostimulatory proteins in combination with systemic Treg inactivation induces tumor-destructive immune responses in mouse models.
Early lung carcinogenesis and tumor microenvironment observed by single-cell transcriptome analysis.
[Allergo-oncology: what allergologists and oncologists can learn from each other : Regulatory T cells in allergy and cancer].
Targeting regulatory T cells by curcumin: A potential for cancer immunotherapy.
[The change of CD4+ CD25high CCR6+ regulatory T cells in breast cancer patients].
[Inhibitory Effect of Feiji Recipe on IDO Induced Immune Escape on the Murine Model of Lewis Lung Carcinoma].
Engineered PD-1/TIGIT dual-activating cell-membrane nanoparticles with dexamethasone act synergistically to shape the effector T cell/Treg balance and alleviate systemic lupus erythematosus.
Foxp3 expression in A549 cells is regulated by Toll-like receptor 4 through nuclear factor- $\kappa$ B.
The number of regular T cells and immature dendritic cells involved in mycosis fungoides is linked to the tumor stage.
Dendritic cell-derived TGF- $\beta$ mediates the induction of mucosal regulatory T-cell response to Helicobacter infection essential for maintenance of immune tolerance in mice.
Discovery and characterization of a selective IKZF2 glue degrader for cancer immunotherapy.
Prognostic value of tumor-infiltrating Foxp3+ T-cell subpopulations in metastatic melanoma.
[Research progress of lymphocyte activation gene 3 (LAG-3) in tumor immunotherapy and parasite disease].
The long noncoding RNA Inc-EGFR stimulates T-regulatory cells differentiation thus promoting hepatocellular carcinoma immune evasion.
Clinical value of Pro-GRP and T lymphocyte subpopulation for the assessment of immune functions of lung cancer patients after DC-CIK biological therapy.
Immunomodulatory Effects of Lenvatinib Plus Anti-Programmed Cell Death Protein 1 in Mice and Rationale for Patient Enrichment in Hepatocellular Carcinoma.
Effector Antitumor and Regulatory T Cell Responses Influence the Development of Nonmelanoma Skin Cancer in Kidney Transplant Patients.
Regulatory T-cell phenotype in association with large cell transformation of mycosis fungoides.
Th17/Treg imbalance in patients with liver cystic echinococcosis.

High and interrelated rates of PD-L1+CD14+ antigen-presenting cells and regulatory T cells mark the microenvironment of metastatic lymph nodes from patients with cervical cancer.
Host immune responses to cervical cancer.
Detection of Tax-specific CTLs in lymph nodes of adult T-cell leukemia/lymphoma patients and its association with Foxp3 positivity of regulatory T-cell function.
The Effect of Antibody Fragments on CD25 Targeted Regulatory T Cell Near-Infrared Photoimmunotherapy.
A Lectin-EGF antibody promotes regulatory T cells and attenuates nephrotoxic nephritis via DC-SIGN on dendritic cells.
Oxygen Sensing by T Cells Establishes an Immunologically Tolerant Metastatic Niche.
Influence of Tumor Location on the Composition of Immune Infiltrate and Its Impact on Patient Survival. Lessons from DCBCL and Animal Models.
CD73-deficient mice have increased antitumor immunity and are resistant to experimental metastasis.
Humoral immune response to adenovirus induce tolerogenic bystander dendritic cells that promote generation of regulatory T cells.
Anti-glucocorticoid-induced Tumor Necrosis Factor-Related Protein (GITR) Therapy Overcomes Radiation-Induced Treg Immunosuppression and Drives Abscopal Effects.
L1CAM overexpression promotes tumor progression through recruitment of regulatory T cells in esophageal carcinoma.
Cancer cell plasticity and MHC-II-mediated immune tolerance promote breast cancer metastasis to lymph nodes.
Immunomodulation for inhibitors in hemophilia A: the important role of Treg cells.
Enhancement of regulatory T cell-like suppressive function in MT-2 by long-term and low-dose exposure to asbestos.
N6-methyladenosine-modified circular RNA QSOX1 promotes colorectal cancer resistance to anti-CTLA-4 therapy through induction of intratumoral regulatory T cells.
Integrated Genomic and Immunophenotypic Classification of Pancreatic Cancer Reveals Three Distinct Subtypes with Prognostic/Predictive Significance.
Immunohistochemical evaluation of immune cell infiltration in canine gliomas.
Nuclear FAK controls chemokine transcription, Tregs, and evasion of anti-tumor immunity.
The parasite cytokine mimic Hp-TGM potently replicates the regulatory effects of TGF- $\beta$ on murine CD4(+) T cells.
Transcriptional regulation of Foxp3 gene: multiple signal pathways on the road.
The Dynamics of Treg/Th17 and the Imbalance of Treg/Th17 in Clonorchis sinensis-Infected Mice.
Deciphering the influence of urinary microbiota on FoxP3+ regulatory T cell infiltration and prognosis in Chinese patients with non-muscle-invasive bladder cancer.
OX40 as a novel target for the reversal of immune escape in colorectal cancer.
Presence of periodontitis may synergistically contribute to cancer progression via Treg and IL-6.
EBV+ tumors exploit tumor cell-intrinsic and -extrinsic mechanisms to produce regulatory T cell-recruiting chemokines CCL17 and CCL22.
Increased infiltration of regulatory T cells in hepatocellular carcinoma of patients with hepatitis B virus pre-S2 mutant.
Dihydroartemisinin inhibits melanoma by regulating CTL/Treg anti-tumor immunity and STAT3-mediated apoptosis via IL-10 dependent manner.
Low doses of cholera toxin and its mediator cAMP induce CTLA-2 secretion by dendritic cells to enhance regulatory T cell conversion.

Synergistic efficacy of PI3K $\delta$ inhibitor with anti-PD-1 mAbs in immune-humanized PDX model of endocrine resistance hormone receptor-positive advanced breast cancer.
Immune Evasion and Drug Resistance Mediated by USP22 in Cancer: Novel Targets and Mechanisms.
Molecular Mechanisms for cAMP-Mediated Immunoregulation in T cells - Role of Anchored Protein Kinase A Signaling Units.
Regulatory T cells mediate maternal tolerance to the fetus.
Induction of immune response against NY-ESO-1 by CHP-NY-ESO-1 vaccination and immune regulation in a melanoma patient.
GARP as an Immune Regulatory Molecule in the Tumor Microenvironment of Glioblastoma Multiforme.
Targeting multiple-myeloma-induced immune dysfunction to improve immunotherapy outcomes.
Investigations on the clinical significance of FOXP3 protein expression in cervical oesophageal cancer and the number of FOXP3+ tumour-infiltrating lymphocytes.
Immunomodulatory effects of radiofrequency ablation in a breast cancer model.
Thalidomide regulation of NF- $\kappa$ B proteins limits Tregs activity in chronic lymphocytic leukemia.
Canine Transforming Growth Factor- $\beta$ Receptor 2-Ig: A Potential Candidate Biologic for Melanoma Treatment That Reverses Transforming Growth Factor- $\beta$ 1 Immunosuppression.
Clinical Significance of Fusobacterium nucleatum Infection and Regulatory T Cell Enrichment in Esophageal Squamous Cell Carcinoma.
Bone marrow infiltrated Lnc-INSR induced suppressive immune microenvironment in pediatric acute lymphoblastic leukemia.
Long non-coding RNA MEG3 mediates the miR-149-3p/FOXP3 axis by reducing p53 ubiquitination to exert a suppressive effect on regulatory T cell differentiation and immune escape in esophageal cancer.
Revealing the evolution of the tumor immune microenvironment in follicular lymphoma patients progressing within 24 months using single-cell imaging mass cytometry.
Mouse Sertoli cells display phenotypical and functional traits of antigen-presenting cells in response to interferon gamma.
Immunohistochemical pattern of T lymphocytes population within bilharzial-associated bladder neoplasm microenvironment.
The AP1-dependent secretion of galectin-1 by Reed Sternberg cells fosters immune privilege in classical Hodgkin lymphoma.
Quantitative analysis of infiltrating immune cells and bovine papillomavirus type 1 E2-positive cells in equine sarcoids.
Immune-Stimulatory Effects of Curcumin on the Tumor Microenvironment in Head and Neck Squamous Cell Carcinoma.
Japanese encephalitis virus expands regulatory T cells by increasing the expression of PD-L1 on dendritic cells.
The expression of Foxp3 and TLR4 in cervical cancer: association with immune escape and clinical pathology.
IDO1-Targeted Therapy Does Not Control Disease Development in the E $\mu$ -TCL1 Mouse Model of Chronic Lymphocytic Leukemia.
Methotrexate up-regulates ecto-5'-nucleotidase/CD73 and reduces the frequency of T lymphocytes in the glioblastoma microenvironment.
The changes of angiogenesis and immune regulations in stromal microenvironment of cutaneous melanomas.

The Influence of the Genetic and Immunologic Context in the Development of Colorectal Adenoma: A Case Series Report.
Association of Foxp3 rs3761548 polymorphism with cytokines concentration in gastric adenocarcinoma patients.
5-Hydroxytryptamine1a receptors on tumour cells induce immune evasion in lung adenocarcinoma patients with depression via autophagy/pSTAT3.
Downregulation of RCAS1 and upregulation of cytotoxic T cells affects synovial proliferation and apoptosis in rheumatoid arthritis.
Membrane-organizing protein moesin controls Treg differentiation and antitumor immunity via TGF- $\beta$ signaling.
ARGX-110, a highly potent antibody targeting CD70, eliminates tumors via both enhanced ADCC and immune checkpoint blockade.
Autoimmune Hepatitis: Tolerogenic Immunological State During Pregnancy and Immune Escape in Post-partum.
Regulatory T Cell Modulation by CBP/EP300 Bromodomain Inhibition.
ICOS-L as a Potential Therapeutic Target for Cancer Immunotherapy.
Elevated IL-35 in bone marrow of the patients with acute myeloid leukemia.
The Effects of Intestinal Nematode L4 Stage on Mouse Experimental Autoimmune Encephalomyelitis.
High density of FOXP3-positive T cells infiltrating colorectal cancers with microsatellite instability.
Endothelin B receptor expression in malignant gliomas: the perivascular immune escape mechanism of gliomas.
Synergistic tumoricidal effect of combined hPD-L1 vaccine and HER2 gene vaccine.
Expression of CXC chemokine receptor-4 and forkhead box 3 in neuroblastoma cells and response to chemotherapy.
Spatial interplay of tissue hypoxia and T-cell regulation in ductal carcinoma in situ.
Pathologic significance of immunoglobulin G4-positive plasma cells in extrahepatic cholangiocarcinoma.
Hashimoto's thyroiditis attenuates progression of papillary thyroid carcinoma: deciphering immunological links.
The GABA and GABA-Receptor System in Inflammation, Anti-Tumor Immune Responses, and COVID-19.
CircUBAP2-mediated competing endogenous RNA network modulates tumorigenesis in pancreatic adenocarcinoma.
RETRACTED ARTICLE: Down-regulation of lncRNA FEZF1-AS1 mediates regulatory T cell differentiation and further blocks immune escape in colon cancer.
Simultaneous Increase in Serum Levels of IL-37 and IL-18 Binding Protein In Low-Grade and High-Grade Brain Tumors.

Table A2-27, Cluster 26

Cluster 26 focuses on human cytomegalovirus disease, with emphasis on countering HCMV immune system evasion and immune escape (309)
[Current status of human cytomegalovirus disease].
The Human Cytomegalovirus Protein UL148A Downregulates the NK Cell-Activating Ligand MICA To Avoid NK Cell Attack.

The expression of interleukin-32 is activated by human cytomegalovirus infection and down regulated by hcmv-miR-UL112-1.
Human Cytomegalovirus (HCMV)-Specific CD4(+) T Cells Are Polyfunctional and Can Respond to HCMV-Infected Dendritic Cells In Vitro.
Dendritic cell biology in human cytomegalovirus infection and the clinical consequences for host immunity and pathology.
Insight for Immunotherapy of HCMV Infection.
Natural killer cells can inhibit the transmission of human cytomegalovirus in cell culture by using mechanisms from innate and adaptive immune responses.
Viral regulation of RANTES expression during human cytomegalovirus infection of endothelial cells.
Battle between Host Immune Cellular Responses and HCMV Immune Evasion.
Subversion of Immune Response by Human Cytomegalovirus.
[Cytomegalovirus (CMV)].
Recent Approaches and Strategies in the Generation of Anti-human Cytomegalovirus Vaccines.
Host Genetics of Cytomegalovirus Pathogenesis.
The use of microRNA by human viruses: lessons from NK cells and HCMV infection.
Development of a Human Cytomegalovirus (HCMV)-Based Therapeutic Cancer Vaccine Uncovers a Previously Unsuspected Viral Block of MHC Class I Antigen Presentation.
Suppression of costimulation by human cytomegalovirus promotes evasion of cellular immune defenses.
Diverse specificities, phenotypes, and antiviral activities of cytomegalovirus-specific CD8+ T cells.
Human cytomegalovirus impairs dendritic cell function: a novel mechanism of human cytomegalovirus immune escape.
What we have learned from animal models of HCMV.
[Research Progress in Mechanisms Associated with Latent Infection in Humans by the Cytomegalovirus].
Genetic Variability of Human Cytomegalovirus Clinical Isolates Correlates With Altered Expression of Natural Killer Cell-Activating Ligands and IFN- $\gamma$ .
Comprehensive Analysis of Human Cytomegalovirus- and HIV-Mediated Plasma Membrane Remodeling in Macrophages.
A Prominent Role of the Human Cytomegalovirus UL8 Glycoprotein in Restraining Proinflammatory Cytokine Production by Myeloid Cells at Late Times during Infection.
Human cytomegalovirus persistence.
Strategy of Human Cytomegalovirus To Escape Interferon Beta-Induced APOBEC3G Editing Activity.
CARs-A New Perspective to HCMV Treatment.
MicroRNA-182 inhibits HCMV replication through activation of type I IFN response by targeting FOXO3 in neural cells.
Development of a Vaccine against Human Cytomegalovirus: Advances, Barriers, and Implications for the Clinical Practice.
microRNA, a Subtle Indicator of Human Cytomegalovirus against Host Immune Cells.
Human cytomegalovirus encoded homologs of cytokines, chemokines and their receptors: roles in immunomodulation.
Cytomegalovirus and the aging population.
Immune evasion proteins gpUS2 and gpUS11 of human cytomegalovirus incompletely protect infected cells from CD8 T cell recognition.
Human cytomegalovirus suppresses Fas expression and function.

Human Cytomegalovirus Encoded miR-US25-1-5p Attenuates CD147/EMMPRIN-Mediated Early Antiviral Response.
Severe Symptomatic Primary Human Cytomegalovirus Infection despite Effective Innate and Adaptive Immune Responses.
Altered expression of host-encoded complement regulators on human cytomegalovirus-infected cells. Cytomegaloviruses use multiple mechanisms to elude the host immune response.
Human Cytomegalovirus miR-US33as-5p Targets IFNAR1 to Achieve Immune Evasion During Both Lytic and Latent Infection.
Cytokine-Mediated Induction and Regulation of Tissue Damage During Cytomegalovirus Infection.
Tuning the Orchestra: HCMV vs. Innate Immunity.
Modulation of HLA expression in human cytomegalovirus immune evasion.
Diverse immune evasion strategies by human cytomegalovirus.
Functional Profile of Human Cytomegalovirus Genes and Their Associated Diseases: A Review.
Cytomegalovirus-Infected Cells Resist T Cell Mediated Killing in an HLA-Recognition Independent Manner.
Human cytomegalovirus modulates monocyte-mediated innate immune responses during short-term experimental latency in vitro.
Impact of donor and recipient human cytomegalovirus status on kidney transplantation.
Targeting the function of mature dendritic cells by human cytomegalovirus: a multilayered viral defense strategy.
Dual inhibition of innate immunity and apoptosis by human cytomegalovirus protein UL37x1 enables efficient virus replication.
Differential expression of chemokines by human retinal pigment epithelial cells infected with cytomegalovirus.
Evaluation of the host transcriptional response to human cytomegalovirus infection.
Human cytomegalovirus encoded microRNAs: hitting targets.
Human Cytomegalovirus Tegument Protein UL82 Inhibits STING-Mediated Signaling to Evade Antiviral Immunity.
Human cytomegalovirus induces and exploits Roquin to counteract the IRF1-mediated antiviral state.
Human Cytomegalovirus DNA Polymerase Subunit UL44 Antagonizes Antiviral Immune Responses by Suppressing IRF3- and NF-κB-Mediated Transcription.
Human cytomegalovirus gene expression in long-term infected glioma stem cells.
Chemokine sequestration by viral chemoreceptors as a novel viral escape strategy: withdrawal of chemokines from the environment of cytomegalovirus-infected cells.
CMV immune evasion and manipulation of the immune system with aging.
Modulation of dendritic cell functions by viral IL-10 encoded by human cytomegalovirus.
Turning the tables on cytomegalovirus: targeting viral Fc receptors by CARs containing mutated CH2-CH3 IgG spacer domains.
Human cytomegalovirus inhibits cytokine-induced macrophage differentiation.
High-Definition Analysis of Host Protein Stability during Human Cytomegalovirus Infection Reveals Antiviral Factors and Viral Evasion Mechanisms.
The human cytomegalovirus protein UL147A downregulates the most prevalent MICA allele: MICA*008, to evade NK cell-mediated killing.
Trehalose, an mTOR-Independent Inducer of Autophagy, Inhibits Human Cytomegalovirus Infection in Multiple Cell Types.
UL40 human cytomegalovirus variability evolution patterns over time in renal transplant recipients.

Identification of a Genetic Variation in ERAP1 Aminopeptidase that Prevents Human Cytomegalovirus miR-UL112-5p-Mediated Immune Evasion.
The CD94/NKG2C+ NK-cell subset on the edge of innate and adaptive immunity to human cytomegalovirus infection.
Human cytomegalovirus tegument proteins (pp65, pp71, pp150, pp28).
All is fair in virus-host interactions: NK cells and cytomegalovirus.
Selective intracellular retention of virally induced NKG2D ligands by the human cytomegalovirus UL16 glycoprotein.
[Modulation of host immune defenses by cytomegalovirus: advanced insights from evolutionary game theory].
Human cytomegalovirus impairs the function of plasmacytoid dendritic cells in lymphoid organs.
The latency-associated UL138 gene product of human cytomegalovirus sensitizes cells to tumor necrosis factor alpha (TNF-alpha) signaling by upregulating TNF-alpha receptor 1 cell surface expression.
Human Cytomegalovirus Immediate Early 86-kDa Protein Blocks Transcription and Induces Degradation of the Immature Interleukin-1 $\beta$ Protein during Virion-Mediated Activation of the AIM2 Inflammasome.
Human cytomegalovirus-infected cells have unstable assembly of major histocompatibility complex class I complexes and are resistant to lysis by cytotoxic T lymphocytes.
Human Cytomegalovirus-Induced Degradation of CYTIP Modulates Dendritic Cell Adhesion and Migration.
Reactive oxygen species-induced parthanatos of immunocytes by human cytomegalovirus-associated substance.
Human Cytomegalovirus Protein UL94 Targets MTA to Evade the Antiviral Immune Response.
Neutralization of rhesus cytomegalovirus IL-10 reduces horizontal transmission and alters long-term immunity.
Expression dynamics of human cytomegalovirus immune evasion genes US3, US6, and US11 in the blood of lung transplant recipients.
The interplay between host and viral factors in shaping the outcome of cytomegalovirus infection.
Rhesus cytomegalovirus contains functional homologues of US2, US3, US6, and US11.
Human cytomegalovirus immunity and immune evasion.
Immune evasion by human cytomegalovirus: lessons in immunology and cell biology.
US28 actions in HCMV infection: lessons from a versatile hijacker.
The role of cell types in cytomegalovirus infection in vivo.
Checks and balances between human cytomegalovirus replication and indoleamine-2,3-dioxygenase.
Frequent detection of human cytomegalovirus in neuroblastoma: a novel therapeutic target?
Regulation of Latency and Reactivation by Human Cytomegalovirus miRNAs.
Cytomegalovirus: the role of CMV post-haematopoietic stem cell transplantation.
Biological relevance of Cytomegalovirus genetic variability in congenitally and postnatally infected children.
Who's Driving? Human Cytomegalovirus, Interferon, and NF $\kappa$ B Signaling.
The US21 viroporin of human cytomegalovirus stimulates cell migration and adhesion.
Hcmv-miR-UL112 attenuates NK cell activity by inhibition type I interferon secretion.
Unravelling the interaction of human cytomegalovirus with dendritic cells by using SuperSAGE.
Large-Scale Screening of HCMV-Seropositive Blood Donors Indicates that HCMV Effectively Escapes from Antibodies by Cell-Associated Spread.



Human Cytomegalovirus UL24 and UL43 Cooperate to Modulate the Expression of Immunoregulatory UL16 Binding Protein 1.
Human Cytomegalovirus MicroRNAs miR-US5-1 and miR-UL112-3p Block Proinflammatory Cytokine Production in Response to NF- $\kappa$ B-Activating Factors through Direct Downregulation of IKK $\alpha$ and IKK $\beta$ .
pUL34 binding near the human cytomegalovirus origin of lytic replication enhances DNA replication and viral growth.
The Abundant Tegument Protein pUL25 of Human Cytomegalovirus Prevents Proteasomal Degradation of pUL26 and Supports Its Suppression of ISGylation.
Increase in the expression of human leukocyte antigen class I in human fibroblasts by soluble factors secreted from human cytomegalovirus-infected cells.
The Role of HCMV and HIV-1 MicroRNAs: Processing, and Mechanisms of Action during Viral Infection.
HCMV-encoded US7 and US8 act as antagonists of innate immunity by distinctively targeting TLR-signaling pathways.
The Viral Tegument Protein pp65 Impairs Transcriptional Upregulation of IL-1 $\beta$ by Human Cytomegalovirus through Inhibition of NF- $\kappa$ B Activity.
Cytomegalovirus US2 destroys two components of the MHC class II pathway, preventing recognition by CD4+ T cells.
Rhesus cytomegalovirus encodes seventeen microRNAs that are differentially expressed in vitro and in vivo.
Human cytomegalovirus protein RL1 degrades the antiviral factor SLFN11 via recruitment of the CRL4 E3 ubiquitin ligase complex.
Supernatants from human cytomegalovirus (HCMV)-infected retinal glial cells increase transepithelial electrical resistance in a cell culture model: evidence of HCMV immune escape in the eye?
Modulation of the NF $\kappa$ B Signalling Pathway by Human Cytomegalovirus.
Mouse Model of Cytomegalovirus Disease and Immunotherapy in the Immunocompromised Host: Predictions for Medical Translation that Survived the "Test of Time".
Cross-presentation of human cytomegalovirus pp65 (UL83) to CD8+ T cells is regulated by virus-induced, soluble-mediator-dependent maturation of dendritic cells.
Rapid Degradation Pathways of Host Proteins During HCMV Infection Revealed by Quantitative Proteomics.
Quantitative temporal viromics: an approach to investigate host-pathogen interaction.
Islands of linkage in an ocean of pervasive recombination reveals two-speed evolution of human cytomegalovirus genomes.
Roseoloviruses and their modulation of host defenses.
Human cytomegalovirus chemokine receptor gene US28 is transcribed in latently infected THP-1 monocytes.
Multiple independent loci within the human cytomegalovirus unique short region down-regulate expression of major histocompatibility complex class I heavy chains.
Degradation of RIG-I following cytomegalovirus infection is independent of apoptosis.
The human cytomegalovirus-encoded chemokine receptor US28 induces caspase-dependent apoptosis.
The pathogenicity of cytomegalovirus.
The Interplay between Human Cytomegalovirus and Pathogen Recognition Receptor Signaling.
Sleepless latency of human cytomegalovirus.
Cytomegalovirus: virological facts for clinicians.
The tiers and dimensions of evasion of the type I interferon response by human cytomegalovirus.

Expression of the UL16 glycoprotein of Human Cytomegalovirus protects the virus-infected cell from attack by natural killer cells.
Human Cytomegalovirus UL23 Antagonizes the Antiviral Effect of Interferon- $\gamma$ by Restraining the Expression of Specific IFN-Stimulated Genes.
Human cytomegalovirus interactome analysis identifies degradation hubs, domain associations and viral protein functions.
The pentameric complex drives immunologically covert cell-cell transmission of wild-type human cytomegalovirus.
Human Cytomegalovirus Protein UL31 Inhibits DNA Sensing of cGAS to Mediate Immune Evasion.
Human cytomegalovirus glycoproteins.
The human cytomegalovirus TRL11/IRL11-encoded Fc $\gamma$ R binds differentially to allelic variants of immunoglobulin G1.
Impact of Natural Occurring ERAP1 Single Nucleotide Polymorphisms within miRNA-Binding Sites on HCMV Infection.
Human cytomegalovirus IE2 protein regulates macrophage-mediated immune escape by upregulating GRB2 expression in UL122 genetically modified mice.
Latency-associated viral interleukin-10 (IL-10) encoded by human cytomegalovirus modulates cellular IL-10 and CCL8 Secretion during latent infection through changes in the cellular microRNA hsa-miR-92a.
Impaired killing of HCMV-infected retinal pigment epithelial cells by anti-pp65 CD8(+) cytotoxic T cells.
Current understanding of cytomegalovirus infection in immunocompetent individuals.
Cytomegalovirus (CMV) research in immune senescence comes of age: overview of the 6th International Workshop on CMV and Immunosenescence.
Incoming human cytomegalovirus pp65 (UL83) contained in apoptotic infected fibroblasts is cross-presented to CD8(+) T cells by dendritic cells.
Controlling cytomegalovirus: helping the immune system take the lead.
The 'indirect' effects of cytomegalovirus infection.
Cytomegalovirus immune evasion.
Ub to no good: How cytomegaloviruses exploit the ubiquitin proteasome system.
The CMV-encoded G protein-coupled receptors M33 and US28 play pleiotropic roles in immune evasion and alter host T cell responses.
Rapid degradation of the heavy chain of class I major histocompatibility complex antigens in the endoplasmic reticulum of human cytomegalovirus-infected cells.
Human cytomegalovirus tegument protein pUL83 inhibits IFI16-mediated DNA sensing for immune evasion.
Virion Glycoprotein-Mediated Immune Evasion by Human Cytomegalovirus: a Sticky Virus Makes a Slick Getaway.
The role of the human cytomegalovirus UL111A gene in down-regulating CD4+ T-cell recognition of latently infected cells: implications for virus elimination during latency.
In vivo impact of cytomegalovirus evasion of CD8 T-cell immunity: facts and thoughts based on murine models.
Murine cytomegalovirus immune evasion proteins operative in the MHC class I pathway of antigen processing and presentation: state of knowledge, revisions, and questions.
Emerging roles of cytomegalovirus-encoded G protein-coupled receptors during lytic and latent infection.
The Human Cytomegalovirus pUL145 Isoforms Act as Viral DDB1-Cullin-Associated Factors to Instruct Host Protein Degradation to Impede Innate Immunity.

Extensive sequence variation exists among isolates of murine cytomegalovirus within members of the m02 family of genes.
Navigating the Host Cell Response during Entry into Sites of Latent Cytomegalovirus Infection.
Human cytomegalovirus (HCMV) US2 protein interacts with human CD1d (hCD1d) and down-regulates invariant NKT (iNKT) cell activity.
HCMV pUL135 remodels the actin cytoskeleton to impair immune recognition of infected cells.
The immunogenicity of human and murine cytomegaloviruses.
The immediate early 2 protein of human cytomegalovirus (HCMV) mediates the apoptotic control in HCMV retinitis through up-regulation of the cellular FLICE-inhibitory protein expression.
Advances in cytomegalovirus (CMV) biology and its relationship to health, diseases, and aging.
Diagnostic approaches to cytomegalovirus infection in bone marrow and organ transplantation.
Down-regulation of the NKG2D ligand MICA by the human cytomegalovirus glycoprotein UL142.
HCMV-Encoded NK Modulators: Lessons From in vitro and in vivo Genetic Variation.
Tegument proteins of human cytomegalovirus.
Effect of human cytomegalovirus on expression of MHC class I-related chains A.
Human cytomegalovirus protein pUL36: A dual cell death pathway inhibitor.
The Expression of Human Cytomegalovirus MicroRNA MiR-UL148D during Latent Infection in Primary Myeloid Cells Inhibits Activin A-triggered Secretion of IL-6.
Human cytomegalovirus retinitis: pathogenicity, immune evasion and persistence.
Human cytomegalovirus-encoded US9 targets MAVS and STING signaling to evade type I interferon immune responses.
Human cytomegalovirus latency and reactivation - a delicate balance between the virus and its host's immune system.
The Human Cytomegalovirus Nonstructural Glycoprotein UL148 Reorganizes the Endoplasmic Reticulum.
Current and new cytomegalovirus antivirals and novel animal model strategies.
Human Cytomegalovirus Tegument Protein pp65 (pUL83) Dampens Type I Interferon Production by Inactivating the DNA Sensor cGAS without Affecting STING.
Structural and Functional Dissection of the Human Cytomegalovirus Immune Evasion Protein US6.
Molecular modelling of the HCMV IL-10 protein isoforms and analysis of their interaction with the human IL-10 receptor.
Viral chemokine receptors and chemokines in human cytomegalovirus trafficking and interaction with the immune system. CMV chemokine receptors.
Human cytomegalovirus UL7, a homologue of the SLAM-family receptor CD229, impairs cytokine production.
Identification and Functional Characterization of a Novel Fc Gamma-Binding Glycoprotein in Rhesus Cytomegalovirus.
Immunomodulation by cytomegaloviruses: manipulative strategies beyond evasion.
Identification of cytomegalovirus-specific cytotoxic T lymphocytes in vitro is greatly enhanced by the use of recombinant virus lacking the US2 to US11 region or modified vaccinia virus Ankara expressing individual viral genes.
Immune escape and exploitation strategies of cytomegaloviruses: impact on and imitation of the major histocompatibility system.
Effects of human cytomegalovirus infection on ligands for the activating NKG2D receptor of NK cells: up-regulation of UL16-binding protein (ULBP)1 and ULBP2 is counteracted by the viral UL16 protein.

The Human Cytomegalovirus US27 Gene Product Constitutively Activates Antioxidant Response Element-Mediated Transcription through G( $\beta$ ) $\gamma$ , Phosphoinositide 3-Kinase, and Nuclear Respiratory Factor 1.
Human cytomegalovirus attenuates interleukin-1 $\beta$ and tumor necrosis factor $\alpha$ proinflammatory signaling by inhibition of NF- $\kappa$ B activation.
Strain Variation and Disease Severity in Congenital Cytomegalovirus Infection: In Search of a Viral Marker.
Neuropathogenesis in cytomegalovirus infection: indication of the mechanisms using mouse models.
Human cytomegalovirus: host immune modulation by the viral US3 gene.
Potent immunosuppressive activities of cytomegalovirus-encoded interleukin-10.
Cutting edge: a novel viral TNF receptor superfamily member in virulent strains of human cytomegalovirus.
Human cytomegalovirus inhibits IFN- $\alpha$ -stimulated antiviral and immunoregulatory responses by blocking multiple levels of IFN- $\alpha$ signal transduction.
The immune response to human CMV.
Stimulatory effects of human cytomegalovirus tegument protein pp71 lead to increased expression of CCL2 (monocyte chemotactic protein-1) during infection.
Sequences of complete human cytomegalovirus genomes from infected cell cultures and clinical specimens.
Interaction between HCMV pUL83 and human AIM2 disrupts the activation of the AIM2 inflammasome.
Complete sequence and genomic analysis of rhesus cytomegalovirus.
Exosomal release of the virus-encoded chemokine receptor US28 contributes to chemokine scavenging.
Congenital cytomegalovirus infection: molecular mechanisms mediating viral pathogenesis.
Preemptive CD8 T-cell immunotherapy of acute cytomegalovirus infection prevents lethal disease, limits the burden of latent viral genomes, and reduces the risk of virus recurrence.
Human cytomegalovirus antagonizes activation of Fc $\gamma$ receptors by distinct and synergizing modes of IgG manipulation.
Clinical and immunologic aspects of cytomegalovirus infection in solid organ transplant recipients.
Monocytes Latently Infected with Human Cytomegalovirus Evade Neutrophil Killing.
Recombinant HCMV UL128 expression and functional identification of PBMC-attracting activity in vitro.
Quantitative competitive NASBA for measuring mRNA expression levels of the immediate early 1, late pp67, and immune evasion genes US3, US6 and US11 in cells infected with human cytomegalovirus.
Receptor expression and responsiveness of human peripheral blood mononuclear cells to a human cytomegalovirus encoded CC chemokine.
Shaping phenotype, function, and survival of dendritic cells by cytomegalovirus-encoded IL-10.
The human cytomegalovirus major immediate-early proteins as antagonists of intrinsic and innate antiviral host responses.
Recombinant human cytomegalovirus (HCMV) RL13 binds human immunoglobulin G Fc.
The UL144 gene product of human cytomegalovirus activates NF $\kappa$ B via a TRAF6-dependent mechanism.
Loss of linkage disequilibrium and accelerated protein divergence in duplicated cytomegalovirus chemokine genes.
Cytomegalovirus following stem cell transplantation: from pharmacologic to immunologic therapy.
Viral immunomodulatory proteins: usurping host genes as a survival strategy.

SOCS and Herpesviruses, With Emphasis on Cytomegalovirus Retinitis.
Two novel human cytomegalovirus NK cell evasion functions target MICA for lysosomal degradation.
Human cytomegalovirus UL55, UL144, and US28 genotype distribution in infants infected congenitally or postnatally.
Susceptibility of immature and mature Langerhans cell-type dendritic cells to infection and immunomodulation by human cytomegalovirus.
Immune modulation by the human cytomegalovirus-encoded molecule UL18, a mystery yet to be solved.
Immune evasion proteins enhance cytomegalovirus latency in the lungs.
Cytomegalovirus retinitis and the acquired immunodeficiency syndrome--bench to bedside: LXVII Edward Jackson Memorial Lecture.
Cytomegalovirus-induced transendothelial cell migration. a closer look at intercellular communication mechanisms.
Self or nonself? That is the question: sensing of cytomegalovirus infection by innate immune receptors.
Human cytomegalovirus interleukin-10 polarizes monocytes toward a deactivated M2c phenotype to repress host immune responses.
Feeling manipulated: cytomegalovirus immune manipulation.
Generation and Application of Mouse Monoclonal Antibody Against Human Cytomegalovirus UL23.
The Roseoloviruses Downregulate the Protein Tyrosine Phosphatase PTPRC (CD45).
Humoral immune responses and cytomegalovirus excretion in children with asymptomatic infection.
A novel CC-chemokine homolog encoded by guinea pig cytomegalovirus.
An Attenuated CMV Vaccine with a Deletion in Tegument Protein GP83 (pp65 Homolog) Protects against Placental Infection and Improves Pregnancy Outcome in a Guinea Pig Challenge Model.
Genetic markers of immunoglobulin G and immunity to cytomegalovirus in patients with breast cancer.
Murine Cytomegalovirus MCK-2 Facilitates In Vivo Infection Transfer from Dendritic Cells to Salivary Gland Acinar Cells.
Host-Adapted Gene Families Involved in Murine Cytomegalovirus Immune Evasion.
Cytomegalovirus immune evasion of myeloid lineage cells.
Viral and Cellular Factors Contributing to the Hematogenous Dissemination of Human Cytomegalovirus via Polymorphonuclear Leukocytes.
Choice of Study Populations for Vaccines.
Diagnosis and treatment of cytomegalovirus infection.
Open reading frames carried on UL/b' are implicated in shedding and horizontal transmission of rhesus cytomegalovirus in rhesus monkeys.
Evidence for positive selection in the extracellular domain of human cytomegalovirus encoded G protein-coupled receptor US28.
Live or let die: manipulation of cellular suicide programs by murine cytomegalovirus.
Human cytomegalovirus chemokine receptor US28 induces migration of cells on a CX3CL1-presenting surface.
The human cytomegalovirus microRNA miR-UL112 acts synergistically with a cellular microRNA to escape immune elimination.
Mouse cytomegalovirus encoded immunoevasins and evolution of Ly49 receptors - Sidekicks or enemies?
Cytomegalovirus: a culprit or protector in multiple sclerosis?
The Natural Selection of Herpesviruses and Virus-Specific NK Cell Receptors.

Transcriptional regulation of the human cytomegalovirus US11 early gene.
Murine Cytomegalovirus Deubiquitinase Regulates Viral Chemokine Levels To Control Inflammation and Pathogenesis.
Human cytomegalovirus evades antibody-mediated immunity through endoplasmic reticulum-associated degradation of the FcRn receptor.
Impact of CMV upon immune aging: facts and fiction.
Vaccine Development for Cytomegalovirus.
A novel human cytomegalovirus locus modulates cell type-specific outcomes of infection.
Structure of human cytomegalovirus UL141 binding to TRAIL-R2 reveals novel, non-canonical death receptor interactions.
Cytomegalovirus protein m154 perturbs the adaptor protein-1 compartment mediating broad-spectrum immune evasion.
Specific recognition of the viral protein UL18 by CD85j/LIR-1/ILT2 on CD8+ T cells mediates the non-MHC-restricted lysis of human cytomegalovirus-infected cells.
LILRB1 polymorphisms influence posttransplant HCMV susceptibility and ligand interactions.
A prospective assessment of cytomegalovirus immune evasion gene transcription profiles in transplant patients with cytomegalovirus infection.
Structural mechanism of CRL4-instructed STAT2 degradation via a novel cytomegaloviral DCAF receptor.
Identification and functional analyses of a cell-death inhibitor encoded by guinea pig cytomegalovirus gp38.1 in cell culture and in animals.
A live guinea pig cytomegalovirus vaccine deleted of three putative immune evasion genes is highly attenuated but remains immunogenic in a vaccine/challenge model of congenital cytomegalovirus infection.
Expanding the Known Functional Repertoire of the Human Cytomegalovirus pp71 Protein.
Guinea pig cytomegalovirus protective T cell antigen GP83 is a functional pp65 homolog for innate immune evasion and pentamer dependent virus tropism.
Cellular proteins bind to sequence motifs in the R1 element between the HCMV immune evasion genes.
The Guinea pig cytomegalovirus GP119.1 gene encodes an IgG-binding glycoprotein that is incorporated into the virion.
Effect of host genetics on the development of cytomegalovirus retinitis in patients with AIDS.
Endoplasmic Reticulum (ER) Reorganization and Intracellular Retention of CD58 Are Functionally Independent Properties of the Human Cytomegalovirus ER-Resident Glycoprotein UL148.
Modulation of innate and adaptive immunity by cytomegaloviruses.
Signaling Lymphocytic Activation Molecule Family Receptor Homologs in New World Monkey Cytomegaloviruses.
Cytomegalovirus encephalitis in an immunocompetent child: a sceptic diagnosis.
Human cytomegalovirus UL40 signal peptide regulates cell surface expression of the NK cell ligands HLA-E and gpUL18.
The m15 Locus of Murine Cytomegalovirus Modulates Natural Killer Cell Responses to Promote Dissemination to the Salivary Glands and Viral Shedding.
The murine cytomegalovirus chemokine homolog, m131/129, is a determinant of viral pathogenicity.
Block of death-receptor apoptosis protects mouse cytomegalovirus from macrophages and is a determinant of virulence in immunodeficient hosts.
An attenuated cytomegalovirus vaccine with a deletion of a viral chemokine gene is protective against congenital CMV transmission in a guinea pig model.

Evasion of the Host Immune Response by Betaherpesviruses.
Cytomegalovirus m154 hinders CD48 cell-surface expression and promotes viral escape from host natural killer cell control.
Characterization of the human cytomegalovirus UL34 gene.
Innate immune responses to cytomegalovirus infection in the developing mouse brain and their evasion by virus-infected neurons.
Compatibility at amino acid position 98 of MICB reduces the incidence of graft-versus-host disease in conjunction with the CMV status.
The carboxy terminal region of the human cytomegalovirus immediate early 1 (IE1) protein disrupts type II inteferon signaling.
The cytomegalovirus m155 gene product subverts natural killer cell antiviral protection by disruption of H60-NKG2D interactions.
Deciphering the role of DC subsets in MCMV infection to better understand immune protection against viral infections.
A slowly cleaved viral signal peptide acts as a protein-integral immune evasion domain.
Stochastic Episodes of Latent Cytomegalovirus Transcription Drive CD8 T-Cell "Memory Inflation" and Avoid Immune Evasion.
Structure of the HCMV UL16-MICB complex elucidates select binding of a viral immunoevasin to diverse NKG2D ligands.
The salivary glands as a privileged site of cytomegalovirus immune evasion and persistence.
Effect of the R1 element on expression of the US3 and US6 immune evasion genes of human cytomegalovirus.
Positivity of cytomegalovirus antibodies predicts a better clinical and radiological outcome in multiple sclerosis patients.
Human cytomegalovirus UL34 binds to multiple sites within the viral genome.
Murine cytomegalovirus infection inhibits tumor necrosis factor alpha responses in primary macrophages.
Cdc42 - A tryst between host cholesterol metabolism and infection.
Detection of Cytomegalovirus Interleukin 10 (cmvIL-10) by Enzyme-Linked Immunosorbent Assay (ELISA).
RANTES binding and down-regulation by a novel human herpesvirus-6 beta chemokine receptor.
Subversion of natural killer cell responses by a cytomegalovirus-encoded soluble CD48 decoy receptor.
The structure of cytomegalovirus immune modulator UL141 highlights structural Ig-fold versatility for receptor binding.
Comparison of vaccination with rhesus CMV (RhCMV) soluble gB with a RhCMV replication-defective virus deleted for MHC class I immune evasion genes in a RhCMV challenge model.
Activator protein 1-mediated transcriptional regulation strategy sustains long-term expression of a xenogeneic gene product in vivo: an implication for gene therapy targeting congenital protein deficiencies.
Chemokine scavenging by the human cytomegalovirus chemokine decoy receptor US28 does not inhibit monocyte adherence to activated endothelium.
Viral GPCR US28 can signal in response to chemokine agonists of nearly unlimited structural degeneracy.
Identification of the functional domains of the essential human cytomegalovirus UL34 proteins.
Characterization of Antibody Bipolar Bridging Mediated by the Human Cytomegalovirus Fc Receptor gp68.

Modulation of Clr Ligand Expression and NKR-P1 Receptor Function during Murine Cytomegalovirus Infection.
Immune evasion by cytomegalovirus--survival strategies of a highly adapted opportunist.
Viruses disregulate cell membrane receptors to become immune fugitives.
Cytomegalovirus restricts ICOSL expression on antigen-presenting cells disabling T cell co-stimulation and contributing to immune evasion.
Divergent Traits and Ligand-Binding Properties of the Cytomegalovirus CD48 Gene Family.
Roles of GP33, a guinea pig cytomegalovirus-encoded G protein-coupled receptor homolog, in cellular signaling, viral growth and inflammation in vitro and in vivo.
Mutations in the temperature-sensitive murine cytomegalovirus (MCMV) mutants tsm5 and tsm30: a study of genes involved in immune evasion, DNA packaging and processing, and DNA replication.
Two glycosaminoglycan-binding domains of the mouse cytomegalovirus-encoded chemokine MCK-2 are critical for oligomerization of the full-length protein.
Vesicular stomatitis virus infection promotes immune evasion by preventing NKG2D-ligand surface expression.
Differentiation and Protective Capacity of Virus-Specific CD8(+) T Cells Suggest Murine Norovirus Persistence in an Immune-Privileged Enteric Niche.
Torquetenovirus (TTV) load is associated with mortality in Italian elderly subjects.
N-glycosylation of mouse TRAIL-R and human TRAIL-R1 enhances TRAIL-induced death.
Rat cytomegalovirus-encoded $\gamma$ -chemokine vXCL1 is a highly adapted, species-specific agonist for rat XCR1-positive dendritic cells.
A herpesvirus entry mediator mutein with selective agonist action for the inhibitory receptor B and T lymphocyte attenuator.
Differential signaling of cmvIL-10 through common variants of the IL-10 receptor 1.

Table A2-28, Cluster 27

Cluster 27 focuses on Mycobacterium tuberculosis, especially its mechanisms for evading the host immune response (450)
Surviving the macrophage: tools and tricks employed by Mycobacterium tuberculosis.
MicroRNA-325-3p Facilitates Immune Escape of Mycobacterium tuberculosis through Targeting LNX1 via NEK6 Accumulation to Promote Anti-Apoptotic STAT3 Signaling.
The role of dendritic cells in Mycobacterium tuberculosis infection.
Immune evasion and provocation by Mycobacterium tuberculosis.
Genes and regulatory networks involved in persistence of Mycobacterium tuberculosis.
Outcome of Mycobacterium tuberculosis and Toll-like receptor interaction: immune response or immune evasion?
[Mycobacterium tuberculosis virulence factors and its immune evasion mechanisms].
Characterization of a secretory hydrolase from Mycobacterium tuberculosis sheds critical insight into host lipid utilization by M. tuberculosis.
Strategies Employed to Evade the Host Immune Response and the Mechanism of Drug Resistance in Mycobacterium tuberculosis: In Search of Finding New Targets.
Immunology of Mycobacterium tuberculosis Infections.
Autophagy in Mycobacterium tuberculosis infection: a passepartout to flush the intruder out?
Evasion of Innate and Adaptive Immunity by Mycobacterium tuberculosis.
Potential Effect of Statins on Mycobacterium tuberculosis Infection.



Dormant Mycobacterium tuberculosis fails to block phagosome maturation and shows unexpected capacity to stimulate specific human T lymphocytes.
Host-pathogen interactions during Mycobacterium tuberculosis infections.
Clinical manifestations and immune response to tuberculosis.
Mycobacterium tuberculosis: Manipulator of Protective Immunity.
Modulation of cell death by M. tuberculosis as a strategy for pathogen survival.
[Advances in understanding of the virulence mechanism of Mycobacterium tuberculosis].
How far have we reached in tuberculosis vaccine development?
Immune evasion by Mycobacterium tuberculosis: living with the enemy.
Tweaking host immune responses for novel therapeutic approaches against Mycobacterium tuberculosis.
Comparative study of interruption of signaling pathways in lung epithelial cell by two different Mycobacterium tuberculosis lineages.
Gene expression profile analysis and target gene discovery of Mycobacterium tuberculosis biofilm.
Immunology of tuberculosis and implications in vaccine development.
Attack on the scourge of tuberculosis: patented drug targets.
Inhibition of Autophagy by MiR-30A Induced by Mycobacteria tuberculosis as a Possible Mechanism of Immune Escape in Human Macrophages.
Latency-associated protein Acr1 impairs dendritic cell maturation and functionality: a possible mechanism of immune evasion by Mycobacterium tuberculosis.
Host evasion and exploitation schemes of Mycobacterium tuberculosis.
Role of glycans and glycoproteins in disease development by Mycobacterium tuberculosis.
Multiplexed Strain Phenotyping Defines Consequences of Genetic Diversity in Mycobacterium tuberculosis for Infection and Vaccination Outcomes.
Keto-mycolic acid-dependent pellicle formation confers tolerance to drug-sensitive Mycobacterium tuberculosis.
[Protective immunity against Mycobacterium tuberculosis].
Phagosomal rupture by Mycobacterium tuberculosis results in toxicity and host cell death.
M. tuberculosis T Cell Epitope Analysis Reveals Paucity of Antigenic Variation and Identifies Rare Variable TB Antigens.
Mycobacterium tuberculosis impairs dendritic cell response by altering CD1b, DC-SIGN and MR profile.
Mycobacterium tuberculosis associated with severe tuberculosis evades cytosolic surveillance systems and modulates IL-1 $\beta$ production.
Orchestration of pulmonary T cell immunity during Mycobacterium tuberculosis infection: immunity interruptus.
Epigenetic orchestration of host immune defences by Mycobacterium tuberculosis.
KefB inhibits phagosomal acidification but its role is unrelated to M. tuberculosis survival in host.
Manipulation and exploitation of host immune system by pathogenic Mycobacterium tuberculosis for its advantage.
Immuno-evasion and immunosuppression of the macrophage by Mycobacterium tuberculosis.
Mechanisms of M. tuberculosis Immune Evasion as Challenges to TB Vaccine Design.
Emerging advances in identifying signal transmission molecules involved in the interaction between Mycobacterium tuberculosis and the host.
Mycobacterium tuberculosis inhibits IFN-gamma transcriptional responses without inhibiting activation of STAT1.

Crosstalk between Mycobacterium tuberculosis and the host cell.
Mycobacterium tuberculosis-macrophage interaction: Molecular updates.
A recombinant attenuated Mycobacterium tuberculosis vaccine strain is safe in immunosuppressed simian immunodeficiency virus-infected infant macaques.
Immune cell interactions in tuberculosis.
The knowns and unknowns of latent Mycobacterium tuberculosis infection.
Human mesenchymal stem cells: New sojourn of bacterial pathogens.
Evolutionary Genetics of Mycobacterium tuberculosis and HIV-1: "The Tortoise and the Hare".
Mycobacterium tuberculosis adhesins: potential biomarkers as anti-tuberculosis therapeutic and diagnostic targets.
Mycobacterium tuberculosis Rv3034c regulates mTORC1 and PPAR-γ dependent pexophagy mechanism to control redox levels in macrophages.
The Immune Escape Mechanisms of Mycobacterium Tuberculosis.
[Insight into tuberculosis pathogenic mechanism from the zebra fish-Mycobacterium marinum model -a review].
Ins and outs of Mycobacterium tuberculosis PPE family in pathogenesis and implications for novel measures against tuberculosis.
Latent tuberculosis: interaction of virulence factors in Mycobacterium tuberculosis.
Mycobacterium tuberculosis evades host immunity by recruiting mesenchymal stem cells.
Toll-like receptor 2-dependent extracellular signal-regulated kinase signaling in Mycobacterium tuberculosis-infected macrophages drives anti-inflammatory responses and inhibits Th1 polarization of responding T cells.
Understanding the early host immune response against Mycobacterium tuberculosis.
TREM2 Promotes Immune Evasion by Mycobacterium tuberculosis in Human Macrophages.
The role of epigenetics, bacterial and host factors in progression of Mycobacterium tuberculosis infection.
Mycobacterial persistence and immunity.
Mycobacterium Tuberculosis and Interactions with the Host Immune System: Opportunities for Nanoparticle Based Immunotherapeutics and Vaccines.
Evasion and subversion of antigen presentation by Mycobacterium tuberculosis.
Enhanced priming of adaptive immunity by a proapoptotic mutant of Mycobacterium tuberculosis.
Evasion of innate immunity by Mycobacterium tuberculosis: is death an exit strategy?
Stealth Strategies of Mycobacterium tuberculosis for Immune Evasion.
Immunotherapy for TB.
Granulocyte chemotactic properties of M. tuberculosis versus M. bovis-infected bovine alveolar macrophages.
Natural and trained innate immunity against Mycobacterium tuberculosis.
Conserved alanine rich protein Rv3878 in Mycobacterium tuberculosis contains sequence polymorphisms.
Initiation and regulation of T-cell responses in tuberculosis.
AIDS associated tuberculosis: a catastrophic collision to evade the host immune system.
An outer membrane channel protein of Mycobacterium tuberculosis with exotoxin activity.
Importance of differential identification of Mycobacterium tuberculosis strains for understanding differences in their prevalence, treatment efficacy, and vaccine development.
Pathological and protective roles of dendritic cells in Mycobacterium tuberculosis infection: Interaction between host immune responses and pathogen evasion.

Cellular and humoral mechanisms involved in the control of tuberculosis.
Macrophage: A Cell With Many Faces and Functions in Tuberculosis.
Host-Pathogen Interaction as a Novel Target for Host-Directed Therapies in Tuberculosis.
[The role of mycobacterial secretory proteins in immune response in tuberculosis].
The PPE2 protein of Mycobacterium tuberculosis translocates to host nucleus and inhibits nitric oxide production.
New insights into the evasion of host innate immunity by Mycobacterium tuberculosis.
Mycobacterium tuberculosis Rv1987 induces Th2 immune responses and enhances Mycobacterium smegmatis survival in mice.
Partial redundancy of the pattern recognition receptors, scavenger receptors, and C-type lectins for the long-term control of Mycobacterium tuberculosis infection.
Regulation of antigen presentation by Mycobacterium tuberculosis: a role for Toll-like receptors.
Codominance of TLR2-dependent and TLR2-independent modulation of MHC class II in Mycobacterium tuberculosis infection in vivo.
Mycobacterium tuberculosis PPE51 Inhibits Autophagy by Suppressing Toll-Like Receptor 2-Dependent Signaling.
Conserved hypothetical protein Rv1977 in Mycobacterium tuberculosis strains contains sequence polymorphisms and might be involved in ongoing immune evasion.
Molecular mechanism of interaction of Mycobacterium tuberculosis with host macrophages under high glucose conditions.
Pathogenesis in tuberculosis: transcriptomic approaches to unraveling virulence mechanisms and finding new drug targets.
Mycobacterium tuberculosis Rv0431 expressed in Mycobacterium smegmatis, a potentially mannosylated protein, mediated the immune evasion of RAW 264.7 macrophages.
Genetic diversity of immune-related antigens in Region of Difference 2 of Mycobacterium tuberculosis strains.
Virulence Factors of Mycobacterium tuberculosis as Modulators of Cell Death Mechanisms.
Sterilization by Adaptive Immunity of a Conditionally Persistent Mutant of Mycobacterium tuberculosis.
Mycobacterium tuberculosis cords within lymphatic endothelial cells to evade host immunity.
Small RNA MTS1338 Configures a Stress Resistance Signature in Mycobacterium tuberculosis.
[Mycobacterium tuberculosis main immune response evasion mechanisms].
Secretory proteins of Mycobacterium tuberculosis and their roles in modulation of host immune responses: focus on therapeutic targets.
Increased expression of Fas ligand on Mycobacterium tuberculosis infected macrophages: a potential novel mechanism of immune evasion by Mycobacterium tuberculosis?
Genome wide approaches discover novel Mycobacterium tuberculosis antigens as correlates of infection, disease, immunity and targets for vaccination.
Is targeting dysregulation in apoptosis splice variants in Mycobacterium tuberculosis (MTB) host interactions and splicing factors resulting in immune evasion by MTB strategies a possibility?
"The Impact of Mycobacterium tuberculosis Immune Evasion on Protective Immunity: Implications for TB Vaccine Design" - Meeting report.
Mycolic acid modification by the mmaA4 gene of M. tuberculosis modulates IL-12 production.
Prime-boost with Mycobacterium smegmatis recombinant vaccine improves protection in mice infected with Mycobacterium tuberculosis.
Antigen 85 complex as a powerful Mycobacterium tuberculosis immunogene: Biology, immune-pathogenicity, applications in diagnosis, and vaccine design.

Mycobacteria manipulate macrophage recruitment through coordinated use of membrane lipids.
The challenge of new drug discovery for tuberculosis.
Mycobacterium tuberculosis may escape helper T cell recognition by infecting human fibroblasts.
GSK-3 $\alpha$ / $\beta$ Activity Negatively Regulates MMP-1/9 Expression to Suppress Mycobacterium tuberculosis Infection.
Efferocytosis is an innate antibacterial mechanism.
Mycobacterium tuberculosis GroEL2 Modulates Dendritic Cell Responses.
Mycobacterium tuberculosis cell wall glycolipids directly inhibit CD4+ T-cell activation by interfering with proximal T-cell-receptor signaling.
pstS1 polymorphisms of Mycobacterium tuberculosis strains may reflect ongoing immune evasion.
The truncated Rv2820c of Mycobacterium tuberculosis Beijing family augments intracellular survival of M. smegmatis by altering cytokine profile and inhibiting NO generation.
The ubiquitin ligase TRIM32 promotes the autophagic response to Mycobacterium tuberculosis infection in macrophages.
Human Pulmonary Tuberculosis: Understanding the Immune Response in the Bronchoalveolar System.
Toll-like receptor 2 in host defense against Mycobacterium tuberculosis: to be or not to be-that is the question.
Single nucleotide polymorphism in Ag85 genes of Mycobacterium tuberculosis complex: analysis of 178 clinical isolates from China and 13 BCG strains.
Fighting Persistence: How Chronic Infections with Mycobacterium tuberculosis Evade T Cell-Mediated Clearance and New Strategies To Defeat Them.
Monocyte-derived dendritic cells early exposed to Mycobacterium tuberculosis induce an enhanced T helper 17 response and transfer mycobacterial antigens.
Structure Based Discovery of Inhibitors of CYP125 and CYP142 from Mycobacterium tuberculosis.
Functional, biochemical and 3D studies of Mycobacterium tuberculosis protein peptides for an effective anti-tuberculosis vaccine.
Modeling phenotypic metabolic adaptations of Mycobacterium tuberculosis H37Rv under hypoxia.
Host-Directed Therapies for Tuberculosis.
Recombinant Rv3261 protein of Mycobacterium tuberculosis induces apoptosis through a mitochondrion-dependent pathway in macrophages and inhibits intracellular bacterial growth.
Perspectives on host adaptation in response to Mycobacterium tuberculosis: modulation of inflammation.
Addressing the Challenges of Tuberculosis: A Brief Historical Account.
Evidence for a rapid rate of molecular evolution at the hypervariable and immunogenic Mycobacterium tuberculosis PPE38 gene region.
[Advances in the study of Mycobacterium tuberculosis protein phosphatase and its inhibitors].
Mycobacterium tuberculosis and the host cell inflammasome: a complex relationship.
The role of IL-10 in immune regulation during M. tuberculosis infection.
Emerging Role of Exosomes in Tuberculosis: From Immunity Regulations to Vaccine and Immunotherapy.
The complete genome sequence of Mycobacterium bovis.
Virulence factors of the Mycobacterium tuberculosis complex.
MiRNAs in tuberculosis: Their decisive role in the fate of TB.
Mycobacterium tuberculosis inhibits maturation of human monocyte-derived dendritic cells in vitro.
A Phenotypic Characterization of Two Isolates of a Multidrug-Resistant Outbreak Strain of Mycobacterium tuberculosis with Opposite Epidemiological Fitness.

Understanding and overcoming the barriers to T cell-mediated immunity against tuberculosis.
Identification and Repurposing of Trisubstituted Harmine Derivatives as Novel Inhibitors of Mycobacterium tuberculosis Phosphoserine Phosphatase.
Human tuberculosis and Mycobacterium tuberculosis complex: A review on genetic diversity, pathogenesis and omics approaches in host biomarkers discovery.
The innate immune response in human tuberculosis.
Molecular basis of mycobacterial survival in macrophages.
Innate immunity in tuberculosis: host defense vs pathogen evasion.
Role of HrcA in stress management in Mycobacterium tuberculosis.
Voltage gated calcium channels negatively regulate protective immunity to Mycobacterium tuberculosis.
Sequencing-relative to hybridization-based transcriptomics approaches better define Mycobacterium tuberculosis stress-response regulons.
Mycobacterial para-Hydroxybenzoic Acid-Derivatives (pHBADs) and Related Structures Induce Macrophage Innate Memory.
The Role of Complement System and the Immune Response to Tuberculosis Infection.
A novel genetically engineered Mycobacterium smegmatis-based vaccine promotes anti-TB immunity.
Mycobacterium tuberculosis Rv1987 protein induces M2 polarization of macrophages through activating the PI3K/Akt1/mTOR signaling pathway.
Genetic diversity of antigens Rv2945c and Rv0309 in Mycobacterium tuberculosis strains may reflect ongoing immune evasion.
Mycobacterium tuberculosis subverts innate immunity to evade specific effectors.
Mycobacterium tuberculosis hijacks ubiquitin to inhibit pyroptosis.
Several Routes to the Same Destination: Inhibition of Phagosome-Lysosome Fusion by Mycobacterium tuberculosis.
Mycobacterium tuberculosis Prolyl Oligopeptidase Induces In vitro Secretion of Proinflammatory Cytokines by Peritoneal Macrophages.
Polymorphisms of human T cell epitopes of Mycobacterium tuberculosis indicate divergence of host immune pressure on different categories of proteins.
Mycobacterium tuberculosis exploits the PPM1A signaling pathway to block host macrophage apoptosis.
The Mycobacterium tuberculosis SecA2 system subverts phagosome maturation to promote growth in macrophages.
Induction of apoptosis in human neutrophils by Mycobacterium tuberculosis is dependent on mature bacterial lipoproteins.
For better or for worse: the immune response against Mycobacterium tuberculosis balances pathology and protection.
Mycobacterium tuberculosis inhibits neutrophil apoptosis, leading to delayed activation of naive CD4 T cells.
The secret trumps, impelling the pathogenicity of tubercle bacilli.
Challenging Mycobacterium tuberculosis dormancy mechanisms and their immunodiagnostic potential.
Roles of PE_PGRS family in Mycobacterium tuberculosis pathogenesis and novel measures against tuberculosis.
The down-regulation of cathepsin G in THP-1 monocytes after infection with Mycobacterium tuberculosis is associated with increased intracellular survival of bacilli.

Annexin1 regulates DC efferocytosis and cross-presentation during Mycobacterium tuberculosis infection.
Mycobacterium tuberculosis Evasion of Guanylate Binding Protein-Mediated Host Defense in Mice Requires the ESX1 Secretion System.
The M. tuberculosis HAD phosphatase (Rv3042c) interacts with host proteins and is inhibited by Clofazimine.
Mycobacterium tuberculosis Exploits a Molecular Off Switch of the Immune System for Intracellular Survival.
Mycobacterial lipolytic enzymes: a gold mine for tuberculosis research.
Mycobacterium tuberculosis Lipoprotein and Lipoglycan Binding to Toll-Like Receptor 2 Correlates with Agonist Activity and Functional Outcomes.
Resistance to cellular autophagy by Mycobacterium tuberculosis Beijing strains.
Mycobacterial Genetic Technologies for Probing the Host-Pathogen Microenvironment.
Differential macrophage response to slow- and fast-growing pathogenic mycobacteria.
Regulatory Mechanisms of Autophagy-Targeted Antimicrobial Therapeutics Against Mycobacterial Infection.
Dual-targeting GroEL/ES chaperonin and protein tyrosine phosphatase B (PtpB) inhibitors: A polypharmacology strategy for treating Mycobacterium tuberculosis infections.
Rv1273c, an ABC transporter of Mycobacterium tuberculosis promotes mycobacterial intracellular survival within macrophages via modulating the host cell immune response.
Intricate Genetic Programs Controlling Dormancy in Mycobacterium tuberculosis.
Lysosome-Mediated Plasma Membrane Repair Is Dependent on the Small GTPase Arl8b and Determines Cell Death Type in Mycobacterium tuberculosis Infection.
A multiple genome analysis of Mycobacterium tuberculosis reveals specific novel genes and mutations associated with pyrazinamide resistance.
Advancing Adjuvants for Mycobacterium tuberculosis Therapeutics.
Mce4A protein of Mycobacterium tuberculosis induces pro inflammatory cytokine response leading to macrophage apoptosis in a TNF- $\alpha$ dependent manner.
TLR-4/microRNA-125a/NF- $\kappa$ B signaling modulates the immune response to Mycobacterium tuberculosis infection.
Linking inflammasome activation and phagosome maturation.
Persistence of Mycobacterium tuberculosis in response to infection burden and host-induced stressors.
Single nucleotide polymorphisms may explain the contrasting phenotypes of two variants of a multidrug-resistant Mycobacterium tuberculosis strain.
Toll-like receptor 2-dependent inhibition of macrophage class II MHC expression and antigen processing by 19-kDa lipoprotein of Mycobacterium tuberculosis.
Methylation in Mycobacterium-host interaction and implications for novel control measures.
Inhibition of Fatty Acid Oxidation Promotes Macrophage Control of Mycobacterium tuberculosis.
The Tim3-galectin 9 pathway induces antibacterial activity in human macrophages infected with Mycobacterium tuberculosis.
Mycobacterium tuberculosis: immune evasion, latency and reactivation.
Cutting edge: Mycobacterium tuberculosis but not nonvirulent mycobacteria inhibits IFN- $\beta$ and AIM2 inflammasome-dependent IL-1 $\beta$ production via its ESX-1 secretion system.
Distinct Strategies Employed by Dendritic Cells and Macrophages in Restricting Mycobacterium tuberculosis Infection: Different Philosophies but Same Desire.
IFN- $\beta$ : A Contentious Player in Host-Pathogen Interaction in Tuberculosis.

Delivery by Dendritic Mesoporous Silica Nanoparticles Enhances the Antimicrobial Activity of a Napsin-Derived Peptide Against Intracellular Mycobacterium tuberculosis.
Macrophage takeover and the host-bacilli interplay during tuberculosis.
DosR antigen Rv1737c induces activation of macrophages dependent on the TLR2 pathway.
Mycobacterium tuberculosis Catalase Inhibits the Formation of Mast Cell Extracellular Traps.
PE_PGRS30 of Mycobacterium tuberculosis mediates suppression of proinflammatory immune response in macrophages through its PGRS and PE domains.
Extracellular adenosine triphosphate affects the response of human macrophages infected with Mycobacterium tuberculosis.
Clinical isolates of Mycobacterium tuberculosis differ in their ability to induce respiratory burst and apoptosis in neutrophils as a possible mechanism of immune escape.
F420H2 Is Required for Phthiocerol Dimycocerosate Synthesis in Mycobacteria.
The Physiology and Genetics of Oxidative Stress in Mycobacteria.
Why macrophages cannot LAP up TB.
Apoptosis is an innate defense function of macrophages against Mycobacterium tuberculosis.
Down-regulation of hsa_circ_0045474 induces macrophage autophagy in tuberculosis via miR-582-5p/TNKS2 axis.
Evidence for the Effect of Vaccination on Host-Pathogen Interactions in a Murine Model of Pulmonary Tuberculosis by Mycobacterium tuberculosis.
Mycobacterium tuberculosis PE25/PPE41 protein complex induces activation and maturation of dendritic cells and drives Th2-biased immune responses.
Predicted Structural Variability of Mycobacterium tuberculosis PPE18 Protein With Immunological Implications Among Clinical Strains.
The 30-kDa and 38-kDa antigens from Mycobacterium tuberculosis induce partial maturation of human dendritic cells shifting CD4(+) T cell responses towards IL-4 production.
Unraveling Mycobacterium tuberculosis genomic diversity and evolution in Lisbon, Portugal, a highly drug resistant setting.
Pathogenic Mycobacterium bovis strains differ in their ability to modulate the proinflammatory activation phenotype of macrophages.
Mycobacterium tuberculosis induces the miR-33 locus to reprogram autophagy and host lipid metabolism.
Evidence for Highly Variable, Region-Specific Patterns of T-Cell Epitope Mutations Accumulating in Mycobacterium tuberculosis Strains.
Current status of anti-tuberculosis therapy: a patent analysis.
DC-SIGN and mannosylated surface structures of Mycobacterium tuberculosis: a deceptive liaison.
Novel miR-1958 Promotes Mycobacterium tuberculosis Survival in RAW264.7 Cells by Inhibiting Autophagy Via Atg5.
Processing of Mycobacterium tuberculosis antigen 85B involves intraphagosomal formation of peptide-major histocompatibility complex II complexes and is inhibited by live bacilli that decrease phagosome maturation.
Mycobacterium bovis PknG R242P Mutation Results in Structural Changes with Enhanced Virulence in the Mouse Model of Infection.
Cell death and autophagy in tuberculosis.
Recombinant Lipoprotein Rv1016c Derived from Mycobacterium tuberculosis Is a TLR-2 Ligand that Induces Macrophages Apoptosis and Inhibits MHC II Antigen Processing.
Reactive oxygen species production by human dendritic cells involves TLR2 and dectin-1 and is essential for efficient immune response against Mycobacteria.

Rv0132c of Mycobacterium tuberculosis encodes a coenzyme F420-dependent hydroxymycolic acid dehydrogenase.
Mycobacterium tuberculosis LprA is a lipoprotein agonist of TLR2 that regulates innate immunity and APC function.
Mycobacterial escape from macrophage phagosomes to the cytoplasm represents an alternate adaptation mechanism.
Mycobacterium tuberculosis Multidrug-Resistant Strain M Induces Low IL-8 and Inhibits TNF- $\alpha$ Secretion by Bronchial Epithelial Cells Altering Neutrophil Effector Functions.
Epitope mapping of the immunodominant antigen TB10.4 and the two homologous proteins TB10.3 and TB12.9, which constitute a subfamily of the esat-6 gene family.
Expression of inhibitory regulators of innate immunity in patients with active tuberculosis.
Mycobacterium tuberculosis EsxH inhibits ESCRT-dependent CD4(+) T-cell activation.
Cloaked dagger: tRNA slicing by an unlikely culprit.
Host-directed therapies to combat tuberculosis and associated non-communicable diseases.
Current efforts and future prospects in the development of live mycobacteria as vaccines.
The formation of the granuloma in tuberculosis infection.
Dying to live: how the death modality of the infected macrophage affects immunity to tuberculosis.
Mycobacterium tuberculosis lipids regulate cytokines, TLR-2/4 and MHC class II expression in human macrophages.
Type VII secretion systems: structure, functions and transport models.
Histone Methyltransferase SET8 Epigenetically Reprograms Host Immune Responses to Assist Mycobacterial Survival.
Formation of Foamy Macrophages by Tuberculous Pleural Effusions Is Triggered by the Interleukin-10/Signal Transducer and Activator of Transcription 3 Axis through ACAT Upregulation.
Genetic-and-Epigenetic Interspecies Networks for Cross-Talk Mechanisms in Human Macrophages and Dendritic Cells during MTB Infection.
Immunogenicity of Mycobacterial Extracellular Vesicles Isolated From Host-Related Conditions Informs About Tuberculosis Disease Status.
An Army Marches on Its Stomach: Metabolic Intermediates as Antimicrobial Mediators in Mycobacterium tuberculosis Infection.
Mycobacterium tuberculosis inhibits the NLRP3 inflammasome activation via its phosphokinase PknF.
Transcriptional profiling of Mycobacterium tuberculosis replicating ex vivo in blood from HIV- and HIV+ subjects.
Natural products against key Mycobacterium tuberculosis enzymatic targets: Emerging opportunities for drug discovery.
Persistent bacterial infections: the interface of the pathogen and the host immune system.
Mycobacterial infection alters host mitochondrial activity.
C5aR contributes to the weak Th1 profile induced by an outbreak strain of Mycobacterium tuberculosis.
Divergent effects of mycobacterial cell wall glycolipids on maturation and function of human monocyte-derived dendritic cells.
Valproic acid promotes a decrease in mycobacterial survival by enhancing nitric oxide production in macrophages stimulated with IFN- $\gamma$ .
The Mycobacterium tuberculosis PE proteins Rv0285 and Rv1386 modulate innate immunity and mediate bacillary survival in macrophages.
Differential immune response of adipocytes to virulent and attenuated Mycobacterium tuberculosis.



De Novo Fatty Acid Synthesis During Mycobacterial Infection Is a Prerequisite for the Function of Highly Proliferative T Cells, But Not for Dendritic Cells or Macrophages.
Hydrophobic Mycobacterial Antigens Elicit Polyfunctional T Cells in Mycobacterium bovis Immunized Cattle: Association With Protection Against Challenge?
How mycobacteria take advantage of the weakness in human immune system in the modern world.
Outbreaks of Mycobacterium tuberculosis MDR strains differentially induce neutrophil respiratory burst involving lipid rafts, p38 MAPK and Syk.
PPE27 in Mycobacterium smegmatis Enhances Mycobacterial Survival and Manipulates Cytokine Secretion in Mouse Macrophages.
Suppression of Toll-like receptor 2-mediated proinflammatory responses by Mycobacterium tuberculosis protein Rv3529c.
Function prediction and analysis of mycobacterium tuberculosis hypothetical proteins.
Mycobacterium tuberculosis strains from ancient and modern lineages induce distinct patterns of immune responses.
Src homology 3-interacting domain of Rv1917c of Mycobacterium tuberculosis induces selective maturation of human dendritic cells by regulating PI3K-MAPK-NF-kappaB signaling and drives Th2 immune responses.
Pro- and anti-inflammatory cytokines against Rv2031 are elevated during latent tuberculosis: a study in cohorts of tuberculosis patients, household contacts and community controls in an endemic setting.
Clinical isolates of the modern Mycobacterium tuberculosis lineage 4 evade host defense in human macrophages through eluding IL-1 $\beta$ -induced autophagy.
Mycobacterium tuberculosis genes involved in regulation of host cell death.
M. tuberculosis Reprograms Hematopoietic Stem Cells to Limit Myelopoiesis and Impair Trained Immunity.
Virulence-Associated Secretion in Mycobacterium abscessus.
The Rv0805 gene from Mycobacterium tuberculosis encodes a 3',5'-cyclic nucleotide phosphodiesterase: biochemical and mutational analysis.
Host MKRN1-Mediated Mycobacterial PPE Protein Ubiquitination Suppresses Innate Immune Response.
Spotlight on Mycobacteria and dendritic cells: will novel targets to fight tuberculosis emerge?
Genetic diversity of antigen 38 kDa in Mycobacterium tuberculosis strains from China.
MicroRNA-26b inhibits the immune response to Mycobacterium tuberculosis (M.tb) infection in THP-1 cells via targeting TGF $\beta$ -activated kinase-1 (TAK1), a promoter of the NF- $\kappa$ B pathway.
The immunology of other mycobacteria: M. ulcerans, M. leprae.
PE_PGRS: Vital proteins in promoting mycobacterial survival and modulating host immunity and metabolism.
Evaluation of new vaccines for tuberculosis in the guinea pig model.
Immune subversion by Mycobacterium tuberculosis through CCR5 mediated signaling: involvement of IL-10.
Th2-type immune response observed in healthy individuals to sonicate antigen prepared from the most prevalent Mycobacterium tuberculosis strain with single copy of IS6110.
Mixed metal oxide nanoparticles inhibit growth of Mycobacterium tuberculosis into THP-1 cells.
ESX secretion systems: mycobacterial evolution to counter host immunity.
Synthesis of multivalent tuberculosis and Leishmania-associated capping carbohydrates reveals structure-dependent responses allowing immune evasion.
Antigenic Variation and Immune Escape in the MTBC.

Understanding delayed T-cell priming, lung recruitment, and airway luminal T-cell responses in host defense against pulmonary tuberculosis.
[Potential of novel antimycobacterial immune factors, SLPI and lipocalin 2].
Playing hide-and-seek with host macrophages through the use of mycobacterial cell envelope phthiocerol dimycocerosates and phenolic glycolipids.
microRNA-20a Inhibits Autophagic Process by Targeting ATG7 and ATG16L1 and Favors Mycobacterial Survival in Macrophage Cells.
Mycobacterium tuberculosis antigen 85A and 85C structures confirm binding orientation and conserved substrate specificity.
An immunomodulatory role for the Mycobacterium tuberculosis region of difference 1 locus proteins PE35 (Rv3872) and PPE68 (Rv3873).
Incorporation of NKT cell-activating glycolipids enhances immunogenicity and vaccine efficacy of Mycobacterium bovis bacillus Calmette-Guerin.
Mycobacterium tuberculosis Rv1096, facilitates mycobacterial survival by modulating the NF- $\kappa$ B/MAPK pathway as peptidoglycan N-deacetylase.
Continuous treatment with recombinant Mycobacterium tuberculosis CFP-10-ESAT-6 protein activated human monocyte while deactivated LPS-stimulated macrophage.
Cell wall-associated alpha-glucan is instrumental for Mycobacterium tuberculosis to block CD1 molecule expression and disable the function of dendritic cell derived from infected monocyte.
Mycobacterium bovis induces mitophagy to suppress host xenophagy for its intracellular survival.
(1)H, (13)C, (15)N backbone and side-chain chemical shift assignments of the polyketide cyclase from Mycobacterium tuberculosis.
Mining large-scale response networks reveals 'topmost activities' in Mycobacterium tuberculosis infection.
Polymorphism of antigen MPT64 in Mycobacterium tuberculosis strains.
Decreased pathology and prolonged survival of human DC-SIGN transgenic mice during mycobacterial infection.
Inhibition of IFN-gamma-induced class II transactivator expression by a 19-kDa lipoprotein from Mycobacterium tuberculosis: a potential mechanism for immune evasion.
Novel genetic polymorphisms identified in the clinical isolates of Mycobacterium tuberculosis PE_PGRS33 gene modulate cytokines expression and promotes survival in macrophages.
Myeloid C-Type Lectin Receptors in Tuberculosis and HIV Immunity: Insights Into Co-infection?
Mycobacterium tuberculosis Rv1096 protein: gene cloning, protein expression, and peptidoglycan deacetylase activity.
Cytotoxic activity of nucleoside diphosphate kinase secreted from Mycobacterium tuberculosis.
Characteristic genes in THP-1 derived macrophages infected with Mycobacterium tuberculosis H37Rv strain identified by integrating bioinformatics methods.
Uncovering New Pathogen-Host Protein-Protein Interactions by Pairwise Structure Similarity.
PPE38 Protein of Mycobacterium tuberculosis Inhibits Macrophage MHC Class I Expression and Dampens CD8(+) T Cell Responses.
Immunobiology of tubercle bacilli and prospects of immunomodulatory drugs to tackle tuberculosis (TB) and other non-tubercular mycobacterial infections.
Mycobacterial lipid logic.
Role of calcium channels in cellular antituberculosis effects: Potential of voltage-gated calcium-channel blockers in tuberculosis therapy.
Mycobacterium tuberculosis Co-operonic PE32/PPE65 Proteins Alter Host Immune Responses by Hampering Th1 Response.

Nanoscale clustering of mycobacterial ligands and DC-SIGN host receptors are key determinants for pathogen recognition.
Effects of Mycobacterium tuberculosis Rv1096 on mycobacterial cell division and modulation on macrophages.
Mycobacterium tuberculosis and myeloid-derived suppressor cells: Insights into caveolin rich lipid rafts.
Mycobacterium tuberculosis Hip1 modulates macrophage responses through proteolysis of GroEL2.
Deletion of BCG Hip1 protease enhances dendritic cell and CD4 T cell responses.
Unraveling the role of the transcriptional regulator VirS in low pH-induced responses of Mycobacterium tuberculosis and identification of VirS inhibitors.
Novel role of PE_PGRS47 in the alteration of mycobacterial cell wall integrity and drug resistance.
Deletion of BCG_2432c from the Bacillus Calmette-Guérin vaccine enhances autophagy-mediated immunity against tuberculosis.
Impaired dendritic cell differentiation of CD16-positive monocytes in tuberculosis: role of p38 MAPK.
Secreted acid phosphatase (SapM) of Mycobacterium tuberculosis is indispensable for arresting phagosomal maturation and growth of the pathogen in guinea pig tissues.
Targeting Autophagy as a Strategy for Developing New Vaccines and Host-Directed Therapeutics Against Mycobacteria.
Polymorphisms in the PE35 and PPE68 antigens in Mycobacterium tuberculosis strains may affect strain virulence and reflect ongoing immune evasion.
Conserved immune recognition hierarchy of mycobacterial PE/PPE proteins during infection in natural hosts.
Infection of Primary Bovine Macrophages with Mycobacterium avium Subspecies paratuberculosis Suppresses Host Cell Apoptosis.
Vaccine potential of ESAT-6 protein fused with consensus CD4(+) T-cell epitopes of PE/PPE proteins against highly pathogenic Mycobacterium tuberculosis strain HN878.
Identification of peptides presented through the MHC-II of dendritic cells stimulated with Mycobacterium avium.
Identification of Unique Key miRNAs, TFs, and mRNAs in Virulent MTB Infection Macrophages by Network Analysis.
The tuberculosis necrotizing toxin kills macrophages by hydrolyzing NAD.
Priming of innate antimycobacterial immunity by heat-killed Listeria monocytogenes induces sterilizing response in the adult zebrafish tuberculosis model.
Targeting the phosphoserine phosphatase MtSerB2 for tuberculosis drug discovery, an hybrid knowledge based /fragment based approach.
The $\alpha/\beta$ Hydrolase Fold Proteins of Mycobacterium tuberculosis, with Reference to their Contribution to Virulence.
Genomic analysis of a Mycobacterium bovis bacillus [corrected] Calmette-Guérin strain isolated from an adult patient with pulmonary tuberculosis.
The mycobacterial cell envelope - a moving target.
Mathematical model of mycobacterium-host interaction describes physiology of persistence.
Biochemical characterization of phosphoserine phosphatase SerB2 from Mycobacterium marinum.
Sequence diversity in the pe_pgrs genes of Mycobacterium tuberculosis is independent of human T cell recognition.
Contrasting persistence strategies in Salmonella and Mycobacterium.
The open pan-genome architecture and virulence landscape of Mycobacterium bovis.

Mycobacterium tuberculosis 19-kDa lipoprotein inhibits IFN-gamma-induced chromatin remodeling of MHC2TA by TLR2 and MAPK signaling.
Mycobacterium tuberculosis lipoprotein LprG binds lipoarabinomannan and determines its cell envelope localization to control phagolysosomal fusion.
Host immunity increases Mycobacterium tuberculosis reliance on cytochrome bd oxidase.
The Mycobacterium tuberculosis PE_PGRS Protein Family Acts as an Immunological Decoy to Subvert Host Immune Response.
Highlights on the Application of Genomics and Bioinformatics in the Fight Against Infectious Diseases: Challenges and Opportunities in Africa.
E3 Ligase FBXW7 Facilitates Mycobacterium Immune Evasion by Modulating TNF- $\alpha$ Expression.
Immune Evasion of Mycoplasma bovis.
RIP3 impedes Mycobacterium tuberculosis survival and promotes p62-mediated autophagy.
Conformational Switch Driven Membrane Pore Formation by Mycobacterium Secretory Protein MPT63 Induces Macrophage Cell Death.
TB: screening for responses to a vile visitor.
Variation of the Mycobacterium tuberculosis PE_PGRS 33 gene among clinical isolates.
Kinetics of chemokine secretion in human macrophages infected with various strains of Mycobacterium tuberculosis.
A paucity of knowledge regarding nontuberculous mycobacterial lipids compared to the tubercle bacillus.
Different macrophage polarization between drug-susceptible and multidrug-resistant pulmonary tuberculosis.
Mycobacterial shuttle vectors designed for high-level protein expression in infected macrophages.
Lessons from Bacillus Calmette-Guérin: Harnessing Trained Immunity for Vaccine Development.
Modern lineages of Mycobacterium tuberculosis exhibit lineage-specific patterns of growth and cytokine induction in human monocyte-derived macrophages.
Functional polymorphisms in CYP2C19 & CYP3A5 genes associated with decreased susceptibility for paediatric tuberculosis.
Mycobacterial mycolic acids trigger inhibitory receptor Clec12A to suppress host immune responses.
Molecular Basis Underlying Host Immunity Subversion by Mycobacterium tuberculosis PE/PPE Family Molecules.
The PE/PPE multigene family codes for virulence factors and is a possible source of mycobacterial antigenic variation: perhaps more?
PPE38 of Mycobacterium marinum triggers the cross-talk of multiple pathways involved in the host response, as revealed by subcellular quantitative proteomics.
Mycobacterium tuberculosis Virulent Factor ESAT-6 Drives Macrophage Differentiation Toward the Pro-inflammatory M1 Phenotype and Subsequently Switches It to the Anti-inflammatory M2 Phenotype.
Manipulation of costimulatory molecules by intracellular pathogens: veni, vidi, vici!!
New insights on the biology of swine respiratory tract mycoplasmas from a comparative genome analysis.
Reduction of host cell mitochondrial activity as Mycobacterium leprae's strategy to evade host innate immunity.
$\alpha$ -Glucan biosynthesis and the GlgE pathway in Mycobacterium tuberculosis.
Identification of a novel role of ESAT-6-dependent miR-155 induction during infection of macrophages with Mycobacterium tuberculosis.

Evasion of peptide, but not lipid antigen presentation, through pathogen-induced dendritic cell maturation.
Long-term survival of <i>Mycoplasma bovis</i> in necrotic lesions and in phagocytic cells as demonstrated by transmission and immunogold electron microscopy in lung tissue from experimentally infected calves.
Expanded polyfunctional T cell response to mycobacterial antigens in TB disease and contraction post-treatment.
MicroRNA-20a-3p regulates the host immune response to facilitate the mycobacterium tuberculosis infection by targeting IKK $\beta$ /NF- $\kappa$ B pathway.
Mitochondrial Transcription Factor A Regulates <i>Mycobacterium bovis</i> -Induced IFN- $\beta$ Production by Modulating Mitochondrial DNA Replication in Macrophages.
<i>Mycobacterium leprae</i> phenolglycolipid-1 expressed by engineered <i>M. bovis</i> BCG modulates early interaction with human phagocytes.
Mass Spectrometry Offers Insight into the Role of Ser/Thr/Tyr Phosphorylation in the Mycobacteria.
Macrophage-T cell interaction in experimental mycobacterial infection. Selective regulation of co-stimulatory molecules on <i>Mycobacterium</i> -infected macrophages and its implication in the suppression of cell-mediated immune response.
Dressed not to kill: CD16+ monocytes impair immune defence against tuberculosis.
Nontuberculous Mycobacteria Show Differential Infectivity and Use Phospholipids to Antagonize LL-37.
Pathogenesis and Virulence of <i>Mycoplasma bovis</i> .
Temporal effects of tumor necrosis factor-alpha on intracellular survival of <i>Mycobacterium paratuberculosis</i> .
Unphosphorylated STAT1 represses apoptosis in macrophages during <i>Mycobacterium tuberculosis</i> infection.
Chronic pneumonia in calves after experimental infection with <i>Mycoplasma bovis</i> strain 1067: characterization of lung pathology, persistence of variable surface protein antigens and local immune response.
The major membrane nuclease MnuA degrades neutrophil extracellular traps induced by <i>Mycoplasma bovis</i> .
<i>Mycobacterium smegmatis</i> Resists the Bactericidal Activity of Hypochlorous Acid Produced in Neutrophil Phagosomes.
Nitric oxide and KLF4 protein epigenetically modify class II transactivator to repress major histocompatibility complex II expression during <i>Mycobacterium bovis</i> bacillus Calmette-Guerin infection.
PET/CT imaging of CSF1R in a mouse model of tuberculosis.
Vaccination with recombinant <i>Mycoplasma bovis</i> GAPDH results in a strong humoral immune response but does not protect feedlot cattle from an experimental challenge with <i>M. bovis</i> .
Independent loss of immunogenic proteins in <i>Mycobacterium ulcerans</i> suggests immune evasion.
Multimodal nanoparticles that provide immunomodulation and intracellular drug delivery for infectious diseases.
<i>Mycoplasma Bovis</i> adhesins and their target proteins.
Inhibition of maturation of human monocyte-derived dendritic cells in a patient with <i>Mycobacterium avium</i> infection.
<i>Mycobacterial</i> immunevasion-Spotlight on the enemy within.
A potential role for complement in immune evasion by <i>Mycobacterium leprae</i> .
T-Regulatory Cells as Part of Strategy of Immune Evasion by Pathogens.

Disruption of immune regulation by microbial pathogens and resulting chronic inflammation.
Identification of a Mycobacterium tuberculosis Cyclic Dinucleotide Phosphodiesterase Inhibitor.
Metabolic Network for the Biosynthesis of Intra- and Extracellular $\alpha$ -Glucans Required for Virulence of Mycobacterium tuberculosis.
Increased survival and proliferation of the epidemic strain Mycobacterium abscessus subsp. massiliense CRM0019 in alveolar epithelial cells.
Long-term persistence of BCG Pasteur in lungs of C57BL/6 mice following intranasal infection.
Virtual screening and crystallographic studies reveal an unexpected $\gamma$ -lactone derivative active against MptpB as a potential antitubercular agent.
The M protein of group A Streptococcus is a key virulence factor and a clinically relevant strain identification marker.
Anti-inflammatory pathways as a host evasion mechanism for pathogens.
Associating H(2)O(2-) and NO-related changes in the proteome of Mycobacterium smegmatis with enhanced survival in macrophage.
Haemagglutinins of pathogenic avian mycoplasmas.
TREM2 is a receptor for non-glycosylated mycolic acids of mycobacteria that limits anti-mycobacterial macrophage activation.
Autophagy Is an Innate Mechanism Associated with Leprosy Polarization.
Partners in Crime: Phenolic Glycolipids and Macrophages.
Revisiting the role of mesenchymal stem cells in tuberculosis and other infectious diseases.
A role for glycogen synthase kinase-3 in antagonizing mycobacterial immune evasion by negatively regulating IL-10 induction.
On deaf ears, Mycobacterium avium paratuberculosis in pathogenesis Crohn's and other diseases.
Crystal structure of the TreS:Pep2 complex, initiating $\alpha$ -glucan synthesis in the GlgE pathway of mycobacteria.
Mycobacterium tuberculosis is protected from NADPH oxidase and LC3-associated phagocytosis by the LCP protein CpsA.
Mycobacterium avium infection of epithelial cells results in inhibition or delay in the release of interleukin-8 and RANTES.
Erythema Nodosum Leprosum Neutrophil Subset Expressing IL-10R1 Transmigrates into Skin Lesions and Responds to IL-10.
Mycoplasma haemocanis--the canine hemoplasma and its feline counterpart in the genomic era.
Looking within the zebrafish to understand the tuberculous granuloma.
Dendritic cell-based vaccination against cancer.
Degradation-resistant protein domains limit host cell processing and immune detection of mycobacteria.
Mycobacterium abscessus cording prevents phagocytosis and promotes abscess formation.
Proteome-scale identification and characterization of mitochondria targeting proteins of Mycobacterium avium subspecies paratuberculosis: Potential virulence factors modulating host mitochondrial function.
PPE38 modulates the innate immune response and is required for Mycobacterium marinum virulence.
Essential Roles of PPARs in Lipid Metabolism during Mycobacterial Infection.
Mycobacterium avium reduces expression of costimulatory/adhesion molecules by human monocytes.
microRNAs in mycobacterial disease: friend or foe?
Causation of Crohn's disease by Mycobacterium avium subspecies paratuberculosis.

Modulation of the host immune response by a transient intracellular stage of <i>Mycobacterium ulcerans</i> : the contribution of endogenous mycolactone toxin.
Analysis of mRNA and circRNA Expression Profiles of Bovine Monocyte-Derived Macrophages Infected With <i>Mycobacterium avium</i> subsp. <i>paratuberculosis</i> .
Hiding behind the mycobacterial cell wall.
Modulation of innate host factors by <i>Mycobacterium avium</i> complex in human macrophages includes interleukin 17.
<i>Leishmania donovani</i> Subverts Host Immune Response by Epigenetic Reprogramming of Macrophage M(Lipopolysaccharides + IFN- $\gamma$ )/M(IL-10) Polarization.
Rab7 controls lipid droplet-phagosome association during mycobacterial infection.
<i>Yersinia enterocolitica</i> Exploits Signal Crosstalk between Complement 5a Receptor and Toll-like Receptor 1/2 and 4 to Avoid the Bacterial Clearance in M cells.
<i>Mycobacterium avium</i> Subspecies <i>paratuberculosis</i> Infects and Replicates within Human Monocyte-Derived Dendritic Cells.
<i>Mycoplasma hyopneumoniae</i> evades phagocytic uptake by porcine alveolar macrophages in vitro.
Gene expression profiles of immune-regulatory genes in whole blood of cattle with a subclinical infection of <i>Mycobacterium avium</i> subsp. <i>paratuberculosis</i> .
[Immunopathology of mycobacterial infection].
Marked Differences in Mucosal Immune Responses Induced in Ileal versus Jejunal Peyer's Patches to <i>Mycobacterium avium</i> subsp. <i>paratuberculosis</i> Secreted Proteins following Targeted Enteric Infection in Young Calves.
Accelerating next-generation vaccine development for global disease prevention.
<i>Mycobacterium avium</i> subsp. <i>paratuberculosis</i> Proteome Changes Profoundly in Milk.
Altered microRNA expression and pre-mRNA splicing events reveal new mechanisms associated with early stage <i>Mycobacterium avium</i> subspecies <i>paratuberculosis</i> infection.
<i>Mycobacterium lepromatosis</i> genome exhibits unusually high CpG dinucleotide content and selection is key force in shaping codon usage.
Manuka honey in combination with azithromycin shows potential for improved activity against <i>Mycobacterium abscessus</i> .
Mycobacterial protein tyrosine kinase, PtkA phosphorylates PtpA at tyrosine residues and the mechanism is stalled by the novel series of inhibitors.
The m(6)A-Binding Protein YTHDF1 Mediates Immune Evasion.
Non-carbohydrate inhibitors of the lectin DC-SIGN.
Binding Sites for Acylated Trehalose Analogs of Glycolipid Ligands on an Extended Carbohydrate Recognition Domain of the Macrophage Receptor Mincle.
Paracellular Pathway-Mediated <i>Mycoplasma hyopneumoniae</i> Migration across Porcine Airway Epithelial Barrier under Air-Liquid Interface Conditions.
The etiologic role of infectious antigens in sarcoidosis pathogenesis.
<i>Mycobacterium avium</i> subspecies <i>paratuberculosis</i> infection in cattle - a review in the context of seasonal pasture-based dairy herds.
Expression of NLRP3 inflammasome in leprosy indicates immune evasion of <i>Mycobacterium leprae</i> .
Genetic Variation in Toll-Interacting Protein Is Associated With Leprosy Susceptibility and Cutaneous Expression of Interleukin 1 Receptor Antagonist.
[ESAT-6 inhibits autophagy in macrophages and promotes the growth of BCG].
Assembly of $\alpha$ -Glucan by GlgE and GlgB in <i>Mycobacteria</i> and <i>Streptomyces</i> .
Host subspecific viral strains in European house mice: Murine cytomegalovirus in the Eastern ( <i>Mus musculus musculus</i> ) and Western house mouse ( <i>Mus musculus domesticus</i> ).

Table A2-29, Cluster 28

Cluster 28 focuses on myeloid-derived suppressor cells in the tumor microenvironment, especially immunotherapy that targets these cells and reduces their suppression of T-cell activity (471)
Immunotherapy Targeting Myeloid-Derived Suppressor Cells (MDSCs) in Tumor Microenvironment.
Myeloid-Derived Suppressor Cells in the Tumor Microenvironment.
Expansion and functions of myeloid-derived suppressor cells in the tumor microenvironment.
Myeloid-derived suppressor cells in cancer patients: a clinical perspective.
Myeloid derived suppressor cells in physiological and pathological conditions: the good, the bad, and the ugly.
Targeting myeloid-derived suppressor cells for cancer immunotherapy.
Insights into Myeloid-Derived Suppressor Cells in Inflammatory Diseases.
Suppression of T cells by myeloid-derived suppressor cells in cancer.
Negative regulation of myeloid-derived suppressor cells in cancer.
On the origin of myeloid-derived suppressor cells.
Myeloid-derived suppressor cells in cancer: therapeutic, predictive, and prognostic implications.
The clinical evidence for targeting human myeloid-derived suppressor cells in cancer patients.
The New Era of Cancer Immunotherapy: Targeting Myeloid-Derived Suppressor Cells to Overcome Immune Evasion.
Myeloid derived suppressor cells-An overview of combat strategies to increase immunotherapy efficacy.
Mechanisms Underlying the Role of Myeloid-Derived Suppressor Cells in Clinical Diseases: Good or Bad.
Myeloid-derived suppressor cells: The green light for myeloma immune escape.
Myeloid-derived suppressor cells as immunosuppressive regulators and therapeutic targets in cancer.
Targeting Myeloid-Derived Suppressor Cells to Enhance the Antitumor Efficacy of Immune Checkpoint Blockade Therapy.
Roles of the Exosomes Derived From Myeloid-Derived Suppressor Cells in Tumor Immunity and Cancer Progression.
Myeloid-derived suppressor cells in cancer: recent progress and prospects.
Can the suppressive activity of myeloid-derived suppressor cells be "chop"ped?
Interleukin 33 in tumor microenvironment is crucial for the accumulation and function of myeloid-derived suppressor cells.
Nanoparticle Systems Modulating Myeloid-Derived Suppressor Cells for Cancer Immunotherapy.
Myeloid-derived suppressor cells-new and exciting players in lung cancer.
Immunotherapy of targeting MDSCs in tumor microenvironment.
The myeloid derived suppressor cells: Who are they? Can they be used as a diagnostic tool to investigate metastasis in veterinary medicine?
Understanding the Differentiation, Expansion, Recruitment and Suppressive Activities of Myeloid-Derived Suppressor Cells in Cancers.
Targeting myeloid-derived suppressor cells in tumor immunotherapy: Current, future and beyond.
Myeloid-derived suppressor cells in cancer immunotherapy-clinical perspectives.
Interaction Between microRNAs and Myeloid-Derived Suppressor Cells in Tumor Microenvironment.
Myeloid-derived suppressor cells: an emerging target for anticancer immunotherapy.
Myeloid-derived suppressor cells and tumor: Current knowledge and future perspectives.



Curdlan blocks the immune suppression by myeloid-derived suppressor cells and reduces tumor burden.
Myeloid-derived suppressor cells: the dark knight or the joker in viral infections?
Chemokines and their receptors promoting the recruitment of myeloid-derived suppressor cells into the tumor.
Pancreatic adenocarcinoma up-regulated factor (PAUF) enhances the accumulation and functional activity of myeloid-derived suppressor cells (MDSCs) in pancreatic cancer.
History of myeloid-derived suppressor cells.
Myeloid-derived suppressor cells in multiple myeloma: pre-clinical research and translational opportunities.
Transcriptomic Analyses of Myeloid-Derived Suppressor Cell Subsets in the Circulation of Colorectal Cancer Patients.
Myeloid-derived suppressor cells (MDSCs) in gliomas and glioma-development.
Myeloid-Derived Suppressor Cells: A New and Pivotal Player in Colorectal Cancer Progression.
Myeloid derived suppressor cells in chronic myeloid leukemia.
Roles of HMGB1 in regulating myeloid-derived suppressor cells in the tumor microenvironment.
Immune suppression: the hallmark of myeloid derived suppressor cells.
Myeloid-derived suppressor cells: Key immunosuppressive regulators and therapeutic targets in cancer.
Immunosuppressive Role of Myeloid-Derived Suppressor Cells and Therapeutic Targeting in Lung Cancer.
Myeloid-derived suppressor cells in B cell malignancies.
Circulating Myeloid-Derived Suppressor Cell Subsets in Patients with Colorectal Cancer - Exploratory Analysis of Their Biomarker Potential.
Epithelial-mesenchymal transition: When tumor cells meet myeloid-derived suppressor cells.
Circulating and tumor-infiltrating arginase 1-expressing cells in gastric adenocarcinoma patients were mainly immature and monocytic Myeloid-derived suppressor cells.
Hepatic myeloid-derived suppressor cells in cancer.
Myeloid-Derived Suppressor Cells in the Tumor Microenvironment: Current Knowledge and Future Perspectives.
TGF- $\beta$ Enhances the Anti-inflammatory Effect of Tumor- Infiltrating CD33+11b+HLA-DR Myeloid-Derived Suppressor Cells in Gastric Cancer: A Possible Relation to MicroRNA-494.
Urokinase-mediated recruitment of myeloid-derived suppressor cells and their suppressive mechanisms are blocked by MUC1/sec.
Myeloid-derived suppressor cells contribute to oral cancer progression in 4NQO-treated mice.
Myeloid-derived suppressor cells in hematologic malignancies: two sides of the same coin.
Transcriptomic profiling disclosed the role of DNA methylation and histone modifications in tumor-infiltrating myeloid-derived suppressor cell subsets in colorectal cancer.
Regulating Tumor Myeloid-Derived Suppressor Cells by MicroRNAs.
Phenotypic and transcriptomic characterization of canine myeloid-derived suppressor cells.
Myeloid-derived suppressor cells in lymphoma: The good, the bad and the ugly.
The Development and Homing of Myeloid-Derived Suppressor Cells: From a Two-Stage Model to a Multistep Narrative.
Myeloid-derived suppressor cells and the efficacy of CD8(+) T-cell immunotherapy.
IL-33 inhibits the differentiation and immunosuppressive activity of granulocytic myeloid-derived suppressor cells in tumor-bearing mice.

IL-18 enhances immunosuppressive responses by promoting differentiation into monocytic myeloid-derived suppressor cells.
The growing diversity and spectrum of action of myeloid-derived suppressor cells.
Monocytic CCR2(+) myeloid-derived suppressor cells promote immune escape by limiting activated CD8 T-cell infiltration into the tumor microenvironment.
Myeloid-derived suppressor cells.
Hypoxia inducible factor HIF-1 promotes myeloid-derived suppressor cells accumulation through ENTPD2/CD39L1 in hepatocellular carcinoma.
HMGB1 enhances immune suppression by facilitating the differentiation and suppressive activity of myeloid-derived suppressor cells.
Myeloid-derived suppressor cells are essential partners for immune checkpoint inhibitors in the treatment of cisplatin-resistant bladder cancer.
Circulating myeloid-derived suppressor cells: An independent prognostic factor in patients with breast cancer.
Myeloid-derived suppressor cells and anti-tumor T cells: a complex relationship.
Breaking bad habits: Targeting MDSCs to alleviate immunosuppression in prostate cancer.
Immunosuppressive myeloid-derived suppressor cells are increased in splenocytes from cancer patients.
Antigen specificity of immune suppression by myeloid-derived suppressor cells.
Myeloid-derived suppressor cell role in tumor-related inflammation.
Myeloid-derived suppressor cells: natural regulators for transplant tolerance.
Renal cancer-derived exosomes induce tumor immune tolerance by MDSCs-mediated antigen-specific immunosuppression.
Long non-coding RNA RUNXOR accelerates MDSC-mediated immunosuppression in lung cancer.
Glioblastoma Myeloid-Derived Suppressor Cell Subsets Express Differential Macrophage Migration Inhibitory Factor Receptor Profiles That Can Be Targeted to Reduce Immune Suppression.
Immunosuppression by monocytic myeloid-derived suppressor cells in patients with pancreatic ductal carcinoma is orchestrated by STAT3.
Myeloid suppressor cells and immune modulation in lung cancer.
The liver is a site for tumor-induced myeloid-derived suppressor cell accumulation and immunosuppression.
AMPK Alpha-1 Intrinsically Regulates the Function and Differentiation of Tumor Myeloid-Derived Suppressor Cells.
Circulating myeloid-derived suppressor cells in patients with pancreatic cancer.
TMPRSS11D/ALR-mediated ER stress regulates the function of myeloid-derived suppressor cells in the cervical cancer microenvironment.
Granulocytic Myeloid-Derived Suppressor Cells as Negative Regulators of Anticancer Immunity.
IRF7 regulates the development of granulocytic myeloid-derived suppressor cells through S100A9 transrepression in cancer.
Targeting Myeloid-Derived Suppressor Cells to Bypass Tumor-Induced Immunosuppression.
Cancer-cell-intrinsic mechanisms regulate MDSCs through cytokine networks.
Characterization of myeloid-derived suppressor cells and cytokines GM-CSF, IL-10 and MCP-1 in dogs with malignant melanoma receiving a GD3-based immunotherapy.
TLR9-Targeted STAT3 Silencing Abrogates Immunosuppressive Activity of Myeloid-Derived Suppressor Cells from Prostate Cancer Patients.
Avoiding phagocytosis-related artifact in myeloid derived suppressor cell T-lymphocyte suppression assays.

Generation of a new therapeutic peptide that depletes myeloid-derived suppressor cells in tumor-bearing mice.
The immune suppressive tumor microenvironment in multiple myeloma: The contribution of myeloid-derived suppressor cells.
Long Non-coding RNAs: Regulators of the Activity of Myeloid-Derived Suppressor Cells.
Critical immunosuppressive effect of MDSC-derived exosomes in the tumor microenvironment.
Comments on the ambiguity of selected surface markers, signaling pathways and omics profiles hampering the identification of myeloid-derived suppressor cells.
Mechanisms overseeing myeloid-derived suppressor cell production in neoplastic disease.
Distinct Myeloid Derived Suppressor Cell Populations Promote Tumor Aggression in Glioblastoma.
Increased frequency and clinical significance of myeloid-derived suppressor cells in human colorectal carcinoma.
Phytochemicals inhibit the immunosuppressive functions of myeloid-derived suppressor cells (MDSC): Impact on cancer and age-related chronic inflammatory disorders.
Aptamer-mediated blockade of IL4R $\alpha$ triggers apoptosis of MDSCs and limits tumor progression.
Tumor-induced tolerance and immune suppression by myeloid derived suppressor cells.
Knowing the myeloid-derived suppressor cells: Another enemy of sarcomas patients.
Clinical perspectives on targeting of myeloid derived suppressor cells in the treatment of cancer.
Myeloid-derived suppressor cells and tumor escape from immune surveillance.
Transforming growth factor-beta1 and myeloid-derived suppressor cells: A cancerous partnership.
Myeloid-Derived Suppressor Cells and Pancreatic Cancer: Implications in Novel Therapeutic Approaches.
Pseudoneutrophil Cytokine Sponges Disrupt Myeloid Expansion and Tumor Trafficking to Improve Cancer Immunotherapy.
Visualization and quantification of in vivo homing kinetics of myeloid-derived suppressor cells in primary and metastatic cancer.
The downregulation of type I IFN signaling in G-MDSCs under tumor conditions promotes their development towards an immunosuppressive phenotype.
Myeloid-Derived Suppressor Cells Promote Metastasis in Breast Cancer After the Stress of Operative Removal of the Primary Cancer.
Role of myeloid-derived suppressor cells in tumor recurrence.
Novel and enhanced anti-melanoma DNA vaccine targeting the tyrosinase protein inhibits myeloid-derived suppressor cells and tumor growth in a syngeneic prophylactic and therapeutic murine model.
Ceramide activates lysosomal cathepsin B and cathepsin D to attenuate autophagy and induces ER stress to suppress myeloid-derived suppressor cells.
MMP-9-cleaved osteopontin isoform mediates tumor immune escape by inducing expansion of myeloid-derived suppressor cells.
Myeloid-derived suppressor cells and pulmonary hypertension.
Subsets of myeloid-derived suppressor cells in hepatocellular carcinoma express chemokines and chemokine receptors differentially.
Cancer cells induce immune escape via glycocalyx changes controlled by the telomeric protein TRF2.
TNF signaling drives myeloid-derived suppressor cell accumulation.
Therapeutic prospects of targeting myeloid-derived suppressor cells and immune checkpoints in cancer.
Serum amyloid A3 exacerbates cancer by enhancing the suppressive capacity of myeloid-derived suppressor cells via TLR2-dependent STAT3 activation.

Monocytic myeloid-derived suppressor cells home to tumor-draining lymph nodes via CCR2 and locally modulate the immune response.
Early Posttransplant Mobilization of Monocytic Myeloid-derived Suppressor Cell Correlates With Increase in Soluble Immunosuppressive Factors and Predicts Cancer in Kidney Recipients.
Expansion and activation of granulocytic, myeloid-derived suppressor cells in childhood precursor B cell acute lymphoblastic leukemia.
GM-CSF signalling blockade and chemotherapeutic agents act in concert to inhibit the function of myeloid-derived suppressor cells in vitro.
Ex vivo generation of myeloid-derived suppressor cells that model the tumor immunosuppressive environment in colorectal cancer.
Tumor microenvironment, histone modifications, and myeloid-derived suppressor cells.
CD38-Expressing Myeloid-Derived Suppressor Cells Promote Tumor Growth in a Murine Model of Esophageal Cancer.
Exosomes Derived from RM-1 Cells Promote the Recruitment of MDSCs into Tumor Microenvironment by Upregulating CXCR4 via TLR2/NF- $\kappa$ B Pathway.
Leukocyte immunoglobulin-like receptors in human diseases: an overview of their distribution, function, and potential application for immunotherapies.
The Functional Crosstalk between Myeloid-Derived Suppressor Cells and Regulatory T Cells within the Immunosuppressive Tumor Microenvironment.
Increase in myeloid-derived suppressor cells (MDSCs) associated with minimal residual disease (MRD) detection in adult acute myeloid leukemia.
Targeting myeloid-derived suppressor cells in combination with tumor cell vaccination predicts anti-tumor immunity and breast cancer dormancy: an in silico experiment.
Myeloid derived suppressor cells (MDSCs) are increased and exert immunosuppressive activity together with polymorphonuclear leukocytes (PMNs) in chronic myeloid leukemia patients.
Tumor-induced tolerance and immune suppression depend on the C/EBPbeta transcription factor.
Immune checkpoint blockade in triple negative breast cancer influenced by B cells through myeloid-derived suppressor cells.
Myeloid-derived suppressor cells and T regulatory cells in tumors: unraveling the dark side of the force.
Hydrogen Sulfide Reduces Myeloid-Derived Suppressor Cell-Mediated Inflammatory Response in a Model of Helicobacter hepaticus-Induced Colitis.
The role of myeloid-derived suppressor cells in liver cancer.
ATP/P2X7 axis modulates myeloid-derived suppressor cell functions in neuroblastoma microenvironment.
Myeloid-derived suppressor cells (MDSCs) in patients with solid tumors: considerations for granulocyte colony-stimulating factor treatment.
Interaction between invariant NKT cells and myeloid-derived suppressor cells in cancer patients: evidence and therapeutic opportunities.
Tumor- and organ-dependent infiltration by myeloid-derived suppressor cells.
Targeting myeloid-derived suppressor cells using a novel adenosine monophosphate-activated protein kinase (AMPK) activator.
Immunostimulatory RNA leads to functional reprogramming of myeloid-derived suppressor cells in pancreatic cancer.
IL-1 $\beta$ regulates a novel myeloid-derived suppressor cell subset that impairs NK cell development and function.

The potential importance of myeloid-derived suppressor cells (MDSCs) in the pathogenesis of Alzheimer's disease.
Arginine regulation by myeloid derived suppressor cells and tolerance in cancer: mechanisms and therapeutic perspectives.
Cross-talk between myeloid-derived suppressor cells (MDSC), macrophages, and dendritic cells enhances tumor-induced immune suppression.
Myeloid-derived suppressor cell and macrophage exert distinct angiogenic and immunosuppressive effects in breast cancer.
Transmembrane tumor necrosis factor- $\alpha$ promotes the recruitment of MDSCs to tumor tissue by upregulating CXCR4 expression via TNFR2.
Myeloid-derived suppressor cells deficient in cholesterol biosynthesis promote tumor immune evasion.
A Call for Epidemiological Research on Myeloid-Derived Suppressor Cells in Ovarian Cancer: A Review of the Existing Immunological Evidence and Suggestions for Moving Forward.
Doxorubicin eliminates myeloid-derived suppressor cells and enhances the efficacy of adoptive T-cell transfer in breast cancer.
Myeloid-derived suppressor cell accumulation in renal cell carcinoma is correlated with CCL2, IL-17 and IL-18 expression in blood and tumors.
Friend or Foe? Recent Strategies to Target Myeloid Cells in Cancer.
Human G-MDSCs are neutrophils at distinct maturation stages promoting tumor growth in breast cancer.
Role of myeloid-derived suppressor cells in metastasis.
Differential response of myeloid-derived suppressor cells to the nonsteroidal anti-inflammatory agent indomethacin in tumor-associated and tumor-free microenvironments.
Myeloid-Derived Suppressor Cells Impair B Cell Responses in Lung Cancer through IL-7 and STAT5.
Blockade of Myd88 signaling by a novel MyD88 inhibitor prevents colitis-associated colorectal cancer development by impairing myeloid-derived suppressor cells.
In vitro differentiation of myeloid suppressor cells (MDSC-like) from an immature myelomonocytic precursor THP-1.
Metformin inhibits the function of granulocytic myeloid-derived suppressor cells in tumor-bearing mice.
microRNAs Shape Myeloid Cell-Mediated Resistance to Cancer Immunotherapy.
The spleen contributes to the increase in PMN-MDSCs in orthotopic H22 hepatoma mice.
Increased frequencies of CD11b(+) CD33(+) CD14(+) HLA-DR(low) myeloid-derived suppressor cells are an early event in melanoma patients.
Quercetin promotes the survival of granulocytic myeloid-derived suppressor cells via the ESR2/STAT3 signaling pathway.
TGF- $\beta$ 1 programmed myeloid-derived suppressor cells (MDSC) acquire immune-stimulating and tumor killing activity capable of rejecting established tumors in combination with radiotherapy.
S100A9 Regulates MDSCs-Mediated Immune Suppression via the RAGE and TLR4 Signaling Pathways in Colorectal Carcinoma.
Peripheral blood myeloid-derived suppressor cells reflect disease status in idiopathic pulmonary fibrosis.
Sunitinib attenuates reactive MDSCs enhancing anti-tumor immunity in HNSCC.
Blockade of CCR5-mediated myeloid derived suppressor cell accumulation enhances anti-PD1 efficacy in gastric cancer.

Targeting myeloid derived suppressor cells with all-trans retinoic acid is highly time-dependent in therapeutic tumor vaccination.
Myeloid derived suppressor cells and the release of micro-metastases from dormancy.
Metabolic and functional reprogramming of myeloid-derived suppressor cells and their therapeutic control in glioblastoma.
Cross-talk between myeloid-derived suppressor cells and macrophages subverts tumor immunity toward a type 2 response.
The stress-response sensor chop regulates the function and accumulation of myeloid-derived suppressor cells in tumors.
Cancer Stem Cell-Secreted Macrophage Migration Inhibitory Factor Stimulates Myeloid Derived Suppressor Cell Function and Facilitates Glioblastoma Immune Evasion.
[Construction and anti-tumor efficacy of a pentameric peptide vaccine that targets S100A8].
Myeloid-Derived Suppressor Cell Survival and Function Are Regulated by the Transcription Factor Nrf2.
Aberrant PGE <sub>2</sub> metabolism in bladder tumor microenvironment promotes immunosuppressive phenotype of tumor-infiltrating myeloid cells.
Low-dose HDACi potentiates anti-tumor activity of macrophages in immunotherapy.
Inhibition of IL-18-mediated myeloid derived suppressor cell accumulation enhances anti-PD1 efficacy against osteosarcoma cancer.
Implications of MDSCs-targeting in lung cancer chemo-immunotherapeutics.
The Metabolic Control of Myeloid Cells in the Tumor Microenvironment.
Arginase I-producing myeloid-derived suppressor cells in renal cell carcinoma are a subpopulation of activated granulocytes.
Pak2 regulates myeloid-derived suppressor cell development in mice.
Human Head and Neck Squamous Cell Carcinoma-Associated Semaphorin 4D Induces Expansion of Myeloid-Derived Suppressor Cells.
Metabolic changes and interaction of tumor cell, myeloid-derived suppressor cell and T cell in hypoxic microenvironment.
Myeloid-Derived Suppressor Cells Endow Stem-like Qualities to Breast Cancer Cells through IL6/STAT3 and NO/NOTCH Cross-talk Signaling.
Polysaccharide Agaricus blazei Murill stimulates myeloid derived suppressor cell differentiation from M2 to M1 type, which mediates inhibition of tumour immune-evasion via the Toll-like receptor 2 pathway.
Cyclophosphamide-induced myeloid-derived suppressor cell population is immunosuppressive but not identical to myeloid-derived suppressor cells induced by growing TC-1 tumors.
Chronic restraint stress promotes the mobilization and recruitment of myeloid-derived suppressor cells through $\beta$ -adrenergic-activated CXCL5-CXCR2-Erk signaling cascades.
Decitabine shows potent anti-myeloma activity by depleting monocytic myeloid-derived suppressor cells in the myeloma microenvironment.
Hierarchy of immunosuppressive strength among myeloid-derived suppressor cell subsets is determined by GM-CSF.
Blockade of CCL2 enhances immunotherapeutic effect of anti-PD1 in lung cancer.
MDSC as a mechanism of tumor escape from sunitinib mediated anti-angiogenic therapy.
Immunotherapeutic modulation of the suppressive liver and tumor microenvironments.
Myeloid-derived suppressor cells promote epithelial ovarian cancer cell stemness by inducing the CSF2/p-STAT3 signalling pathway.
Myeloid cells' evasion of melanoma immunity.

CpG-mediated modulation of MDSC contributes to the efficacy of Ad5-TRAIL therapy against renal cell carcinoma.
STAT3 inhibition induces Bax-dependent apoptosis in liver tumor myeloid-derived suppressor cells.
Targeting tumor-associated macrophages and granulocytic myeloid-derived suppressor cells augments PD-1 blockade in cholangiocarcinoma.
Oncolytic Reovirus Inhibits Immunosuppressive Activity of Myeloid-Derived Suppressor Cells in a TLR3-Dependent Manner.
Melanoma-induced immunosuppression and its neutralization.
Biophysical heterogeneity of myeloid-derived microenvironment to regulate resistance to cancer immunotherapy.
Infiltration of alternatively activated macrophages in cancer tissue is associated with MDSC and Th2 polarization in patients with esophageal cancer.
Regulatory T Cells in the Tumor Microenvironment and Cancer Progression: Role and Therapeutic Targeting.
Tumor-infiltrating myeloid cells induce tumor cell resistance to cytotoxic T cells in mice.
PI3K $\gamma$ Activates Integrin $\alpha(4)$ and Promotes Immune Suppressive Myeloid Cell Polarization during Tumor Progression.
HMGB1 promotes myeloid-derived suppressor cells and renal cell carcinoma immune escape.
Common extracellular matrix regulation of myeloid cell activity in the bone marrow and tumor microenvironments.
Targeting glutamine metabolism enhances tumor-specific immunity by modulating suppressive myeloid cells.
Prim-O-glucosylcimifugin enhances the antitumour effect of PD-1 inhibition by targeting myeloid-derived suppressor cells.
Targeting myeloid cells in the tumor microenvironment enhances vaccine efficacy in murine epithelial ovarian cancer.
Fc $\gamma$ RIIB potentiates differentiation of myeloid-derived suppressor cells to mediate tumor immunoescape.
Timing of CSF-1/CSF-1R signaling blockade is critical to improving responses to CTLA-4 based immunotherapy.
LMP1-mediated glycolysis induces myeloid-derived suppressor cell expansion in nasopharyngeal carcinoma.
Melanoma induced immunosuppression is mediated by hematopoietic dysregulation.
Targeting KIT on innate immune cells to enhance the antitumor activity of checkpoint inhibitors.
Targeting and exploitation of tumor-associated neutrophils to enhance immunotherapy and drug delivery for cancer treatment.
Fatty Acid Metabolism in Myeloid-Derived Suppressor Cells and Tumor-Associated Macrophages: Key Factor in Cancer Immune Evasion.
Polymorphonuclear MDSCs are enriched in the stroma and expanded in metastases of prostate cancer.
Gemcitabine directly inhibits myeloid derived suppressor cells in BALB/c mice bearing 4T1 mammary carcinoma and augments expansion of T cells from tumor-bearing mice.
Mechanism of all-trans retinoic acid effect on tumor-associated myeloid-derived suppressor cells.
Tetraspanin CD81 promotes tumor growth and metastasis by modulating the functions of T regulatory and myeloid-derived suppressor cells.
Global stability and parameter analysis reinforce therapeutic targets of PD-L1-PD-1 and MDSCs for glioblastoma.

Palmitoylated Proteins on AML-Derived Extracellular Vesicles Promote Myeloid-Derived Suppressor Cell Differentiation via TLR2/Akt/mTOR Signaling.
Targeted depletion of an MDSC subset unmasks pancreatic ductal adenocarcinoma to adaptive immunity.
Size-tunable Nanoregulator-Based Radiofrequency Ablation Suppresses MDSCs And Their Compensatory Immune Evasion in Hepatocellular Carcinoma.
The receptor for advanced glycation end products promotes pancreatic carcinogenesis and accumulation of myeloid-derived suppressor cells.
Myeloid-derived suppressor cells adhere to physiologic STAT3- vs STAT5-dependent hematopoietic programming, establishing diverse tumor-mediated mechanisms of immunologic escape.
Targeting MDSC for Immune-Checkpoint Blockade in Cancer Immunotherapy: Current Progress and New Prospects.
Myeloid-derived suppressor cell inhibition of the IFN response in tumor-bearing mice.
IL-6 regulates CCR5 expression and immunosuppressive capacity of MDSC in murine melanoma.
Poly (I: C) modulates the immunosuppressive activity of myeloid-derived suppressor cells in a murine model of breast cancer.
Increased circulating myeloid-derived suppressor cells correlate with clinical cancer stage, metastatic tumor burden, and doxorubicin-cyclophosphamide chemotherapy.
MDSC and TGF $\beta$ Are Required for Facilitation of Tumor Growth in the Lungs of Mice Exposed to Carbon Nanotubes.
Gangliosides drive the tumor infiltration and function of myeloid-derived suppressor cells.
Strategies to overcome myeloid cell induced immune suppression in the tumor microenvironment.
Does placental MDSC-mediated modulation of arginine levels help protect the foetus from auxotrophic pathogens?
Turn Back the TIME: Targeting Tumor Infiltrating Myeloid Cells to Revert Cancer Progression.
HDAC inhibition potentiates anti-tumor activity of macrophages and enhances anti-PD-L1-mediated tumor suppression.
Th1 polarization in the tumor microenvironment upregulates the myeloid-derived suppressor-like function of macrophages.
Phosphodiesterase-5 inhibition augments endogenous antitumor immunity by reducing myeloid-derived suppressor cell function.
Cancer cell-intrinsic XBP1 drives immunosuppressive reprogramming of intratumoral myeloid cells by promoting cholesterol production.
Myeloid derived suppressor and dendritic cell subsets are related to clinical outcome in prostate cancer patients treated with prostate GVAX and ipilimumab.
Metabolomic profiling of tumor-infiltrating macrophages during tumor growth.
Myeloid NEMO deficiency promotes tumor immunosuppression partly via MCP1-CCR2 axis.
ARID1A loss induces polymorphonuclear myeloid-derived suppressor cell chemotaxis and promotes prostate cancer progression.
Tolerance and immune suppression in the tumor microenvironment.
Myeloid-Derived Suppressor Cells as Target of Phosphodiesterase-5 Inhibitors in Host-Directed Therapeutics for Tuberculosis.
Flagellin induces myeloid-derived suppressor cells: implications for Pseudomonas aeruginosa infection in cystic fibrosis lung disease.
IRF8: Mechanism of Action and Health Implications.
Cancer Cells Resistance Shaping by Tumor Infiltrating Myeloid Cells.



Targeted Deletion of CXCR2 in Myeloid Cells Alters the Tumor Immune Environment to Improve Antitumor Immunity.
Tumor-Derived Myeloid Cell Chemoattractants and T Cell Exclusion in Pancreatic Cancer.
Human prosthetic joint infections are associated with myeloid-derived suppressor cells (MDSCs): Implications for infection persistence.
Effective combinatorial immunotherapy for castration-resistant prostate cancer.
Disruption of CXCR2-mediated MDSC tumor trafficking enhances anti-PD1 efficacy.
MUC1-induced immunosuppression in colon cancer can be reversed by blocking the PD1/PDL1 signaling pathway.
Obesity Promotes Tumor Immune Evasion in Ovarian Cancer Through Increased Production of Myeloid-Derived Suppressor Cells via IL-6.
A Nanoplatfrom to Amplify Apoptosis-to-Pyroptosis Immunotherapy via Immunomodulation of Myeloid-Derived Suppressor Cells.
Blockade of A2b adenosine receptor reduces tumor growth and immune suppression mediated by myeloid-derived suppressor cells in a mouse model of melanoma.
Tumor NLRP3-Derived IL-1 $\beta$ Drives the IL-6/STAT3 Axis Resulting in Sustained MDSC-Mediated Immunosuppression.
Transmembrane TNF- $\alpha$ promotes suppressive activities of myeloid-derived suppressor cells via TNFR2.
Cellular immune suppressor mechanisms in patients with hepatocellular carcinoma.
The impact of hypoxia on immune state in cancer.
Tumor-reactive immune cells protect against metastatic tumor and induce immunoediting of indolent but not quiescent tumor cells.
Immune microenvironment of gliomas.
Myeloid-derived suppressor cells and regulatory T cells share common immunoregulatory pathways-related microRNAs that are dysregulated by acute lymphoblastic leukemia and chemotherapy.
IL-12 triggers a programmatic change in dysfunctional myeloid-derived cells within mouse tumors.
Mesenchymal Stem Cells (MSC) Regulate Activation of Granulocyte-Like Myeloid Derived Suppressor Cells (G-MDSC) in Chronic Myeloid Leukemia Patients.
Role of immature myeloid cells in mechanisms of immune evasion in cancer.
Potential Novel Ovarian Cancer Treatment Targeting Myeloid-Derived Suppressor Cells.
TIE-2 Signaling Activation by Angiopoietin 2 On Myeloid-Derived Suppressor Cells Promotes Melanoma-Specific T-cell Inhibition.
Fibrocytes represent a novel MDSC subset circulating in patients with metastatic cancer.
Increased circulating GrMyeloid-derived suppressor cells correlated with tumor burden and survival in locally advanced cervical cancer patient.
If we build it they will come: targeting the immune response to breast cancer.
Oleate but not stearate induces the regulatory phenotype of myeloid suppressor cells.
Atovaquone Suppresses Triple-Negative Breast Tumor Growth by Reducing Immune-Suppressive Cells.
Glutamine Deprivation Promotes the Generation and Mobilization of MDSCs by Enhancing Expression of G-CSF and GM-CSF.
Targeting tumor-derived NLRP3 reduces melanoma progression by limiting MDSCs expansion.
L-glutamine is a key parameter in the immunosuppression phenomenon.
TNF Receptor-2 Facilitates an Immunosuppressive Microenvironment in the Liver to Promote the Colonization and Growth of Hepatic Metastases.

Glioma-derived CCL2 and CCL7 mediate migration of immune suppressive CCR2(+)/CX3CR1(+) M-MDSCs into the tumor microenvironment in a redundant manner.
Hypoxia-Induced VISTA Promotes the Suppressive Function of Myeloid-Derived Suppressor Cells in the Tumor Microenvironment.
Hepatic stellate cells promote tumor progression by enhancement of immunosuppressive cells in an orthotopic liver tumor mouse model.
Effect of tumor-derived cytokines and growth factors on differentiation and immune suppressive features of myeloid cells in cancer.
Prospects of combinatorial synthetic peptide vaccine-based immunotherapy against cancer.
Breaking immunotolerance of tumors: a new perspective for dendritic cell therapy.
Renal transplant recipients have elevated frequencies of circulating myeloid-derived suppressor cells.
CSF1/CSF1R blockade reprograms tumor-infiltrating macrophages and improves response to T-cell checkpoint immunotherapy in pancreatic cancer models.
LPS converts Gr-1(+)/CD115(+) myeloid-derived suppressor cells from M2 to M1 via P38 MAPK.
CD200 in CNS tumor-induced immunosuppression: the role for CD200 pathway blockade in targeted immunotherapy.
The addition of recombinant vaccinia HER2/neu to oncolytic vaccinia-GM-CSF given into the tumor microenvironment overcomes MDSC-mediated immune escape and systemic anergy.
Small molecule immunomodulation: the tumor microenvironment and overcoming immune escape.
CaMKK2 in myeloid cells is a key regulator of the immune-suppressive microenvironment in breast cancer.
Differential Targeting of Gr-MDSCs, T Cells and Prostate Cancer Cells by Dactolisib and Dasatinib.
Vascular endothelial growth factor-trap overcomes defects in dendritic cell differentiation but does not improve antigen-specific immune responses.
CD40 controls CXCR5-induced recruitment of myeloid-derived suppressor cells to gastric cancer.
Overcoming the Immunosuppressive Tumor Microenvironment in Multiple Myeloma.
Crosstalk between Tumor-Infiltrating Immune Cells and Cancer-Associated Fibroblasts in Tumor Growth and Immunosuppression of Breast Cancer.
Immune System Evasion as Hallmark of Melanoma Progression: The Role of Dendritic Cells.
Fas signal promotes lung cancer growth by recruiting myeloid-derived suppressor cells via cancer cell-derived PGE2.
Chromatin Regulator CHD1 Remodels the Immunosuppressive Tumor Microenvironment in PTEN-Deficient Prostate Cancer.
Coordinated regulation of myeloid cells by tumours.
Immune response in glioma's microenvironment.
Pancreatic cancer-associated stellate cells promote differentiation of myeloid-derived suppressor cells in a STAT3-dependent manner.
Myeloid Cells as Clinical Biomarkers for Immune Checkpoint Blockade.
Tumor-induced neurogenesis and immune evasion as targets of innovative anti-cancer therapies.
Multifaceted Roles of Chemokines and Chemokine Receptors in Tumor Immunity.
Unwelcome complement.
Checkpoint Inhibition in Head and Neck Cancer: Immune Therapeutic Options, Limitations, and Beyond.
The Role of Tumor-Stroma Interactions in Drug Resistance Within Tumor Microenvironment.
Knockdown of serine/threonine-protein kinase 24 promotes tumorigenesis and myeloid-derived suppressor cell expansion in an orthotopic immunocompetent gastric cancer animal model.

Cathepsin C promotes colorectal cancer metastasis by regulating immune escape through upregulating CSF1.
Immune consequences of penfluridol treatment associated with inhibition of glioblastoma tumor growth.
Extracellular S100A9 Protein in Bone Marrow Supports Multiple Myeloma Survival by Stimulating Angiogenesis and Cytokine Secretion.
Advances in radiotherapy and targeted therapies for rectal cancer.
Role of myeloid cells in tumor angiogenesis and growth.
Bone marrow myeloid cells in regulation of multiple myeloma progression.
Standard therapies versus novel therapies in Hodgkin lymphoma.
Glycan-Lectin Interactions as Novel Immunosuppression Drivers in Glioblastoma.
Microbiota modulation of myeloid cells in cancer therapy.
Role of the Immune Response in Disease Progression and Therapy in Multiple Myeloma.
Activities of stromal and immune cells in HPV-related cancers.
Deletion of p53 and Hyper-Activation of PIK3CA in Keratin-15(+) Stem Cells Lead to the Development of Spontaneous Squamous Cell Carcinoma.
Therapeutic activity of sunitinib for Her2/neu induced mammary cancer in FVB mice.
Cellular and molecular mechanisms in cancer immune escape: a comprehensive review.
Immune system: a double-edged sword in cancer.
Therapeutic Tumor Control of HER2 DNA Vaccines Is Achieved by an Alteration of Tumor Cells and Tumor Microenvironment by Gemcitabine and Anti-Gr-1 Ab Treatment in a HER2-Expressing Tumor Model.
HPV16 E2 protein promotes innate immunity by modulating immunosuppressive status.
Phenylboronic acid modified nanoparticles simultaneously target pancreatic cancer and its metastasis and alleviate immunosuppression.
Role of immature myeloid Gr-1+ cells in the development of antitumor immunity.
[Research Advance on the Immune Escape of Acute Myeloid Leukemia--Review].
Analyzing One Cell at a TIME: Analysis of Myeloid Cell Contributions in the Tumor Immune Microenvironment.
Capecitabine reverses tumor escape from anti-VEGF through the eliminating CD11b(high)/Gr1(high) myeloid cells.
Extracellular matrix and the myeloid-in-myeloma compartment: balancing tolerogenic and immunogenic inflammation in the myeloma niche.
Inhibition of arginase by CB-1158 blocks myeloid cell-mediated immune suppression in the tumor microenvironment.
Using chemo-drugs or irradiation to break immune tolerance and facilitate immunotherapy in solid cancer.
Perspectives of Immune Suppression in the Tumor Microenvironment Promoting Oral Malignancy.
Immune suppression in gliomas.
TLR9 expression and secretion of LIF by prostate cancer cells stimulates accumulation and activity of polymorphonuclear MDSCs.
Epithelial-to-mesenchymal Transition Heterogeneity of Circulating Tumor Cells and Their Correlation With MDSCs and Tregs in HER2-negative Metastatic Breast Cancer Patients.
Human papillomavirus-driven immune deviation: challenge and novel opportunity for immunotherapy.
Tumor-released microvesicles as vehicles of immunosuppression.
Bone Marrow Immune Microenvironment in Myelodysplastic Syndromes.

Myeloid-derived cells in prostate cancer progression: phenotype and prospective therapies.
Combination of a STAT3 inhibitor with anti-PD-1 immunotherapy is an effective treatment regimen for a vemurafenib-resistant melanoma.
Ferroptosis promotes anti-tumor immune response by inducing immunogenic exposure in HNSCC.
The role of myeloid cells in cancer therapies.
Cellular constituents of immune escape within the tumor microenvironment.
The Immunosuppressive Effect of TNFR2 Expression in the Colorectal Cancer Microenvironment.
IDO is a nodal pathogenic driver of lung cancer and metastasis development.
Regulation of tumor immune suppression and cancer cell survival by CXCL1/2 elevation in glioblastoma multiforme.
Splenic Hematopoietic and Stromal Cells in Cancer Progression.
Overcoming Hypoxia-Mediated Tumor Progression: Combinatorial Approaches Targeting pH Regulation, Angiogenesis and Immune Dysfunction.
Immunosuppressive parameters in serum of ovarian cancer patients change during the disease course.
A rationally designed combined treatment with an alphavirus-based cancer vaccine, sunitinib and low-dose tumor irradiation completely blocks tumor development.
Suppression of canine myeloid cells by soluble factors from cultured canine tumor cells.
DCLK1 Suppresses Tumor-Specific Cytotoxic T Lymphocyte Function Through Recruitment of MDSCs via the CXCL1-CXCR2 Axis.
Stromal Reprogramming through Dual PDGFR $\alpha$ / $\beta$ Blockade Boosts the Efficacy of Anti-PD-1 Immunotherapy in Fibrotic Tumors.
Prediction of anti-CD25 and 5-FU treatments efficacy for pancreatic cancer using a mathematical model.
Innate Immune Defense Mechanisms by Myeloid Cells That Hamper Cancer Immunotherapy.
Myeloid cells: Prime drivers of tumor progression.
Rapid deletion and inactivation of CTLs upon recognition of a number of target cells over a critical threshold.
The Yin-Yang of tumor-associated macrophages in neoplastic progression and immune surveillance.
Apl6/AlM/Sp $\alpha$ /CD5L overexpression in alveolar type II epithelial cells induces spontaneous lung adenocarcinoma.
Immunotherapy in Myeloproliferative Diseases.
Mapping the immunosuppressive environment in uterine tumors: implications for immunotherapy.
Interleukin-33 in tumorigenesis, tumor immune evasion, and cancer immunotherapy.
The tyrosine kinase inhibitors imatinib and dasatinib reduce myeloid suppressor cells and release effector lymphocyte responses.
Nitric oxide-producing myeloid-derived suppressor cells inhibit vascular E-selectin expression in human squamous cell carcinomas.
Immunological and biological dissection of normal and tumoral salivary glands.
Tumor-induced myeloid dysfunction and its implications for cancer immunotherapy.
Immunomodulation in cancer.
The Sabotaging Role of Myeloid Cells in Anti-Angiogenic Therapy: Coordination of Angiogenesis and Immune Suppression by Hypoxia.
Hydrogen-Peroxide Synthesis and LDL-Uptake Controls Immunosuppressive Properties in Monocyte-Derived Dendritic Cells.
Exploring the regulatory role of lncRNA in cancer immunity.

Intratumoral Hypoxia and Mechanisms of Immune Evasion Mediated by Hypoxia-Inducible Factors.
GOLM1 Drives Colorectal Cancer Metastasis by Regulating Myeloid-derived Suppressor Cells.
The emerging role of triggering receptor expressed on myeloid cell-2 in malignant tumor.
TNF Receptor 2 Makes Tumor Necrosis Factor a Friend of Tumors.
Cancer associated fibroblasts have phenotypic and functional characteristics similar to the fibrocytes that represent a novel MDSC subset.
A tumor-myeloid cell axis, mediated via the cytokines IL-1 $\alpha$ and TSLP, promotes the progression of breast cancer.
G9a promotes immune suppression by targeting the Fbxw7/Notch pathway in glioma stem cells.
Mild Hyperthermia Induced by Water-Filtered Infrared A Irradiation: A Potent Strategy to Foster Immune Recognition and Anti-Tumor Immune Responses in Superficial Cancers?
Adenosine-generating ovarian cancer cells attract myeloid cells which differentiate into adenosine-generating tumor associated macrophages - a self-amplifying, CD39- and CD73-dependent mechanism for tumor immune escape.
The Reversal of Immune Exclusion Mediated by Tadalafil and an Anti-tumor Vaccine Also Induces PDL1 Upregulation in Recurrent Head and Neck Squamous Cell Carcinoma: Interim Analysis of a Phase I Clinical Trial.
Emerging Role of Helicobacter pylori in the Immune Evasion Mechanism of Gastric Cancer: An Insight Into Tumor Microenvironment-Pathogen Interaction.
Podoplanin Positive Myeloid Cells Promote Glioma Development by Immune Suppression.
Epigenetic therapy regulates the expression of ALDH1 and immunologic response: Relevance to the prognosis of oral cancer.
Factors Influencing the Differentiation of Human Monocytic Myeloid-Derived Suppressor Cells Into Inflammatory Macrophages.
Mycobacterium tuberculosis exploits MPT64 to generate myeloid-derived suppressor cells to evade the immune system.
HDACi Delivery Reprograms Tumor-Infiltrating Myeloid Cells to Eliminate Antigen-Loss Variants.
Increased production of immature myeloid cells in cancer patients: a mechanism of immunosuppression in cancer.
Myeloid-derived suppressor cells (MDSC) facilitate distant metastasis of malignancies by shielding circulating tumor cells (CTC) from immune surveillance.
An Exploration of Oral-Gut Pathogens Mediating Immune Escape of Pancreatic Cancer via miR-21/PTEN Axis.
Tumor Immune Evasion Induced by Dysregulation of Erythroid Progenitor Cells Development.
CD38 and Regulation of the Immune Response Cells in Cancer.
Tumor-derived osteopontin suppresses antitumor immunity by promoting extramedullary myelopoiesis.
WDR5-H3K4me3 epigenetic axis regulates OPN expression to compensate PD-L1 function to promote pancreatic cancer immune escape.
Potential differentiation of tumor bearing mouse CD11b+Gr-1+ immature myeloid cells into both suppressor macrophages and immunostimulatory dendritic cells.
Heterogeneity in Immune Cell Content in Malignant Pleural Mesothelioma.
$\beta$ -glucan restores tumor-educated dendritic cell maturation to enhance antitumor immune responses.
Elevated lactate dehydrogenase (LDH) can be a marker of immune suppression in cancer: Interplay between hematologic and solid neoplastic clones and their microenvironments.
Glioma-derived galectin-1 regulates innate and adaptive antitumor immunity.

Preparation and functional analysis of tumor-infiltrating stroma cells using bone marrow chimera mice.
The Role of ERO1 $\alpha$ in Modulating Cancer Progression and Immune Escape.
All-trans-retinoic acid eliminates immature myeloid cells from tumor-bearing mice and improves the effect of vaccination.
The Immune Microenvironment in Human Papilloma Virus-Induced Cervical Lesions-Evidence for Estrogen as an Immunomodulator.
Immune checkpoints in hematologic malignancies: What made the immune cells and clinicians exhausted!
Upregulation of Atypical Cadherin FAT1 Promotes an Immunosuppressive Tumor Microenvironment via TGF- $\beta$ .
A Tumor Microenvironments-Adapted Polypeptide Hydrogel/Nanogel Composite Boosts Antitumor Molecularly Targeted Inhibition and Immunoactivation.
Kras-driven intratumoral heterogeneity triggers infiltration of M2 polarized macrophages via the circHIPK3/PTK2 immunosuppressive circuit.
From gatekeepers to providers: regulation of immune functions by cancer-associated fibroblasts.
Immunological Evasion in Glioblastoma.
Dickkopf-1: A Promising Target for Cancer Immunotherapy.
Minimal residual disease in advanced or metastatic solid cancers: The G0-G1 state and immunotherapy are key to unwinding cancer complexity.
CCL2/CCR2 pathway mediates recruitment of myeloid suppressor cells to cancers.
The DNA Methylcytosine Dioxygenase Tet2 Sustains Immunosuppressive Function of Tumor-Infiltrating Myeloid Cells to Promote Melanoma Progression.
Chronic Myeloid Leukemia: Immunobiology and Novel Immunotherapeutic Approaches.
SIRP $\alpha$ blockade improves the antitumor immunity of radiotherapy in colorectal cancer.
Effects of glioblastoma-derived extracellular vesicles on the functions of immune cells.
A novel probe for the non-invasive detection of tumor-associated inflammation.
Sunitinib-induced myeloid lineage redistribution in renal cell cancer patients: CD1c+ dendritic cell frequency predicts progression-free survival.
Immune microenvironment profiles of tumor immune equilibrium and immune escape states of mouse sarcoma.
Expansion of CCR8(+) inflammatory myeloid cells in cancer patients with urothelial and renal carcinomas.
Pharmacological Wnt ligand inhibition overcomes key tumor-mediated resistance pathways to anti-PD-1 immunotherapy.
Phenotypes, Functions, and Clinical Relevance of Regulatory B Cells in Cancer.
Heterogeneity of tumor-infiltrating myeloid cells in era of single-cell genomics.
KRAS-IRF2 Axis Drives Immune Suppression and Immune Therapy Resistance in Colorectal Cancer.
Effects of helicobacter pylori on tumor microenvironment and immunotherapy responses.
All-trans-retinoic acid improves differentiation of myeloid cells and immune response in cancer patients.
Arid5a Promotes Immune Evasion by Augmenting Tryptophan Metabolism and Chemokine Expression.
Inflammatory Microenvironment and Specific T Cells in Myeloproliferative Neoplasms: Immunopathogenesis and Novel Immunotherapies.
Monitoring dendritic cell and cytokine biomarkers during remission prior to relapse in patients with FLT3-ITD acute myeloid leukemia.

Potential of memory T cells in bridging preoperative chemoradiation and immunotherapy in rectal cancer.
Tumor-educated myeloid cells: impact the micro- and macroenvironment.
Blood monocytes from mammary tumor-bearing mice: early targets of tumor-induced immune suppression?
Persistent <i>Cryptosporidium parvum</i> Infection Leads to the Development of the Tumor Microenvironment in an Experimental Mouse Model: Results of a Microarray Approach.
Myeloid STAT3 promotes formation of colitis-associated colorectal cancer in mice.
Red Blood Cell Membrane as a Biomimetic Nanocoating for Prolonged Circulation Time and Reduced Accelerated Blood Clearance.
Overexpression of Arginase 1 is linked to DNMT3A and TET2 mutations in lower-grade myelodysplastic syndromes and chronic myelomonocytic leukemia.
De-O-acylated lipooligosaccharide of <i>E. coli</i> B reduces the number of metastatic foci via downregulation of myeloid cell activity.
Cancer makes new friends with old tricks.
Efficacy of intravesical Bacillus Calmette-Guérin therapy against tumor immune escape in an orthotopic model of bladder cancer.
Potent GCN2 Inhibitor Capable of Reversing MDSC-Driven T Cell Suppression Demonstrates In Vivo Efficacy as a Single Agent and in Combination with Anti-Angiogenesis Therapy.
Tumor-associated myeloid cells in cancer immunotherapy.
Saturated fatty acids dampen the immunogenicity of cancer by suppressing STING.
ARG1 functions as a tumor suppressor in breast cancer.
Induction of immunoregulatory CD271+ cells by metastatic tumor cells that express human endogenous retrovirus H.
Peroxisome proliferator-activated receptor gamma (PPARgamma) ligands reverse CTL suppression by alternatively activated (M2) macrophages in cancer.
IFITM6 expression is increased in macrophages of tumor-bearing mice.
Microglia/macrophages express alternative proangiogenic factors depending on granulocyte content in human glioblastoma.
Acute Myeloid Leukemia Presenting as Myeloid Sarcoma with a Predisposition to the Gynecologic Tract.
Transcriptomic profiling of peripheral blood cells in HPV-associated carcinoma patients receiving combined valproic acid and avelumab.
Lactylation-driven METTL3-mediated RNA m(6)A modification promotes immunosuppression of tumor-infiltrating myeloid cells.
[Hematopoietic growth factors induce proliferation of non-hematopoietic tumor cells].
The immune microenvironment of myeloma.
iNKT cell-neutrophil crosstalk promotes colorectal cancer pathogenesis.
Role of Germline Predisposition to Therapy-Related Myeloid Neoplasms.
Macrophage migration inhibitory factor (MIF) is induced by cytotoxic drugs and is involved in immune escape and migration in childhood rhabdomyosarcoma.
Reduced frequencies and functional impairment of dendritic cell subsets and non-classical monocytes in myelodysplastic syndromes.
PRMT1 reverts the immune escape of necroptotic colon cancer through RIP3 methylation.
Epstein-Barr virus Zta-induced immunomodulators from nasopharyngeal carcinoma cells upregulate interleukin-10 production from monocytes.

Human myeloid dendritic cells treated with supernatants of rotavirus infected Caco-2 cells induce a poor Th1 response.
CD71(+) Erythroid Suppressor Cells Promote Fetomaternal Tolerance through Arginase-2 and PDL-1.

Table A2-30, Cluster 29

Cluster 29 focuses on 1) tumor MHC Class I downregulation, 2) the consequent reduction in tumor cell immunosurveillance, and 3) immunotherapies that can reverse downregulation of MHC-I to aid in anti-tumor immunity (360)
Tumour MHC class I downregulation and immunotherapy (Review).
MHC heterogeneity and response of metastases to immunotherapy.
MHC class I antigens and immune surveillance in transformed cells.
In vivo priming of two distinct antitumor effector populations: the role of MHC class I expression.
Down-regulation of the MHC class I antigen-processing machinery after oncogenic transformation of murine fibroblasts.
MHC class I down-regulation: tumour escape from immune surveillance? (review).
Molecular mechanisms of MHC class I abnormalities and APM components in human tumors.
"Hard" and "soft" lesions underlying the HLA class I alterations in cancer cells: implications for immunotherapy.
MHC presentation via autophagy and how viruses escape from it.
The regulatory network behind MHC class I expression.
The urgent need to recover MHC class I in cancers for effective immunotherapy.
MHC class I transactivator NLRC5 in host immunity, cancer and beyond.
Immune escape phenotype of HPV16-associated tumours: MHC class I expression changes during progression and therapy.
The link between MHC class I abnormalities of tumors, oncogenes, tumor suppressor genes, and transcription factors.
Suppression of MHC class I antigens in oncogenic transformants: association with decreased recognition by cytotoxic T lymphocytes.
NLRC5/MHC class I transactivator is a target for immune evasion in cancer.
Influence of nNav1.5 on MHC class I expression in breast cancer.
CD99 regulates the transport of MHC class I molecules from the Golgi complex to the cell surface.
Tumor escape from immune recognition: loss of HLA-A2 melanoma cell surface expression is associated with a complex rearrangement of the short arm of chromosome 6.
Major histocompatibility complex antigens in normal, acanthotic and neoplastic ovine skin: an association between tumor invasiveness and low level MHC class I expression.
MHC class I-deficient metastatic tumor variants immunoselected by T lymphocytes originate from the coordinated downregulation of APM components.
Novel insights into the molecular mechanisms of HLA class I abnormalities.
Analysis of the major histocompatibility complex class I antigen presentation machinery in normal and malignant renal cells: evidence for deficiencies associated with transformation and progression.
MHC class I antigens, immune surveillance, and tumor immune escape.
Antigenic Hsp70-peptide upregulate altered cell surface MHC class I expression in TAMs and increases anti-tumor function in Dalton's lymphoma bearing mice.
NLRC5 expression in tumors and its role as a negative prognostic indicator in stage III non-small-cell lung cancer patients.



MHC class I-independent activation of virtual memory CD8 T cells induced by chemotherapeutic agent-treated cancer cells.
Major histocompatibility complex class I molecule expression by pancreatic cancer cells is regulated by activation and inhibition of the epidermal growth factor receptor.
Recipients with in utero induction of tolerance upregulated MHC class I in the engrafted donor skin.
Playing hide and seek: Tumor cells in control of MHC class I antigen presentation.
Proto-oncogene PML controls genes devoted to MHC class I antigen presentation.
The selection of tumor variants with altered expression of classical and nonclassical MHC class I molecules: implications for tumor immune escape.
MHC class I antigen processing and presenting machinery: organization, function, and defects in tumor cells.
The MHC Class-I Transactivator NLRC5: Implications to Cancer Immunology and Potential Applications to Cancer Immunotherapy.
Non-immune functions of MHC class I glycoproteins in normal and malignant cells.
MHC class II genotype- and MHC class I and II phenotype-related parameters in sporadic colorectal cancer.
Loss of antigen-presenting molecules (MHC class I and TAP-1) in lung cancer.
Altered MHC class I antigens in tumors.
The tumour suppressor Fhit positively regulates MHC class I expression on cancer cells.
Generation of MHC class I diversity in primary tumors and selection of the malignant phenotype.
MHC Class I Deficiency in Solid Tumors and Therapeutic Strategies to Overcome It.
Clustering of Major Histocompatibility Complex-Class I Molecules in Healthy and Cancer Colon Cells Revealed from Their Nanomechanical Properties.
The escape of cancer from T lymphocytes: immunoselection of MHC class I loss variants harboring structural-irreversible "hard" lesions.
Autophagy-dependent regulation of MHC-I molecule presentation.
Characterization of MHC Class I and $\beta$ -2-Microglobulin Expression in Pediatric Solid Malignancies to Guide Selection of Immune-Based Therapeutic Trials.
Forcing tumor cells to present their own tumor antigens to the immune system: a necessary design for an efficient tumor immunotherapy.
IFN $\gamma$ producing CD8(+) T cells modified to resist major immune checkpoints induce regression of MHC class I-deficient melanomas.
Loss of functional beta 2-microglobulin in metastatic melanomas from five patients receiving immunotherapy.
Cancer Immune Evasion Through Loss of MHC Class I Antigen Presentation.
Immune selective pressure and HLA class I antigen defects in malignant lesions.
Rejection versus escape: the tumor MHC dilemma.
Restoration of MHC-I on Tumor Cells by Fhit Transfection Promotes Immune Rejection and Acts as an Individualized Immunotherapeutic Vaccine.
Induction of immunogenicity of a human renal-cell carcinoma cell line by TAP1-gene transfer.
Immune escape of melanoma: first evidence of structural alterations in two distinct components of the MHC class I antigen processing pathway.
An Evolutionarily Conserved Function of Polycomb Silences the MHC Class I Antigen Presentation Pathway and Enables Immune Evasion in Cancer.
MHC class I molecules act as tumor suppressor genes regulating the cell cycle gene expression, invasion and intrinsic tumorigenicity of melanoma cells.

Enhanced expression of interferon-gamma-induced antigen-processing machinery components in a spontaneously occurring cancer.
Metabolic stress in cancer cells induces immune escape through a PI3K-dependent blockade of IFN $\gamma$ receptor signaling.
APLP2 regulates the expression of MHC class I molecules on irradiated Ewing's sarcoma cells.
Antigen-specific tumor vaccine efficacy in vivo against prostate cancer with low class I MHC requires competent class II MHC.
Trogocytosis of MHC-I/peptide complexes derived from tumors and infected cells enhances dendritic cell cross-priming and promotes adaptive T cell responses.
Tumor immune evasion through loss of MHC class-I antigen presentation.
Coordinate downregulation of multiple MHC class I antigen processing genes in chemical-induced murine tumor cell lines of distinct origin.
Functional deficiencies of components of the MHC class I antigen pathway in human tumors of epithelial origin.
PD-L1 and MHC Class I Expression in High-grade Ovarian Cancers, Including Platinum-resistant Recurrences Treated With Checkpoint Inhibitor Therapy.
[Role of MHC class I molecules in anti-tumoral mechanisms in human malignant melanoma].
Viral MHC class I inhibition evades CD8+ T-cell effector responses in vivo but not CD8+ T-cell priming.
Modulation of MHC class I surface expression in B16F10 melanoma cells by methylseleninic acid.
CD8(+) T cells maintain killing of MHC-I-negative tumor cells through the NKG2D-NKG2DL axis.
Human cytomegalovirus UL18 utilizes US6 for evading the NK and T-cell responses.
The MHC class I-LILRB1 signalling axis as a promising target in cancer therapy.
Tumor CEMIP drives immune evasion of colorectal cancer via MHC-I internalization and degradation.
NLRCS regulates expression of MHC-I and provides a target for anti-tumor immunity in transmissible cancers.
MHC-class I antigen expression on micrometastases in bone marrow of patients with head and neck squamous cell cancer.
Downregulation of tapasin expression in progressive human malignant melanoma.
Selective cytotoxic T-lymphocyte targeting of tumor immune escape variants.
Human herpesvirus 7 u21 downregulates classical and nonclassical class I major histocompatibility complex molecules from the cell surface.
NLRCS, a promising new entry in tumor immunology.
MHC class I TCR engineered anti-tumor CD4 T cells: implications for cancer immunotherapy.
Cytokine therapy reverses NK cell anergy in MHC-deficient tumors.
PD-1/PD-L1 Blockade Therapy for Tumors with Downregulated MHC Class I Expression.
Mechanisms of antigen presentation.
Deficient expression of components of the MHC class I antigen processing machinery in human cervical carcinoma.
Hypoxia and hypoxia-inducible factor (HIF) downregulate antigen-presenting MHC class I molecules limiting tumor cell recognition by T cells.
Antigen processing and immune regulation in the response to tumours.
Cytomegalovirus MHC class I homologues and natural killer cells: an overview.
MHC class I antigen processing regulated by cytosolic proteolysis-short cuts that alter peptide generation.
Endosomal processing for antigen presentation mediated by CD1 and Class I major histocompatibility complex: roads to display or destruction.

ER quality control in the biogenesis of MHC class I molecules.
Correction of defects responsible for impaired Qa-2 class Ib MHC expression on melanoma cells protects mice from tumor growth.
Targeting tumor-associated antigens to the MHC class I presentation pathway.
NLRCS/CITA: A Key Player in Cancer Immune Surveillance.
A membrane-associated MHC-I inhibitory axis for cancer immune evasion.
Blocking oncogenic RAS enhances tumour cell surface MHC class I expression but does not alter susceptibility to cytotoxic lymphocytes.
Genome-wide Screens Identify Lineage- and Tumor-Specific Genes Modulating MHC-I- and MHC-II-Restricted Immunosurveillance of Human Lymphomas.
HLA-dependent tumour development: a role for tumour associated macrophages?
Secretogranin II influences the assembly and function of MHC class I in melanoma.
Activated SUMOylation restricts MHC class I antigen presentation to confer immune evasion in cancer.
Loss of interferon-gamma inducibility of the MHC class II antigen processing pathway in head and neck cancer: evidence for post-transcriptional as well as epigenetic regulation.
Autophagy promotes immune evasion of pancreatic cancer by degrading MHC-I.
Comparison of Automated and Conventional IHC Visual Scoring Analysis for MHC Class I and Tapasin Expression in Cervical Carcinoma.
High-risk human papillomavirus E7 expression reduces cell-surface MHC class I molecules and increases susceptibility to natural killer cells.
BRAFV600E Co-opts a Conserved MHC Class I Internalization Pathway to Diminish Antigen Presentation and CD8+ T-cell Recognition of Melanoma.
A few good peptides: MHC class I-based cancer immunosurveillance and immunoevasion.
Monitoring peptide processing for MHC class I molecules in the endoplasmic reticulum.
Complex interplay of activating and inhibitory signals received by Vgamma9Vdelta2 T cells revealed by target cell beta2-microglobulin knockdown.
Biglycan-mediated upregulation of MHC class I expression in HER-2/neu-transformed cells.
LC3 and NLRCS interaction inhibits NLRCS-mediated MHC class I antigen presentation pathway in endometrial cancer.
Selective autophagy of MHC-I promotes immune evasion of pancreatic cancer.
Positive Role of the MHC Class-I Antigen Presentation Regulator m04/gp34 of Murine Cytomegalovirus in Antiviral Protection by CD8 T Cells.
Cathepsin G-mediated proteolytic degradation of MHC class I molecules to facilitate immune detection of human glioblastoma cells.
Activation of NF- $\kappa$ B and p300/CBP potentiates cancer chemoimmunotherapy through induction of MHC-I antigen presentation.
Differential MHC class II component expression in HPV-positive cervical cancer cells: implication for immune surveillance.
Downregulation of the constitutive tapasin expression in human tumor cells of distinct origin and its transcriptional upregulation by cytokines.
Aberrant Expression of MHC Class II in Melanoma Attracts Inflammatory Tumor-Specific CD4+ T- Cells, Which Dampen CD8+ T-cell Antitumor Reactivity.
Regulation of the antigen presentation machinery in cancer and its implication for immune surveillance.
Can viruses help us to understand and classify the MHC class I molecules at the maternal-fetal interface?

Downregulation of MHC-I expression is prevalent but reversible in Merkel cell carcinoma.
Activation of CXCR4 triggers ubiquitination and down-regulation of major histocompatibility complex class I (MHC-I) on epithelioid carcinoma HeLa cells.
Characterizing MHC-I Genotype Predictive Power for Oncogenic Mutation Probability in Cancer Patients.
Targeting the MHC Class II antigen presentation pathway in cancer immunotherapy.
Enriched HLA-DQ3 phenotype and decreased class I major histocompatibility complex antigen expression in recurrent respiratory papillomatosis.
Open MHC Class I Conformers: A Look through the Looking Glass.
Mechanical stress downregulates MHC class I expression on human cancer cell membrane.
Engineering Tolerance toward Allogeneic CAR-T Cells by Regulation of MHC Surface Expression with Human Herpes Virus-8 Proteins.
miR-148a-3p silences the CANX/MHC-I pathway and impairs CD8(+) T cell-mediated immune attack in colorectal cancer.
Promotion on NLRC5 upregulating MHC-I expression by IFN- $\gamma$ in MHC-I-deficient breast cancer cells.
Metformin rescues cell surface major histocompatibility complex class I (MHC-I) deficiency caused by oncogenic transformation.
Generation of cancer vaccine immunogens derived from major histocompatibility complex (MHC) class I molecules using variable epitope libraries.
Differential downregulation of endoplasmic reticulum-residing chaperones calnexin and calreticulin in human metastatic melanoma.
Adenoviral-mediated gene transfer of ICP47 inhibits major histocompatibility complex class I expression on vascular cells in vitro.
T cell- and perforin-dependent depletion of B cells in vivo by staphylococcal enterotoxin A.
Genetic mechanisms of HLA-I loss and immune escape in diffuse large B cell lymphoma.
Microenvironmental stresses induce HLA-E/Qa-1 surface expression and thereby reduce CD8(+) T-cell recognition of stressed cells.
Local expression of TAP-1 and MHC-I molecules and their relationship in condyloma acuminatum.
MHC Class II Antigen Presentation by Lymphatic Endothelial Cells in Tumors Promotes Intratumoral Regulatory T cell-Suppressive Functions.
Transcriptional regulation of the major histocompatibility complex (MHC) class I heavy chain, TAP1 and LMP2 genes by the human papillomavirus (HPV) type 6b, 16 and 18 E7 oncoproteins.
The varicellovirus-encoded TAP inhibitor UL49.5 regulates the presentation of CTL epitopes by Qa-1b1.
Mass Spectrometric Identification and Molecular Modeling of Glycopeptides Presented by MHC Class I and II Processing Pathways.
Loss of Major Histocompatibility Complex Class I, CD8 + Tumor-infiltrating Lymphocytes, and PD-L1 Expression in Ovarian Clear Cell Carcinoma.
T cells specific for a TAP-independent self-peptide remain naïve in tumor-bearing mice and are fully exploitable for therapy.
Enhanced MHC class I and costimulatory molecules on B16F10 cells by Ganoderma lucidum polysaccharides.
Analysis of Class I Major Histocompatibility Complex Gene Transcription in Human Tumors Caused by Human Papillomavirus Infection.
Transcriptional control of MHC genes in fetal trophoblast cells.
Seeking Cellular Fitness and Immune Evasion: Autophagy in Pancreatic Carcinoma.

Active immunosurveillance in the tumor microenvironment of colorectal cancer is associated with low frequency tumor budding and improved outcome.
Tumor necrosis factor overcomes immune evasion in p53-mutant medulloblastoma.
Major histocompatibility complex class I expression impacts on patient survival and type and density of immune cells in biliary tract cancer.
The transfer of host MHC class I protein protects donor cells from NK cell and macrophage-mediated rejection during hematopoietic stem cell transplantation and engraftment in mice.
Selected HLA-B allotypes are resistant to inhibition or deficiency of the transporter associated with antigen processing (TAP).
T cell infiltration and MHC I and II expression in the presence of tumor antigens: An immunohistochemical study in patients with serous epithelial ovarian cancer.
MHC class II restricted neoantigen: A promising target in tumor immunotherapy.
Restoration of the expression of transporters associated with antigen processing in lung carcinoma increases tumor-specific immune responses and survival.
A tumor-associated glycoprotein that blocks MHC class II-dependent antigen presentation by dendritic cells.
In vivo CRISPR screens reveal the landscape of immune evasion pathways across cancer.
HER-2/neu-mediated regulation of components of the MHC class I antigen-processing pathway.
Down-regulation of the transporter for antigen presentation, proteasome subunits, and class I major histocompatibility complex in tumor cell lines.
Altered expression of TAP-1 and major histocompatibility complex class I in laryngeal papillomatosis: correlation of TAP-1 with disease.
A novel category of antigens enabling CTL immunity to tumor escape variants: Cinderella antigens.
MHC class I loss is a frequent mechanism of immune escape in papillary thyroid cancer that is reversed by interferon and selumetinib treatment in vitro.
CD8 T cell-evasive functions of human cytomegalovirus display pervasive MHC allele specificity, complementarity, and cooperativity.
Loss of transporter protein, encoded by the TAP-1 gene, is highly correlated with loss of HLA expression in cervical carcinomas.
Alterations in the expression of MHC class I glycoproteins by B16BL6 melanoma cells modulate insulin receptor-regulated signal transduction and augment [correction of augments] resistance to apoptosis.
Frequent Loss of IRF2 in Cancers Leads to Immune Evasion through Decreased MHC Class I Antigen Presentation and Increased PD-L1 Expression.
Characterisation of Viscum album L. effect on immune escape proteins PD-L1, PD-L2 and MHC-I in prostate, colon, lung and breast cancer cells.
Immunologic targeting of FOXP3 in inflammatory breast cancer cells.
Suppression of inducible CD4 regulatory cells by MHC class I-restricted human tumor epitope specific TCR engineered multifunctional CD4 T cells.
HIV-1 infection leads to increased HLA-E expression resulting in impaired function of natural killer cells.
TAP-inhibitors from old world primate 1-herpesviruses and their use: WO2009008713.
DUX4 Suppresses MHC Class I to Promote Cancer Immune Evasion and Resistance to Checkpoint Blockade.
Induction of protective CTL immunity against peptide transporter TAP-deficient tumors through dendritic cell vaccination.
Immunologic mechanisms of antitumor activity.

Multifront assault on antigen presentation by Japanese encephalitis virus subverts CD8+ T cell responses.
Interferons and Resistance Mechanisms in Tumors and Pathogen-Driven Diseases-Focus on the Major Histocompatibility Complex (MHC) Antigen Processing Pathway.
Compartmentalization of class II antigen presentation: contribution of cytoplasmic and endosomal processing.
IL-17A increases MHC class I expression and promotes T cell activation in papillary thyroid cancer patients with coexistent Hashimoto's thyroiditis.
Gamma Irradiation Triggers Immune Escape in Glioma-Propagating Cells.
MHC class II engagement by its ligand LAG-3 (CD223) contributes to melanoma resistance to apoptosis.
Metastatic melanoma cells evade immune detection by silencing STAT1.
High Levels of Class I Major Histocompatibility Complex mRNA Are Present in Epstein-Barr Virus-Associated Gastric Adenocarcinomas.
Cancer immunotherapy in clinical oncology.
T cell recognition of HLA-A2 restricted tumor antigens is impaired by the oncogene HER2.
TLR2 Promotes Glioma Immune Evasion by Downregulating MHC Class II Molecules in Microglia.
The immune-related role of beta-2-microglobulin in melanoma.
Involvement of STAT3 in immune evasion during lung tumorigenesis.
A novel approach to rescue immune escape in oral squamous cell carcinoma: Combined use of interferon- $\gamma$ and LY294002.
Proto-oncogene PML enhances antigen presentation by MHC class I molecules in human lung cancer cells.
TAP-independent self-peptides enhance T cell recognition of immune-escaped tumors.
TAP-ing into TIEPPs for cancer immunotherapy.
Suppression of tumor antigen presentation during aneuploid tumor evolution contributes to immune evasion.
Tumor escape from immune surveillance.
IRF1 and NF- $\kappa$ B restore MHC class I-restricted tumor antigen processing and presentation to cytotoxic T cells in aggressive neuroblastoma.
CD8+ T cell responses against TAP-inhibited cells are readily detected in the human population.
Principles of tumor immunosurveillance and implications for immunotherapy.
Brucella abortus induces intracellular retention of MHC-I molecules in human macrophages down-modulating cytotoxic CD8(+) T cell responses.
High frequency of functionally active Melan-a-specific T cells in a patient with progressive immunoproteasome-deficient melanoma.
Exploitation of herpesvirus immune evasion strategies to modify the immunogenicity of human mesenchymal stem cell transplants.
Quantitative correlation between HLA class I allele expression and recognition of melanoma cells by antigen-specific cytotoxic T lymphocytes.
Strategies for the development of vaccines to treat breast cancer.
Tumor necrosis factor $\alpha$ -induced hypoxia-inducible factor 1 $\alpha$ - $\beta$ -catenin axis regulates major histocompatibility complex class I gene activation through chromatin remodeling.
CTL-defined cancer vaccines: perspectives for active immunotherapeutic interventions in minimal residual disease.
Peptide Vaccination in Clinical Oncology.
Lymphocyte-melanoma interaction: role of surface molecules.

Lymphocyte activation gene 3: a novel therapeutic target in chronic lymphocytic leukemia.
Tapasin-mediated editing of the MHC I immunopeptidome is epitope specific and dependent on peptide off-rate, abundance, and level of tapasin expression.
Spotlight on TAP and its vital role in antigen presentation and cross-presentation.
Tumor-associated antigen profiling in breast and ovarian cancer: mRNA, protein or T cell recognition?
NLR5 deficiency has a moderate impact on immunodominant CD8(+) T-cell responses during rotavirus infection of adult mice.
Tumour immunology, vaccination and escape strategies.
The MHC Class I Cancer-Associated Neoepitope Trh4 Linked with Impaired Peptide Processing Induces a Unique Noncanonical TCR Conformer.
Reversible epigenetic down-regulation of MHC molecules by devil facial tumour disease illustrates immune escape by a contagious cancer.
Immune responses to tumour antigens: implications for antigen specific immunotherapy of cancer.
gp100/pmel 17 is a murine tumor rejection antigen: induction of "self"-reactive, tumoricidal T cells using high-affinity, altered peptide ligand.
Expression of class I MHC molecule, HSP70 and TAP in human hepatocellular carcinoma.
Inactivation of tumor-specific CD8 <sup>+</sup> CTLs by tumor-infiltrating tolerogenic dendritic cells.
Molecular and Genetic Characterization of MHC Deficiency Identifies EZH2 as Therapeutic Target for Enhancing Immune Recognition.
A Cell-Surface MHC-I Inhibitory Axis Promotes Immune Evasion in Cancer.
Consequences of antigen self-presentation by tumor-specific cytotoxic T cells.
Evaluation of cancer testis antigen (CT10, PRAME) and MHC I expression in high-grade urothelial carcinoma of the bladder.
Genome-wide expression profiling reveals EBV-associated inhibition of MHC class I expression in nasopharyngeal carcinoma.
RARbeta involvement in enhancement of lung tumor cell immunogenicity revealed by array analysis.
[Genetic techniques to overcome antitumor anergy].
How tumors escape immune destruction and what we can do about it.
Epigenetic modification and preliminary investigation of the mechanism of the immune evasion of HL-60 cells.
Mouse models of efficient and inefficient anti-tumor immunity, with emphasis on minimal residual disease and tumor escape.
The newly-arisen Devil facial tumour disease 2 (DFT2) reveals a mechanism for the emergence of a contagious cancer.
The role of the Major Histocompatibility Complex in the spread of contagious cancers.
Nonclassical MHC class Ib-restricted cytotoxic T cells monitor antigen processing in the endoplasmic reticulum.
A novel immune evasion mechanism of LMP-1, an EBV-primary oncogene, in nasopharyngeal carcinoma.
Reduced recognition of metastatic melanoma cells by autologous MART-1 specific CTL: relationship to TAP expression.
Genomic Alterations in CIITA Are Frequent in Primary Mediastinal Large B Cell Lymphoma and Are Associated with Diminished MHC Class II Expression.
Immune system evasion by peripheral nerve sheath tumor.
Inhibition of serine-peptidase activity enhances the generation of a survivin-derived HLA-A2-presented CTL epitope in colon-carcinoma cells.
Selective autophagy of NLR5 promotes immune evasion of endometrial cancer.

A tumor escape variant that has lost one major histocompatibility complex class I restriction element induces specific CD8+ T cells to an antigen that no longer serves as a target.
Crosstalk between tumor T lymphocytes and reactive T lymphocytes in cutaneous T cell lymphomas.
Implication of the $\beta 2$ -microglobulin gene in the generation of tumor escape phenotypes.
Metabolic Stress Triggers Immune Escape by Tumors.
CXCL14 suppresses human papillomavirus-associated head and neck cancer through antigen-specific CD8(+) T-cell responses by upregulating MHC-I expression.
Protein Kinase D3 Promotes the Reconstruction of OSCC Immune Escape Niche Via Regulating MHC-I and Immune Inhibit Molecules Expression.
Expression of the Nonclassical MHC Class I, Saha-UD in the Transmissible Cancer Devil Facial Tumour Disease (DFTD).
Tumor escape mechanisms from immunosurveillance: induction of unresponsiveness in a specific MHC-restricted CD4+ human T cell clone by the autologous MHC class II+ melanoma.
MHC class I antigen processing pathway defects, ras mutations and disease stage in colorectal carcinoma.
TNF $\alpha$ driven HIF-1 $\alpha$ -hexokinase II axis regulates MHC-I cluster stability through actin cytoskeleton.
Induction of regular cytolytic T cell synapses by bispecific single-chain antibody constructs on MHC class I-negative tumor cells.
Loss of tapasin in human lung and colon cancer cells and escape from tumor-associated antigen-specific CTL recognition.
Aspects of cancer immunotherapy.
Cell-mediated immunity and expression of MHC class I and class II molecules in dogs naturally infected by canine transmissible venereal tumor: Is there complete spontaneous regression outside the experimental CTVT?
Temozolomide increases MHC-I expression via NF- $\kappa$ B signaling in glioma stem cells.
The effects of IFITM1 and IFITM3 gene deletion on IFN $\gamma$ stimulated protein synthesis.
Inhibition of antigen presentation by Brucella: many more than many ways.
Deficiency of transporter for antigen presentation (TAP) in tumor cells allows evasion of immune surveillance and increases tumorigenesis.
Introduction.
Reduced MHC class II expression in medullary thyroid cancer identifies patients with poor prognosis.
Impacts of the MHC class I-like XNC10 and innate-like T cells on tumor tolerance and rejection in the amphibian Xenopus.
The E6E7 oncoproteins of cutaneous human papillomavirus type 38 interfere with the interferon pathway.
Coexpression of CD40L and CD70 by semiallogenic tumor cells induces anti-tumor immunity.
Rationally designed inhibitor targeting antigen-trimming aminopeptidases enhances antigen presentation and cytotoxic T-cell responses.
Roles of tumour localization, second signals and cross priming in cytotoxic T-cell induction.
Flt3-L gene therapy enhances immunocytokine-mediated antitumor effects and induces long-term memory.
Immune Escape Mechanisms in Intravascular Large B-Cell Lymphoma: A Molecular Cytogenetic and Immunohistochemical Study.
Peroxyntirite promotes immune evasion by reducing tumor antigenicity.
Mechanisms of tumor escape: role of tumor microenvironment in inducing apoptosis of cytolytic effector cells.



A double recombinant adenovirus expressing the costimulatory molecule B7-1 (murine) and human IL-2 induces complete tumor regression in a murine breast adenocarcinoma model.
IL-10 Restores MHC Class I Expression and Interferes With Immunity in Papillary Thyroid Cancer With Hashimoto Thyroiditis.
Loss of tapasin correlates with diminished CD8(+) T-cell immunity and prognosis in colorectal cancer.
Cytosolic Processing Governs TAP-Independent Presentation of a Critical Melanoma Antigen.
The HDAC Inhibitor Domatinostat Promotes Cell-Cycle Arrest, Induces Apoptosis, and Increases Immunogenicity of Merkel Cell Carcinoma Cells.
The mercy of adrenocortical tumor cells on lymphocytes.
TAP expression level in tumor cells defines the nature and processing of MHC class I peptides for recognition by tumor-specific cytotoxic T lymphocytes.
Small-molecule MHC-II inducers promote immune detection and anti-cancer immunity via editing cancer metabolism.
NLRCS: new cancer buster?
Immunologic and clinical outcomes of a randomized phase II trial of two multi-peptide vaccines for melanoma in the adjuvant setting.
Combination of intratumoral injections of vaccinia virus MVA expressing GM-CSF and immunization with DNA vaccine prolongs the survival of mice bearing HPV16 induced tumors with downregulated expression of MHC class I molecules.
Melanoma-Derived Extracellular Vesicles Bear the Potential for the Induction of Antigen-Specific Tolerance.
Tim-3 Promotes <i>Listeria monocytogenes</i> Immune Evasion by Suppressing Major Histocompatibility Complex Class I.
The mechanism of de novo expression of programmed cell death-ligand 1 in squamous cell carcinoma of the lung.
Restoration of immune response gene induction in trophoblast tumor cells associated with cellular senescence.
Epigenetic enhancement of antigen processing and presentation promotes immune recognition of tumors.
Immune Escape of Relapsed AML Cells after Allogeneic Transplantation.
Qa-1-Restricted CD8(+) T Cells Can Compensate for the Absence of Conventional T Cells during Viral Infection.
Constitutive expression of TGF- $\beta$ 1, interleukin-6 and interleukin-8 by tumor cells as a major component of immune escape in human ovarian carcinoma.
Prognostic significance of activated cytotoxic T-lymphocytes in primary nodal diffuse large B-cell lymphomas.
Cytomegalovirus inhibition of extrinsic apoptosis determines fitness and resistance to cytotoxic CD8 T cells.
Ectopic transplantation of equine invasive trophoblast.
MC32 tumor cells acquire Ag-specific CTL resistance through the loss of CEA in a colon cancer model.
Ex vivo-activated MHC-unrestricted immune effectors for cancer adoptive immunotherapy.
Human Papillomavirus 16 E6 Suppresses Transporter Associated with Antigen-Processing Complex in Human Tongue Keratinocyte Cells by Activating Lymphotoxin Pathway.
The role of MHC genes in contagious cancer: the story of Tasmanian devils.
Reversal of epigenetic silencing of MHC class I chain-related protein A and B improves immune recognition of Merkel cell carcinoma.

Oncoprotein SND1 hijacks nascent MHC-I heavy chain to ER-associated degradation, leading to impaired CD8(+) T cell response in tumor.
Phosphoinositide 3-Kinase Signaling Can Modulate MHC Class I and II Expression.
MHC Class II is Induced by IFN $\gamma$ and Follows Three Distinct Patterns of Expression in Colorectal Cancer Organoids.
Interactions of host IL-6 and IFN-gamma and cancer-derived TGF-beta1 on MHC molecule expression during tumor spontaneous regression.
Incomplete tolerance to the tumour-associated antigen MDM2.
CDK4/6 inhibitors sensitize gammaherpesvirus-infected tumor cells to T-cell killing by enhancing expression of immune surface molecules.
Insertion of exogenous epitopes in the E3-19K of oncolytic adenoviruses to enhance TAP-independent presentation and immunogenicity.
Targeting MDM2 enhances antileukemia immunity after allogeneic transplantation via MHC-II and TRAIL-R1/2 upregulation.
Effective adoptive therapy of tap-deficient lymphoma using diverse high avidity alloreactive T cells.
The hemochromatosis protein HFE 20 years later: An emerging role in antigen presentation and in the immune system.
Immune escape from a graft-versus-leukemia effect may play a role in the relapse of myeloid leukemias following allogeneic bone marrow transplantation.
B16 melanomas evade antitumor immunity by the loss of epitope presentation and the acquisition of tumor resistance to granzyme B.
Tumour-induced suppression of immune response and its correction.
Morphological evidence of an activated cytotoxic T-cell infiltrate in EBV-positive gastric carcinoma preventing lymph node metastases.
T-cell anergy induced by antigen presenting cells treated with the hemolysin of <i>Listeria monocytogenes</i> .
Immunoproteasome subunits LMP2 and LMP7 downregulation in primary malignant melanoma lesions: association with lack of spontaneous regression.
Impairment of Macrophage Presenting Ability and Viability by <i>Echinococcus granulosus</i> Antigens.
E1A-based determinants of oncogenicity in human adenovirus groups A and C.
Cancer immunoediting from immune surveillance to immune escape.
Vascular endothelial cells have impaired capacity to present immunodominant, antigenic peptides: a mechanism of cell type-specific immune escape.
Cellular senescence and the host immune system in aging and age-related disorders.
Cell surface sphingomyelin: key role in cancer initiation, progression, and immune evasion.
A critical role of non-classical MHC in tumor immune evasion in the amphibian <i>Xenopus</i> model.
Identification of mechanisms underlying transporter associated with antigen processing deficiency in metastatic murine carcinomas.
Evaluation of cellular tumour rejection mechanisms in the peritumoral bladder wall after bacillus Calmette-Guérin treatment.
Biglycan as a potential regulator of tumorigenicity and immunogenicity in K-RAS-transformed cells.
Immunotherapeutic potential of whole tumour cells.
CML cells actively evade host immune surveillance through cytokine-mediated downregulation of MHC-II expression.
Cell-dose-dependent increases in circulating levels of immune effector cells in rhesus macaques following intracranial injection of allogeneic MSCs.
Altered patterns of T cell cytokine production induced by relapsed pre-B ALL cells.

Human embryonic stem cells and their differentiated derivatives are less susceptible to immune rejection than adult cells.
Retroviral expression of MIR2 decreases both surface MHC class I and the alloimmune CTL response.
Enhanced expression of LAG-3 on lymphocyte subpopulations from persistently lymphocytotic cattle infected with bovine leukemia virus.
Indirect CD4+ T-cell-mediated elimination of MHC II(NEG) tumor cells is spatially restricted and fails to prevent escape of antigen-negative cells.
T cell receptor binding kinetics required for T cell activation depend on the density of cognate ligand on the antigen-presenting cell.
Update on Kaposi's sarcoma and other HHV8 associated diseases. Part 2: pathogenesis, Castleman's disease, and pleural effusion lymphoma.
Discovery of a Metastatic Immune Escape Mechanism Initiated by the Loss of Expression of the Tumour Biomarker Interleukin-33.
Hypoimmunogenic derivatives of induced pluripotent stem cells evade immune rejection in fully immunocompetent allogeneic recipients.
Spheres derived from the human SK-RC-42 renal cell carcinoma cell line are enriched in cancer stem cells.
Neoantigen Fitness Model Predicts Lower Immune Recognition of Cutaneous Squamous Cell Carcinomas Than Actinic Keratoses.
Human preprocalcitonin self-antigen generates TAP-dependent and -independent epitopes triggering optimised T-cell responses toward immune-escaped tumours.
Immunization Strategies Producing a Humoral IgG Immune Response against Devil Facial Tumor Disease in the Majority of Tasmanian Devils Destined for Wild Release.
Brain activation of monocyte-lineage cells: involvement of interleukin-6.
Lack of CIITA expression is central to the absence of antigen presentation functions of trophoblast cells and is caused by methylation of the IFN-gamma inducible promoter (PIV) of CIITA.
Role and regulation of CD1d in normal and pathological B cells.
Antigen processing and remodeling of the endosomal pathway: requirements for antigen cross-presentation.
The impact of interferon gamma receptor expression on the mechanism of escape from host immune surveillance in hepatocellular carcinoma.
Immune evasion by neocartilage-derived chondrocytes: Implications for biologic repair of joint articular cartilage.
Comparative transcriptomic profiling in HPV-associated cervical carcinogenesis: Implication of MHC class II and immunoglobulin heavy chain genes.
Identification of the neoplastically transformed cells in Marek's disease herpesvirus-induced lymphomas: recognition by the monoclonal antibody AV37.
[Construction of $\beta 2m$ knockout mice].
Expression and clinical value of the soluble major histocompatibility complex class I-related chain A molecule in the serum of patients with renal tumors.
Oncogenic drivers dictate immune control of acute myeloid leukemia.
A 'hairy' privilege.
Interferon- $\beta$ Modulates the Innate Immune Response against Glioblastoma Initiating Cells.
Expression of thimet oligopeptidase (THOP) modulated by oxidative stress in human multidrug resistant (MDR) leukemia cells.
Loss of antigen-processing molecules in primary orbital melanoma.

Signaling and transcriptional changes critical for transformation of human cells by simian virus 40 small tumor antigen or protein phosphatase 2A B56gamma knockdown.
Immunology of naturally transmissible tumours.
Optineurin Guards IFN $\gamma$ Signaling in Cancer Cells.
Nitric oxide signaling pathway activation inhibits the immune escape of pancreatic carcinoma cells.
Human cytomegalovirus-derived protein UL18 alters the phenotype and function of monocyte-derived dendritic cells.
In vitro atrazine exposure affects the phenotypic and functional maturation of dendritic cells.
Placental trophoblast and endothelial cells as target of maternal immune response.
Expression of heat shock protein 70 and c-myc in cervical carcinoma.
Gene therapy of murine teratocarcinoma: separate functions for insulin-like growth factors I and II in immunogenicity and differentiation.
CD40 expression in Wehi-164 cell line.
Immune evasion strategies of pediatric precursor-B acute lymphoblastic leukemia after allogeneic bone marrow transplantation-a case study.
Corneal transplantation and immune privilege.
Type 2 human papillomavirus E7 attenuates E-cadherin expression in human keratinocytes.

Table A2-31, Cluster 30

Cluster 30 focuses on variant surface glycoprotein coat density in African trypanosomes, emphasizing the mechanisms responsible for this parasite's evasion of its host's immune response (305)
DNA metabolism and genetic diversity in Trypanosomes.
DNA Recombination Strategies During Antigenic Variation in the African Trypanosome.
Antigenic diversity is generated by distinct evolutionary mechanisms in African trypanosome species.
African trypanosomes: the genome and adaptations for immune evasion.
The genome of the African trypanosome <i>Trypanosoma brucei</i> .
Quantitative sequencing confirms VSG diversity as central to immune evasion by <i>Trypanosoma brucei</i> .
Antigenic variation in African trypanosomes.
Variant surface glycoprotein density defines an immune evasion threshold for African trypanosomes undergoing antigenic variation.
VSG 117 gene is conservatively present and early expressed in <i>Trypanosma evansi</i> YNB stock.
Evolution of the variant surface glycoprotein family in African trypanosomes.
Genome research and evolution in trypanosomes.
Trypanosomes expressing a mosaic variant surface glycoprotein coat escape early detection by the immune system.
Regulation of Antigenic Variation by <i>Trypanosoma brucei</i> Telomere Proteins Depends on Their Unique DNA Binding Activities.
A histone methyltransferase modulates antigenic variation in African trypanosomes.
Variant antigen diversity in <i>Trypanosoma vivax</i> is not driven by recombination.
To the Surface and Back: Exo- and Endocytic Pathways in <i>Trypanosoma brucei</i> .
Phylogenetic and syntenic data support a single horizontal transference to a <i>Trypanosoma</i> ancestor of a prokaryotic proline racemase implicated in parasite evasion from host defences.
Competition among variants is predictable and contributes to the antigenic variation dynamics of African trypanosomes.

Beyond immune escape: a variant surface glycoprotein causes suramin resistance in <i>Trypanosoma brucei</i> .
Histone deacetylases play distinct roles in telomeric VSG expression site silencing in African trypanosomes.
A role for RAD51 and homologous recombination in <i>Trypanosoma brucei</i> antigenic variation.
Blocking Synthesis of the Variant Surface Glycoprotein Coat in <i>Trypanosoma brucei</i> Leads to an Increase in Macrophage Phagocytosis Due to Reduced Clearance of Surface Coat Antibodies.
The anatomy and transcription of a monocistronic expression site for a metacyclic variant surface glycoprotein gene in <i>Trypanosoma brucei</i> .
TelAP1 links telomere complexes with developmental expression site silencing in African trypanosomes.
<i>Trypanosoma brucei</i> Orc1 is essential for nuclear DNA replication and affects both VSG silencing and VSG switching.
Turnover of Variant Surface Glycoprotein in <i>Trypanosoma brucei</i> Is Not Altered in Response to Specific Silencing.
African trypanosomes evade immune clearance by O-glycosylation of the VSG surface coat.
Combinatorial selection of high affinity RNA ligands to live African trypanosomes.
Evasion of immune responses by <i>Trypanosoma cruzi</i> , the etiological agent of Chagas disease.
Mechanisms of <i>Trypanosoma cruzi</i> persistence in Chagas disease.
Telomere length affects the frequency and mechanism of antigenic variation in <i>Trypanosoma brucei</i> .
Molecular mechanisms underlying the control of antigenic variation in African trypanosomes.
Escape mechanisms of African trypanosomes: why trypanosomiasis is keeping us awake.
African trypanosomes expressing multiple VSGs are rapidly eliminated by the host immune system.
Developmental changes and metabolic reprogramming during establishment of infection and progression of <i>Trypanosoma brucei</i> through its insect host.
Escaping Deleterious Immune Response in Their Hosts: Lessons from Trypanosomatids.
Is It Possible to Intervene in the Capacity of <i>Trypanosoma cruzi</i> to Elicit and Evade the Complement System?
Mono-allelic VSG expression by RNA polymerase I in <i>Trypanosoma brucei</i> : expression site control from both ends?
Antigenic variation in the African trypanosome: molecular mechanisms and phenotypic complexity.
Multifunctional class I transcription in <i>Trypanosoma brucei</i> depends on a novel protein complex.
Point mutations are associated with a gene duplication leading to the bloodstream reexpression of a trypanosome metacyclic VSG.
Visualisation of experimentally determined and predicted protein N-glycosylation and predicted glycosylphosphatidylinositol anchor addition in <i>Trypanosoma brucei</i> .
Intrinsic DNA curvature in trypanosomes.
Transcriptional regulation of metacyclic variant surface glycoprotein gene expression during the life cycle of <i>Trypanosoma brucei</i> .
Life and times: synthesis, trafficking, and evolution of VSG.
Two pathways of homologous recombination in <i>Trypanosoma brucei</i> .
<i>Trypanosoma brucei</i> DMC1 does not act in DNA recombination, repair or antigenic variation in bloodstream stage cells.
DNA double-strand breaks and telomeres play important roles in <i>trypanosoma brucei</i> antigenic variation.
A DOT1B/Ribonuclease H2 Protein Complex Is Involved in R-Loop Processing, Genomic Integrity, and Antigenic Variation in <i>Trypanosoma brucei</i> .

Insufficient TLR activation contributes to the slow development of CD8+ T cell responses in <i>Trypanosoma cruzi</i> infection.
Adaptin evolution in kinetoplastids and emergence of the variant surface glycoprotein coat in African trypanosomatids.
The Dialogue of the Host-Parasite Relationship: <i>Leishmania</i> spp. and <i>Trypanosoma cruzi</i> Infection.
Activity of a trypanosome metacyclic variant surface glycoprotein gene promoter is dependent upon life cycle stage and chromosomal context.
Mechanisms of Infectivity and Evasion Derived from Microvesicles Cargo Produced by <i>Trypanosoma cruzi</i> .
Comprehensive glycoprofiling of the epimastigote and trypomastigote stages of <i>Trypanosoma cruzi</i> .
Evasion of the Immune Response by <i>Trypanosoma cruzi</i> during Acute Infection.
Evaluation of mechanisms that may generate DNA lesions triggering antigenic variation in African trypanosomes.
Does DNA replication direct locus-specific recombination during host immune evasion by antigenic variation in the African trypanosome?
Variable Surface Glycoprotein from <i>Trypanosoma brucei</i> Undergoes Cleavage by Matrix Metalloproteinases: An in silico Approach.
Application of long read sequencing to determine expressed antigen diversity in <i>Trypanosoma brucei</i> infections.
Genome hyperevolution and the success of a parasite.
Multiple mechanisms of immune evasion by African trypanosomes.
[Antigenic variation and the problem of vaccines against African trypanosomes].
Comparative Analysis of Virulence Mechanisms of Trypanosomatids Pathogenic to Humans.
Distinct roles for two RAD51-related genes in <i>Trypanosoma brucei</i> antigenic variation.
<i>Trypanosoma cruzi</i> evades the protective role of interferon-gamma-signaling in parasite-infected cells.
Variant antigen repertoires in <i>Trypanosoma congolense</i> populations and experimental infections can be profiled from deep sequence data using universal protein motifs.
Immune Evasion Strategies of <i>Trypanosoma cruzi</i> .
Analysis of a donor gene region for a variant surface glycoprotein and its expression site in African trypanosomes.
Evasion and Immuno-Endocrine Regulation in Parasite Infection: Two Sides of the Same Coin in Chagas Disease?
Role of expression site switching in the development of resistance to human Trypanosome Lytic Factor-1 in <i>Trypanosoma brucei brucei</i> .
Strain-specific genome evolution in <i>Trypanosoma cruzi</i> , the agent of Chagas disease.
Monoallelic expression and epigenetic inheritance sustained by a <i>Trypanosoma brucei</i> variant surface glycoprotein exclusion complex.
<i>Trypanosoma brucei</i> ATR Links DNA Damage Signaling during Antigenic Variation with Regulation of RNA Polymerase I-Transcribed Surface Antigens.
CD8+ T cells in <i>Trypanosoma cruzi</i> infection.
CRISPR Genome Editing and the Study of Chagas Disease.
Persistent DNA Damage Foci and DNA Replication with a Broken Chromosome in the African Trypanosome.
High affinity nanobodies against the Trypanosome <i>brucei</i> VSG are potent trypanolytic agents that block endocytosis.
<i>Trypanosoma brucei</i> TIF2 suppresses VSG switching by maintaining subtelomere integrity.

A trypanosome metacyclic VSG gene promoter with two functionally distinct, life cycle stage-specific activities.
Trypanosoma cruzi Evades the Complement System as an Efficient Strategy to Survive in the Mammalian Host: The Specific Roles of Host/Parasite Molecules and Trypanosoma cruzi Calreticulin.
Multigene families in Trypanosoma cruzi and their role in infectivity.
Immunity and immune modulation in Trypanosoma cruzi infection.
Control of Variant Surface Glycoprotein Expression by CFB2 in Trypanosoma brucei and Quantitative Proteomic Connections to Translation and Cytokinesis.
Trypanosoma cruzi immune evasion mediated by host cell-derived microvesicles.
The effect of the biflavonoid 2",3"-dihydrochonaflavone on Trypanosoma cruzi Y strain.
Transcriptome analysis of the bloodstream stage from the parasite Trypanosoma vivax.
Bromodomain Proteins Contribute to Maintenance of Bloodstream Form Stage Identity in the African Trypanosome.
Implications of conserved structural motifs in disparate trypanosome surface proteins.
Ku is important for telomere maintenance, but not for differential expression of telomeric VSG genes, in African trypanosomes.
Turnover of Variant Surface Glycoprotein in Trypanosoma brucei Is a Bimodal Process.
Prospects for vaccination against pathogenic African trypanosomes.
Human B cells infected by Trypanosoma cruzi undergo F-actin disruption and cell death via caspase-7 activation and cleavage of phospholipase C $\gamma$ 1.
Bioactive lipids in Trypanosoma cruzi infection.
Role of Trypanosoma cruzi Trans-sialidase on the Escape from Host Immune Surveillance.
Isolation of the repertoire of VSG expression site containing telomeres of Trypanosoma brucei 427 using transformation-associated recombination in yeast.
How the African trypanosomes evade host immune killing.
Molecular diversity of the Trypanosoma cruzi TcSMUG family of mucin genes and proteins.
Parasite-host glycan interactions during Trypanosoma cruzi infection: trans-Sialidase rides the show.
CD8+ T-Cell responses to Trypanosoma cruzi are highly focused on strain-variant trans-sialidase epitopes.
The flagellum of Trypanosoma brucei: new tricks from an old dog.
Life Stage-Specific Cargo Receptors Facilitate Glycosylphosphatidylinositol-Anchored Surface Coat Protein Transport in Trypanosoma brucei.
Immunodominant surface epitopes power immune evasion in the African trypanosome.
Intracellular DNA replication and differentiation of Trypanosoma cruzi is asynchronous within individual host cells in vivo at all stages of infection.
Developmental variation in Rab11-dependent trafficking in Trypanosoma brucei.
Implication of Apoptosis for the Pathogenesis of Trypanosoma cruzi Infection.
Specializations in a successful parasite: what makes the bloodstream-form African trypanosome so deadly?
Subtelomeric I-SceI-Mediated Double-Strand Breaks Are Repaired by Homologous Recombination in Trypanosoma cruzi.
Examination of the telomere G-overhang structure in Trypanosoma brucei.
Modulatory Effect of Trypanosoma cruzi Infective Stages in Different Dendritic Cell Populations in vitro.
Telomere and Subtelomere R-loops and Antigenic Variation in Trypanosomes.
Antigenic variation in trypanosomes: secrets surface slowly.

Immune Evasion Strategies of Trypanosoma brucei within the Mammalian Host: Progression to Pathogenicity.
Transcriptomic analysis reveals metabolic switches and surface remodeling as key processes for stage transition in Trypanosoma cruzi.
The GPI biosynthetic pathway as a therapeutic target for African sleeping sickness.
The within-host dynamics of African trypanosome infections.
Repeat-Driven Generation of Antigenic Diversity in a Major Human Pathogen, Trypanosoma cruzi.
Suppression of subtelomeric VSG switching by Trypanosoma brucei TRF requires its TTAGGG repeat-binding activity.
Identification of clinically approved small molecules that inhibit growth and affect transcript levels of developmentally regulated genes in the African trypanosome.
Antigenic variation in African trypanosomes: monitoring progress.
The role of B-cells and IgM antibodies in parasitemia, anemia, and VSG switching in Trypanosoma brucei-infected mice.
Trypanosoma cruzi Invasion into Host Cells: A Complex Molecular Targets Interplay.
Modulation of Virulence Factors during Trypanosoma cruzi Differentiation.
Trypanosoma cruzi trans-sialidase as a drug target against Chagas disease (American trypanosomiasis).
From silent genes to noisy populations-dialogue between the genotype and phenotypes of antigenic variation.
A structural classification of the variant surface glycoproteins of the African trypanosome.
The Recombinant Form of Trypanosoma cruzi P21 Controls Infection by Modulating Host Immune Response.
The epigenome of Trypanosoma brucei: a regulatory interface to an unconventional transcriptional machine.
TbTRF suppresses the TERRA level and regulates the cell cycle-dependent TERRA foci number with a TERRA binding activity in its C-terminal Myb domain.
Differences in inferred genome-wide signals of positive selection during the evolution of Trypanosoma cruzi and Leishmania spp. lineages: A result of disparities in host and tissue infection ranges?
African trypanosomes.
Identification of potential protein partners that bind to the variant surface glycoprotein in Trypanosoma equiperdum.
N-linked glycans containing linear poly-N-acetylglucosamine as sorting signals in endocytosis in Trypanosoma brucei.
TcTASV: a novel protein family in trypanosoma cruzi identified from a subtractive trypomastigote cDNA library.
Ribonuclease H1-targeted R-loops in surface antigen gene expression sites can direct trypanosome immune evasion.
The Trypanosomal Transferrin Receptor of Trypanosoma Brucei-A Review.
Host Immune Responses and Immune Evasion Strategies in African Trypanosomiasis.
Critical interplay between parasite differentiation, host immunity, and antigenic variation in trypanosome infections.
Extracellular Vesicles from Trypanosoma brucei Mediate Virulence Factor Transfer and Cause Host Anemia.
Efficient targeting of conserved cryptic epitopes of infectious agents by single domain antibodies. African trypanosomes as paradigm.



Differential endocytic functions of <i>Trypanosoma brucei</i> Rab5 isoforms reveal a glycosylphosphatidylinositol-specific endosomal pathway.
Rab5 and Rab11 mediate transferrin and anti-variant surface glycoprotein antibody recycling in <i>Trypanosoma brucei</i> .
Theft and Reception of Host Cell's Sialic Acid: Dynamics of <i>Trypanosoma Cruzi</i> Trans-sialidases and Mucin-Like Molecules on Chagas' Disease Immunomodulation.
Macromolecular trafficking and immune evasion in african trypanosomes.
Chemical Inhibition of Bromodomain Proteins in Insect-Stage African Trypanosomes Perturbs Silencing of the Variant Surface Glycoprotein Repertoire and Results in Widespread Changes in the Transcriptome.
L-arginine metabolism during interaction of <i>Trypanosoma cruzi</i> with host cells.
The transferrin receptor genes of <i>Trypanosoma equiperdum</i> are less diverse in their transferrin binding site than those of the broad-host range <i>Trypanosoma brucei</i> .
<i>Trypanosoma cruzi</i> Calreticulin: Immune Evasion, Infectivity, and Tumorigenesis.
Early Immune Response Elicited by Different <i>Trypanosoma cruzi</i> Infective Stages.
<i>Trypanosoma brucei</i> Secreted Aromatic Ketoacids Activate the Nrf2/HO-1 Pathway and Suppress Pro-inflammatory Responses in Primary Murine Glia and Macrophages.
<i>Trypanosoma cruzi</i> proline racemases are involved in parasite differentiation and infectivity.
Proteomics on the rims: insights into the biology of the nuclear envelope and flagellar pocket of trypanosomes.
Phylogenomics from transcriptomic "bycatch" clarify the origins and diversity of avian trypanosomes in North America.
Extracellular vesicles of <i>Trypanosoma cruzi</i> tissue-culture cell-derived trypomastigotes: Induction of physiological changes in non-parasitized culture cells.
Extracellular Vesicles Shed By <i>Trypanosoma cruzi</i> Potentiate Infection and Elicit Lipid Body Formation and PGE(2) Production in Murine Macrophages.
POLIE suppresses telomerase-mediated telomere G-strand extension and helps ensure proper telomere C-strand synthesis in trypanosomes.
Mucosal Heterologous Prime/Boost Vaccination Induces Polyfunctional Systemic Immunity, Improving Protection Against <i>Trypanosoma cruzi</i> .
Endogenous CD4(+) CD25(+) regulatory T cells have a limited role in the control of <i>Trypanosoma cruzi</i> infection in mice.
<i>Trypanosoma brucei</i> metabolite indolepyruvate decreases HIF-1 $\alpha$ and glycolysis in macrophages as a mechanism of innate immune evasion.
Invariant surface glycoprotein 65 of <i>Trypanosoma brucei</i> is a complement C3 receptor.
Synthesis of Galactosylated Glycosylphosphatidylinositol Derivatives from <i>Trypanosoma brucei</i> .
Phosphoinositides, kinases and adaptors coordinating endocytosis in <i>Trypanosoma brucei</i> .
DNA double strand break position leads to distinct gene expression changes and regulates VSG switching pathway choice.
African Trypanosomes Undermine Humoral Responses and Vaccine Development: Link with Inflammatory Responses?
The inositol-1,2-cyclic phosphate moiety of the cross-reacting determinant, carbohydrate chains, and proteinaceous components are all responsible for the cross-reactivity of trypanosome variant surface glycoproteins.
The C-terminal region of <i>Trypanosoma cruzi</i> MASPs is antigenic and secreted via exovesicles.
Adaptation and Therapeutic Exploitation of the Plasma Membrane of African Trypanosomes.

Molecular players of homologous recombination in protozoan parasites: implications for generating antigenic variation.
Recombination-driven generation of the largest pathogen repository of antigen variants in the protozoan <i>Trypanosoma cruzi</i> .
Extracellular vesicles of trypomastigotes of <i>Trypanosoma cruzi</i> induce changes in ubiquitin-related processes, cell-signaling pathways and apoptosis.
The major surface glycoprotein of <i>Trypanosoma cruzi</i> amastigotes are ligands of the human serum mannose-binding protein.
Gene and Chromosomal Copy Number Variations as an Adaptive Mechanism Towards a Parasitic Lifestyle in Trypanosomatids.
Deletion of a Golgi protein in <i>Trypanosoma cruzi</i> reveals a critical role for Mn2+ in protein glycosylation needed for host cell invasion and intracellular replication.
Infections With Extracellular Trypanosomes Require Control by Efficient Innate Immune Mechanisms and Can Result in the Destruction of the Mammalian Humoral Immune System.
Developmental competence and antigen switch frequency can be uncoupled in <i>Trypanosoma brucei</i> .
Trypanosomiasis-induced B cell apoptosis results in loss of protective anti-parasite antibody responses and abolishment of vaccine-induced memory responses.
Endosomal localization of the serum resistance-associated protein in African trypanosomes confers human infectivity.
Keeping Balance Between Genetic Stability and Plasticity at the Telomere and Subtelomere of <i>Trypanosoma brucei</i> .
The nuclear envelope and gene organization in parasitic protozoa: Specializations associated with disease.
Large-scale conformational changes of <i>Trypanosoma cruzi</i> proline racemase predicted by accelerated molecular dynamics simulation.
Developmental and morphological regulation of clathrin-mediated endocytosis in <i>Trypanosoma brucei</i> .
Natural programmed cell death in <i>T. cruzi</i> epimastigotes maintained in axenic cultures.
TbRAP1 has an unusual duplex DNA binding activity required for its telomere localization and VSG silencing.
Immuno-metabolic profile of human macrophages after <i>Leishmania</i> and <i>Trypanosoma cruzi</i> infection.
Inflammatory and Pro-resolving Lipids in Trypanosomatid Infections: A Key to Understanding Parasite Control.
Effect of the Tc13Tul antigen from <i>Trypanosoma cruzi</i> on splenocytes from naïve mice.
Evidence for recycling of invariant surface transmembrane domain proteins in African trypanosomes.
The History of the ABC Proteins in Human Trypanosomiasis Pathogens.
Involvement of SNARE protein Ykt6 in glycosome biogenesis in <i>Trypanosoma brucei</i> .
Cohesin regulates VSG monoallelic expression in trypanosomes.
The Unsolved Jigsaw Puzzle of the Immune Response in Chagas Disease.
Does <i>Trypanosoma cruzi</i> calreticulin modulate the complement system and angiogenesis?
Peroxioredoxins play a major role in protecting <i>Trypanosoma cruzi</i> against macrophage- and endogenously-derived peroxynitrite.
A proteomics approach reveals molecular manipulators of distinct cellular processes in the salivary glands of <i>Glossina m. morsitans</i> in response to <i>Trypanosoma b. brucei</i> infections.
Oral Versus Intragastric Inoculation: Similar Pathways of <i>Trypanosoma cruzi</i> Experimental Infection? From Target Tissues, Parasite Evasion, and Immune Response.

The Trypanosome-Derived Metabolite Indole-3-Pyruvate Inhibits Prostaglandin Production in Macrophages by Targeting COX2.
Telomeric expression sites are highly conserved in <i>Trypanosoma brucei</i> .
<i>T. brucei</i> infections abrogate diverse plasma cell-mediated effector B cell responses, independently of their specificity, affinity and host genetic background.
A phosphoproteomic approach towards the understanding of the role of TGF- $\beta$ in <i>Trypanosoma cruzi</i> biology.
Identification of immunodominant epitopes in <i>Trypanosoma cruzi</i> trypomastigote surface antigen-1 protein that mask protective epitopes.
Immune evasion strategies of trypanosomes: a review.
VEX1 controls the allelic exclusion required for antigenic variation in trypanosomes.
Role of Virulence Factors of Trypanosomatids in the Insect Vector and Putative Genetic Events Involved in Surface Protein Diversity.
Polymorphism in the subtelomeric regions of chromosomes of Kinetoplastida.
A recombinant protein based on <i>Trypanosoma cruzi</i> P21 enhances phagocytosis.
The <i>Trypanosoma cruzi</i> protease cruzain mediates immune evasion.
Inhibition of <i>Trypanosoma cruzi</i> proline racemase affects host-parasite interactions and the outcome of in vitro infection.
Identification and Characterization of the <i>Trypanosoma cruzi</i> B-cell Superantigen Tc24.
Cardiomyocyte diffusible redox mediators control <i>Trypanosoma cruzi</i> infection: role of parasite mitochondrial iron superoxide dismutase.
Clinical and Neuropathogenetic Aspects of Human African Trypanosomiasis.
Autoimmunity in Chronic Chagas Disease: A Road of Multiple Pathways to Cardiomyopathy?
Cryo-EM structures of <i>Trypanosoma brucei</i> gambiense ISG65 with human complement C3 and C3b and their roles in alternative pathway restriction.
Microvesicles released during the interaction between <i>Trypanosoma cruzi</i> TcI and TcII strains and host blood cells inhibit complement system and increase the infectivity of metacyclic forms of host cells in a strain-independent process.
Identification of paralogous life-cycle stage specific cytoskeletal proteins in the parasite <i>Trypanosoma brucei</i> .
Antigenic variation: extending the reach of telomeric silencing.
The Trypanosome Exocyst: A Conserved Structure Revealing a New Role in Endocytosis.
Cruzipain, the major cysteine protease of <i>Trypanosoma cruzi</i> : a sulfated glycoprotein antigen as relevant candidate for vaccine development and drug target. A review.
<i>Trypanosoma brucei</i> RAP1 Has Essential Functional Domains That Are Required for Different Protein Interactions.
The Interactions of Parasite Calreticulin With Initial Complement Components: Consequences in Immunity and Virulence.
Clathrin-mediated endocytosis is essential in <i>Trypanosoma brucei</i> .
Silencing subtelomeric VSGs by <i>Trypanosoma brucei</i> RAP1 at the insect stage involves chromatin structure changes.
The importance of mosaic genes to trypanosome survival.
Recombinant glycoprotein 63 (Gp63) of <i>Trypanosoma carassii</i> suppresses antimicrobial responses of goldfish ( <i>Carassius auratus</i> L.) monocytes and macrophages.
Interactions among <i>Trypanosoma brucei</i> RAD51 paralogues in DNA repair and antigenic variation.
A nuclear enterprise: zooming in on nuclear organization and gene expression control in the African trypanosome.

The Mitogen-Activated Protein Kinase (MAPK) Pathway: Role in Immune Evasion by Trypanosomatids.
ATR Kinase Is a Crucial Player Mediating the DNA Damage Response in <i>Trypanosoma brucei</i> .
Mouse models for pathogenic African trypanosomes: unravelling the immunology of host-parasite-vector interactions.
Trypanosomal immune evasion, chronicity and transmission: an elegant balancing act.
Chagas disease: the challenge of polyparasitism?
Role of membrane-bound IgM in <i>Trypanosoma cruzi</i> evasion from immune clearance.
Extracellular vesicles in the context of chagas disease - A systematic review.
An exploration of the genetic robustness landscape of surface protein families in the human protozoan parasite <i>Trypanosoma cruzi</i> .
The MASP family of <i>Trypanosoma cruzi</i> : changes in gene expression and antigenic profile during the acute phase of experimental infection.
PCR-based diagnosis of surra-targeting VSG gene: experimental studies in small laboratory rodents and buffalo.
Trans-sialidase recombinant protein mixed with CpG motif-containing oligodeoxynucleotide induces protective mucosal and systemic trypanosoma cruzi immunity involving CD8+ CTL and B cell-mediated cross-priming.
Alteration of macrophage function by a <i>Trypanosoma cruzi</i> membrane mucin.
Genome maintenance functions of a putative <i>Trypanosoma brucei</i> translesion DNA polymerase include telomere association and a role in antigenic variation.
Biochemical characterization of proline racemases from the human protozoan parasite <i>Trypanosoma cruzi</i> and definition of putative protein signatures.
TatD DNases of African trypanosomes confer resistance to host neutrophil extracellular traps.
High-throughput hit-squad tackles trypanosomes.
Activation of endocytosis as an adaptation to the mammalian host by trypanosomes.
A comparative analysis of trypanosomatid SNARE proteins.
The structure of serum resistance-associated protein and its implications for human African trypanosomiasis.
An immunoinformatic approach for identification of <i>Trypanosoma cruzi</i> HLA-A2-restricted CD8(+) T cell epitopes.
IFN- $\gamma$ mediates early B-cell loss in experimental African trypanosomiasis.
Structure of trypanosome coat protein VSG <sub>sur</sub> and function in suramin resistance.
Microbial antigenic variation mediated by homologous DNA recombination.
Linking the antigen archive structure to pathogen fitness in African trypanosomes.
A monoallelic deletion of the TcCRT gene increases the attenuation of a cultured <i>Trypanosoma cruzi</i> strain, protecting against an in vivo virulent challenge.
Genome-wide analysis reveals extensive functional interaction between DNA replication initiation and transcription in the genome of <i>Trypanosoma brucei</i> .
Revisiting trypanosomatid nucleoside diphosphate kinases.
History of sleeping sickness (African trypanosomiasis).
The <i>Trypanosoma brucei</i> -Derived Ketoacids, Indole Pyruvate and Hydroxyphenylpyruvate, Induce HO-1 Expression and Suppress Inflammatory Responses in Human Dendritic Cells.
Trypanosome motion represents an adaptation to the crowded environment of the vertebrate bloodstream.
cAMP signalling in trypanosomatids: role in pathogenesis and as a drug target.
mSphere of Influence: Expanding the CRISPR Sphere with Single-Locus Proteomics.

The role of invariant surface glycoprotein 75 in xenobiotic acquisition by African trypanosomes.
High-throughput chemical screening for antivirulence developmental phenotypes in <i>Trypanosoma brucei</i> .
Effect of the saliva from different triatomine species on the biology and immunity of TLR-4 ligand and <i>Trypanosoma cruzi</i> -stimulated dendritic cells.
Secreted trypanosome cyclophilin inactivates lytic insect defense peptides and induces parasite calcineurin activation and infectivity.
Structure of the trypanosome transferrin receptor reveals mechanisms of ligand recognition and immune evasion.
Mitochondrial dual-coding genes in <i>Trypanosoma brucei</i> .
Expression of co-stimulatory molecules CD80 and CD86 is altered in CD14 + HLA-DR + monocytes from patients with Chagas disease following induction by <i>Trypanosoma cruzi</i> recombinant antigens.
Nanobodies As Tools to Understand, Diagnose, and Treat African Trypanosomiasis.
Identification of genes encoding hypothetical proteins in open-reading frame expressed sequence tags from mammalian stages of <i>Trypanosoma cruzi</i> .
Ubiquitylation and developmental regulation of invariant surface protein expression in trypanosomes.
Spatial integration of transcription and splicing in a dedicated compartment sustains monogenic antigen expression in African trypanosomes.
A B-cell mitogen from a pathogenic trypanosome is a eukaryotic proline racemase.
Preclinical advances and the immunophysiology of a new therapeutic Chagas disease vaccine.
ENTH and ANTH domain proteins participate in AP2-independent clathrin-mediated endocytosis.
Mechanisms of complement lectin pathway activation and resistance by trypanosomatid parasites.
DNA Double-Strand Breaks: A Double-Edged Sword for Trypanosomatids.
Genome organization and DNA accessibility control antigenic variation in trypanosomes.
<i>Trypanosoma brucei</i> ribonuclease H2A is an essential R-loop processing enzyme whose loss causes DNA damage during transcription initiation and antigenic variation.
Segregation of minichromosomes in trypanosomes: implications for mitotic mechanisms.
<i>Trypanosoma cruzi</i> trans-sialidase prevents elicitation of Th1 cell response via interleukin 10 and downregulates Th1 effector cells.
Shedding light on lipid metabolism in Kinetoplastida: A phylogenetic analysis of phospholipase D protein homologs.
Relevance of the diversity among members of the <i>Trypanosoma cruzi</i> trans-sialidase family analyzed with camelids single-domain antibodies.
F(ab') <sub>2</sub> antibody fragments against <i>Trypanosoma cruzi</i> calreticulin inhibit its interaction with the first component of human complement.
Mucin AgC10 from <i>Trypanosoma cruzi</i> Interferes with L-selectin-mediated monocyte adhesion.
The endocytic activity of the flagellar pocket in <i>Trypanosoma brucei</i> is regulated by an adjacent phosphatidylinositol phosphate kinase.
Do trypanosome turncoats wait before they commit?
The trypanosome flagellar pocket.
<i>Trypanosoma brucei</i> gambiense excreted/secreted factors impair lipopolysaccharide-induced maturation and activation of human monocyte-derived dendritic cells.
Immunosuppression in Experimental Chagas Disease Is Mediated by an Alteration of Bone Marrow Stromal Cell Function During the Acute Phase of Infection.
Hepatocyte-derived IL-10 plays a crucial role in attenuating pathogenicity during the chronic phase of <i>T. congolense</i> infection.
Intracellular trafficking in the trypanosomatids.

Identification of a novel UDP-sugar pyrophosphorylase with a broad substrate specificity in <i>Trypanosoma cruzi</i> .
Multiparameter Optimization of Trypanocidal Cruzain Inhibitors With In Vivo Activity and Favorable Pharmacokinetics.
Experimental murine <i>Trypanosoma congolense</i> infections. II. Role of splenic adherent CD3+Thy1.2+ TCR-alpha beta- gamma delta- CD4+8- and CD3+Thy1.2+ TCR-alpha beta- gamma delta- CD4-8- cells in the production of IL-4, IL-10, and IFN-gamma and in trypanosome-elicited immunosuppression.
Sexual reproduction in a natural <i>Trypanosoma cruzi</i> population.
Gene conversion is a convergent strategy for pathogen antigenic variation.
Molecular Cloning, Expression and Characterization of Para Flagellar Rod Protein 1 of <i>Trypanosoma evansi</i> .
Membrane domains and flagellar pocket boundaries are influenced by the cytoskeleton in African trypanosomes.
Differentiation-associated surface antigen variation in the ancient eukaryote <i>Giardia lamblia</i> .
Identification of a novel base J binding protein complex involved in RNA polymerase II transcription termination in trypanosomes.
Inhibition of <i>Trypanosoma evansi</i> Protein-Tyrosine Phosphatase by Myristic Acid Analogues Isolated from <i>Khaya senegalensis</i> and <i>Tamarindus indica</i> .
Discordant antibody and cellular responses to <i>Pneumocystis</i> major surface glycoprotein variants in mice.
<i>Trypanosoma evansi</i> evades host innate immunity by releasing extracellular vesicles to activate TLR2-AKT signaling pathway.
Evidence of the Red-Queen Hypothesis from Accelerated Rates of Evolution of Genes Involved in Biotic Interactions in <i>Pneumocystis</i> .
Diversity and Complexity of the Large Surface Protein Family in the Compacted Genomes of Multiple <i>Pneumocystis</i> Species.
Histone Modifications and Other Facets of Epigenetic Regulation in Trypanosomatids: Leaving Their Mark.
Characterization of <i>pneumocystis</i> major surface glycoprotein gene (msg) promoter activity in <i>Saccharomyces cerevisiae</i> .
The G4 genome.

Table A2-32, Cluster 31

Cluster 31 focuses on the malaria parasite <i>Plasmodium falciparum</i> , emphasizing pathogenicity and immune evasion (509)
Small variant surface antigens and <i>Plasmodium</i> evasion of immunity.
More than just immune evasion: Hijacking complement by <i>Plasmodium falciparum</i> .
The selection landscape of malaria parasites.
Sticking for a Cause: The <i>Falciparum</i> Malaria Parasites Cytoadherence Paradigm.
Molecules on the surface of the <i>Plasmodium falciparum</i> infected erythrocyte and their role in malaria pathogenesis and immune evasion.
PfEMP1: an antigen that plays a key role in the pathogenicity and immune evasion of the malaria parasite <i>Plasmodium falciparum</i> .
Recombination and Diversification of the Variant Antigen Encoding Genes in the Malaria Parasite <i>Plasmodium falciparum</i> .

Epigenetic memory at malaria virulence genes.
Successful Profiling of Plasmodium falciparum var Gene Expression in Clinical Samples via a Custom Capture Array.
A coordinated transcriptional switching network mediates antigenic variation of human malaria parasites.
The epigenetic control of antigenic variation in Plasmodium falciparum.
Pathogenesis of Plasmodium falciparum malaria: the roles of parasite adhesion and antigenic variation.
Mosquito Vectors and the Globalization of Plasmodium falciparum Malaria.
Antigenic variation and immune evasion in Plasmodium falciparum malaria.
PfSETvs methylation of histone H3K36 represses virulence genes in Plasmodium falciparum.
PfSET2 Is Involved in Genome Organization of Var Gene Family in Plasmodium falciparum.
Parasite adhesion and immune evasion in placental malaria.
Frequent ectopic recombination of virulence factor genes in telomeric chromosome clusters of P. falciparum.
A var gene promoter controls allelic exclusion of virulence genes in Plasmodium falciparum malaria.
Plasmodium falciparum: epigenetic control of var gene regulation and disease.
Surface antigens of Plasmodium falciparum-infected erythrocytes as immune targets and malaria vaccine candidates.
The molecular basis for the cytoadherence of Plasmodium falciparum-infected erythrocytes to endothelium.
Transcriptional variation in the malaria parasite Plasmodium falciparum.
The varieties of gene amplification, diversification and hypervariability in the human malaria parasite, Plasmodium falciparum.
Prognostic indicators of life-threatening malaria are associated with distinct parasite variant antigen profiles.
Immune evasion of Plasmodium falciparum by RIFIN via inhibitory receptors.
Mutually exclusive expression of virulence genes by malaria parasites is regulated independently of antigen production.
Antigenic Variation in Plasmodium falciparum.
Regulation of antigenic variation in Plasmodium falciparum: censoring freedom of expression?
Molecular basis for evasion of host immunity and pathogenesis in malaria.
Conserved associations between G-quadruplex-forming DNA motifs and virulence gene families in malaria parasites.
Heterochromatin protein 1 secures survival and transmission of malaria parasites.
Variant antigen gene expression in malaria.
Plasmodium falciparum PfRUVBL proteins bind at the TARE region and var gene promoter located in the subtelomeric region.
A Unique Virulence Gene Occupies a Principal Position in Immune Evasion by the Malaria Parasite Plasmodium falciparum.
Molecular mechanisms of Plasmodium falciparum placental adhesion.
Mechanisms underlying mutually exclusive expression of virulence genes by malaria parasites.
The effects of a partitioned var gene repertoire of Plasmodium falciparum on antigenic diversity and the acquisition of clinical immunity.
Malaria vaccines.
Variant surface antigens of Plasmodium falciparum and their roles in severe malaria.

Genome sequence of the human malaria parasite <i>Plasmodium falciparum</i> .
Antigenic variation at the infected red cell surface in malaria.
Mechanisms for Host Immune Evasion Mediated by <i>Plasmodium falciparum</i> -Infected Erythrocyte Surface Antigens.
Three Is a Crowd - New Insights into Rosetting in <i>Plasmodium falciparum</i> .
Host-Parasite Interactions in Human Malaria: Clinical Implications of Basic Research.
<i>Plasmodium falciparum</i> erythrocyte invasion: combining function with immune evasion.
The human malaria parasite <i>Plasmodium falciparum</i> can sense environmental changes and respond by antigenic switching.
Ectopic recombination of a malaria var gene during mitosis associated with an altered var switch rate.
<i>Plasmodium falciparum</i> transcriptome analysis reveals pregnancy malaria associated gene expression.
Transcriptome profiling reveals functional variation in <i>Plasmodium falciparum</i> parasites from controlled human malaria infection studies.
Malaria: immune evasion by parasites.
<i>Plasmodium knowlesi</i> -mediated zoonotic malaria: A challenge for elimination.
Fine-scale genetic characterization of <i>Plasmodium falciparum</i> chromosome 7 encompassing the antigenic var and the drug-resistant pfcrt genes.
Host immune evasion strategies of malaria blood stage parasite.
Heterochromatin silencing and locus repositioning linked to regulation of virulence genes in <i>Plasmodium falciparum</i> .
Uncovering the role of Rad51 in homologous recombination-mediated antigenic diversification in the human malaria parasite <i>Plasmodium falciparum</i> .
VAR2CSA-Mediated Host Defense Evasion of <i>Plasmodium falciparum</i> Infected Erythrocytes in Placental Malaria.
Generation of antigenic diversity in <i>Plasmodium falciparum</i> by structured rearrangement of Var genes during mitosis.
New var reconstruction algorithm exposes high var sequence diversity in a single geographic location in Mali.
A large gene family for putative variant antigens shared by human and rodent malaria parasites.
<i>Plasmodium falciparum</i> Secretome in Erythrocyte and Beyond.
Malaria parasite interactions with the human host.
Malaria pathogenesis.
Global genetic diversity and evolution of var genes associated with placental and severe childhood malaria.
Acquisition of human plasminogen facilitates complement evasion by the malaria parasite <i>Plasmodium falciparum</i> .
<i>Plasmodium falciparum</i> virulence determinants unveiled.
Antigenic variation in <i>Plasmodium falciparum</i> malaria involves a highly structured switching pattern.
Host-parasite interactions during <i>Plasmodium</i> infection: Implications for immunotherapies.
Some strains of <i>Plasmodium falciparum</i> , a human malaria parasite, evade the complement-like system of <i>Anopheles gambiae</i> mosquitoes.
Immunoglobulin M: Restraint of Inflammation and Mediator of Immune Evasion by <i>Plasmodium falciparum</i> Malaria.
Characterization of a <i>Plasmodium falciparum</i> PHISTc protein, PF3D7_0801000, in blood- stage malaria parasites.
Insulator-like pairing elements regulate silencing and mutually exclusive expression in the malaria parasite <i>Plasmodium falciparum</i> .



Selecting Plasmodium falciparum-Infected Erythrocytes for Adhesion to Recombinant Receptors Under Flow Conditions.
A greedy promoter controls malarial var-iations.
Immune Response and Evasion Mechanisms of Plasmodium falciparum Parasites.
Revisiting the multigene families: Plasmodium var and vir genes.
Intra-cluster recombination and var transcription switches in the antigenic variation of Plasmodium falciparum.
Role of Pfs47 in the dispersal of ancestral Plasmodium falciparum malaria through adaptation to different anopheline vectors.
Genetic diversity of Plasmodium falciparum erythrocyte membrane protein 1 in field isolates from central Myanmar.
Analysis of subtelomeric virulence gene families in Plasmodium falciparum by comparative transcriptional profiling.
Variant Gene Expression and Antigenic Variation by Malaria Parasites.
[Immunology in medical practice. XXVII. Vaccines against malaria: new perspectives].
Mitotic evolution of Plasmodium falciparum shows a stable core genome but recombination in antigen families.
Cross-reactive immune responses as primary drivers of malaria chronicity.
RBC barcoding allows for the study of erythrocyte population dynamics and P. falciparum merozoite invasion.
Binding of human serum proteins to Plasmodium falciparum-infected erythrocytes and its association with malaria clinical presentation.
Whole-genome analysis of Malawian Plasmodium falciparum isolates identifies possible targets of allele-specific immunity to clinical malaria.
The role of epigenetics and chromatin structure in transcriptional regulation in malaria parasites.
The remarkable journey of adaptation of the Plasmodium falciparum malaria parasite to New World anopheline mosquitoes.
Changes in genome organization of parasite-specific gene families during the Plasmodium transmission stages.
Major Histocompatibility Complex and Malaria: Focus on Plasmodium vivax Infection.
Characterization of the repertoire diversity of the Plasmodium falciparum stevor multigene family in laboratory and field isolates.
Analysis of Plasmodium falciparum Rh2b deletion polymorphism across different transmission areas.
Variable var transition rates underlie antigenic variation in malaria.
In vivo transcriptome of Plasmodium falciparum reveals overexpression of transcripts that encode surface proteins.
Towards an anti-disease malaria vaccine.
Identification of a cis-acting DNA-protein interaction implicated in singular var gene choice in Plasmodium falciparum.
Positive selection of Plasmodium falciparum parasites with multiple var2csa-type PfEMP1 genes during the course of infection in pregnant women.
Regulation of immune response by Plasmodium-infected red blood cells.
Evolutionary analysis of the most polymorphic gene family in falciparum malaria.
Erythrocyte-binding antigens of Plasmodium falciparum are targets of human inhibitory antibodies and function to evade naturally acquired immunity.
Antigenic variation in Babesia bovis: how similar is it to that in Plasmodium falciparum?
Expression of P. falciparum var genes involves exchange of the histone variant H2A.Z at the promoter.

Assaying Interactions Between Neutrophils and Plasmodium falciparum-Infected Red Blood Cells.
Strand-specific RNA sequencing in Plasmodium falciparum malaria identifies developmentally regulated long non-coding RNA and circular RNA.
Plasmodium berghei-Released Factor, PbTIP, Modulates the Host Innate Immune Responses.
Programmed transcription of the var gene family, but not of stevor, in Plasmodium falciparum gametocytes.
CRISPR Interference of a Clonally Variant GC-Rich Noncoding RNA Family Leads to General Repression of var Genes in Plasmodium falciparum.
Plasmodium evasion of mosquito immunity and global malaria transmission: The lock-and-key theory.
Immune Escape Strategies of Malaria Parasites.
Population genomics of the immune evasion (var) genes of Plasmodium falciparum.
CRISPR/Cas9 Genome Editing Reveals That the Intron Is Not Essential for var2csa Gene Activation or Silencing in Plasmodium falciparum.
The Medicines for Malaria Venture Malaria Box contains inhibitors of protein secretion in Plasmodium falciparum blood stage parasites.
The evolutionary consequences of blood-stage vaccination on the rodent malaria Plasmodium chabaudi.
Antibody Levels to Plasmodium falciparum Erythrocyte Membrane Protein 1-DBLy11 and DBLδ-1 Predict Reduction in Parasite Density.
Differential expression of var subgroups and PfSir2a genes in afebrile Plasmodium falciparum malaria: a matched case-control study.
Malaria Vaccine Development: The Need for Novel Approaches: A Review Article.
Comprehensive analysis of antibody responses to Plasmodium falciparum erythrocyte membrane protein 1 domains.
Identification of Plasmodium falciparum var1CSA and var2CSA domains that bind IgM natural antibodies.
Immune evasion by malaria parasites: a challenge for vaccine development.
Novel insights into the role of long non-coding RNA in the human malaria parasite, Plasmodium falciparum.
Allelic polymorphisms in apical membrane antigen-1 are responsible for evasion of antibody-mediated inhibition in Plasmodium falciparum.
Additional Feeding Reveals Differences in Immune Recognition and Growth of Plasmodium Parasites in the Mosquito Host.
Increased melanizing activity in Anopheles gambiae does not affect development of Plasmodium falciparum.
'2TM proteins': an antigenically diverse superfamily with variable functions and export pathways.
Genome sequence, transcriptome, and annotation of rodent malaria parasite Plasmodium yoelii nigeriensis N67.
A nuclear redox sensor modulates gene activation and var switching in Plasmodium falciparum.
Exonuclease-mediated degradation of nascent RNA silences genes linked to severe malaria.
Plasmodium gene regulation: far more to factor in.
Parasite antigens on the infected red cell surface are targets for naturally acquired immunity to malaria.
Immune evasion by Babesia bovis and Plasmodium falciparum: cliff-dwellers of the parasite world.
Unveiling the Sugary Secrets of Plasmodium Parasites.
Comparative Genomics and Systems Biology of Malaria Parasites Plasmodium.
Plasmodium P47: a key gene for malaria transmission by mosquito vectors.

Rapid turnover of Plasmodium falciparum var gene transcripts and genotypes during natural non-symptomatic infections.
A systematic map of genetic variation in Plasmodium falciparum.
Variant surface antigens of malaria parasites: functional and evolutionary insights from comparative gene family classification and analysis.
Genome-wide patterns of genetic polymorphism and signatures of selection in Plasmodium vivax.
Genome-wide SNP analysis of Plasmodium falciparum shows differentiation at drug-resistance-associated loci among malaria transmission settings in southern Mali.
Antisense long noncoding RNAs regulate var gene activation in the malaria parasite Plasmodium falciparum.
Patterns and dynamics of genetic diversity in Plasmodium falciparum: what past human migrations tell us about malaria.
PfAP2-MRP DNA-binding protein is a master regulator of parasite pathogenesis during malaria parasite blood stages.
Structure of the Pf12 and Pf41 heterodimeric complex of Plasmodium falciparum 6-cysteine proteins.
[Genotyping of the Plasmodium falciparum msp1 (block 2) and dhfr (codon 108) genes in field samples collected in four endemic Colombian localities].
The immunological balance between host and parasite in malaria.
Silence, activate, poise and switch! Mechanisms of antigenic variation in Plasmodium falciparum.
Phylogeography of var gene repertoires reveals fine-scale geospatial clustering of Plasmodium falciparum populations in a highly endemic area.
The Architectural Factor HMGB1 Is Involved in Genome Organization in the Human Malaria Parasite Plasmodium falciparum.
Structural and immunological correlations between the variable blocks of the VAR2CSA domain DBL6 $\epsilon$ from two Plasmodium falciparum parasite lines.
Variability and exclusion in host and parasite: epigenetic regulation of Ig and var expression.
Variation in susceptibility of African Plasmodium falciparum malaria parasites to TEP1 mediated killing in Anopheles gambiae mosquitoes.
Inferring malaria parasite population structure from serological networks.
Nuclear pores and perinuclear expression sites of var and ribosomal DNA genes correspond to physically distinct regions in Plasmodium falciparum.
Novel Strategies for Malaria Vaccine Design.
PIMMS43 is required for malaria parasite immune evasion and sporogonic development in the mosquito vector.
Variation in use of erythrocyte invasion pathways by Plasmodium falciparum mediates evasion of human inhibitory antibodies.
Low-Complexity Repetitive Epitopes of Plasmodium falciparum Are Decoys for Humoural Immune Responses.
Phenotypic variation of Plasmodium falciparum merozoite proteins directs receptor targeting for invasion of human erythrocytes.
Complement in malaria: immune evasion strategies and role in protective immunity.
Identification of a vir-orthologous immune evasion gene family from primate malaria parasites.
Recombinant Duffy binding-like-alpha domains of Plasmodium falciparum erythrocyte membrane protein 1 elicit antibodies in rats that recognise conserved epitopes.
Latent transcriptional variations of individual Plasmodium falciparum uncovered by single-cell RNA-seq and fluorescence imaging.
Plasmodium falciparum var Gene Is Activated by Its Antisense Long Noncoding RNA.

The surface variant antigens of Plasmodium falciparum contain cross-reactive epitopes.
STEVR is a Plasmodium falciparum erythrocyte binding protein that mediates merozoite invasion and rosetting.
STRIDE: a command-line HMM-based identifier and sub-classifier of Plasmodium falciparum RIFIN and STEVR variant surface antigen families.
Accounting for red blood cell accessibility reveals distinct invasion strategies in Plasmodium falciparum strains.
Frequency-Dependent Competition Between Strains Imparts Persistence to Perturbations in a Model of Plasmodium falciparum Malaria Transmission.
Functional and immunological characterization of a Duffy binding-like alpha domain from Plasmodium falciparum erythrocyte membrane protein 1 that mediates rosetting.
Why functional pre-erythrocytic and bloodstage malaria vaccines fail: a meta-analysis of fully protective immunizations and novel immunological model.
Sequence polymorphism, segmental recombination and toggling amino acid residues within the DBL3X domain of the VAR2CSA placental malaria antigen.
The Form of an Antigen and Its Molecular Context Do Matter: Infectious versus Attenuated Plasmodium Sporozoites.
A Specific PfEMP1 Is Expressed in P. falciparum Sporozoites and Plays a Role in Hepatocyte Infection.
Chromodomain Protein Interacts with H3K9me3 and Controls RBC Rosette Formation by Regulating the Expression of a Subset of RIFINs in the Malaria Parasite.
Targeting SUMOylation in Plasmodium as a Potential Target for Malaria Therapy.
Transcription of the var genes from a freshly-obtained field isolate of Plasmodium falciparum shows more variable switching patterns than long laboratory-adapted isolates.
Manipulation of host innate immune responses by the malaria parasite.
Strategies for designing and monitoring malaria vaccines targeting diverse antigens.
Plasmodium species: master renovators of their host cells.
Rapid whole genome optical mapping of Plasmodium falciparum.
Structural basis for Duffy recognition by the malaria parasite Duffy-binding-like domain.
Selective expression of variant surface antigens enables Plasmodium falciparum to evade immune clearance in vivo.
Recruitment of Factor H as a Novel Complement Evasion Strategy for Blood-Stage Plasmodium falciparum Infection.
Plasmodium falciparum heterochromatin protein 1 marks genomic loci linked to phenotypic variation of exported virulence factors.
Virulence in malaria: an evolutionary viewpoint.
Malaria parasites require TLR9 signaling for immune evasion by activating regulatory T cells.
Attenuated P. falciparum Parasite Shows Cytokine Variations in Humanized Mice.
Extent and Dynamics of Polymorphism in the Malaria Vaccine Candidate Plasmodium falciparum Reticulocyte-Binding Protein Homologue-5 in Kalifabougou, Mali.
Protein microarray analysis of antibody responses to Plasmodium falciparum in western Kenyan highland sites with differing transmission levels.
Host age and expression of genes involved in red blood cell invasion in Plasmodium falciparum field isolates.
Identification and Characterization of the Antigens Expressed On the Surface of Human Erythrocytes Infected With Plasmodium falciparum.
Single-Cell RNA Sequencing Reveals Cellular Heterogeneity and Stage Transition under Temperature Stress in Synchronized Plasmodium falciparum Cells.

Malaria: could its unusual epigenome be the weak spot?
Origins and evolution of antigenic diversity in malaria parasites.
Transcriptional memory and switching in the Plasmodium falciparum rif gene family.
'Original antigenic sin', T cell memory, and malaria sporozoite immunity: an hypothesis for immune evasion.
Antigenic variation and the generation of diversity in malaria parasites.
The Plasmodium falciparum parasitophorous vacuole protein P113 interacts with the parasite protein export machinery and maintains normal vacuole architecture.
Plasmodium's journey through the Anopheles mosquito: A comprehensive review.
Dual regulatory effects of non-coding GC-rich elements on the expression of virulence genes in malaria parasites.
Epigenetic Variation and Regulation in Malaria Parasites.
Rifins: a second family of clonally variant proteins expressed on the surface of red cells infected with Plasmodium falciparum.
Antibodies to Peptides in Semiconserved Domains of RIFINs and STEVORs Correlate with Malaria Exposure.
The apical organelles of malaria merozoites: host cell selection, invasion, host immunity and immune evasion.
Exploring the virulence gene interactome with CRISPR/dCas9 in the human malaria parasite.
Merozoite surface protein 1, immune evasion, and vaccines against asexual blood stage malaria.
Comparative and functional genomics of the innate immune system in the malaria vector Anopheles gambiae.
Plasmodium falciparum parasites causing cerebral malaria share variant surface antigens, but are they specific?
Variable expression of the 235 kDa rhoptry protein of Plasmodium yoelii mediate host cell adaptation and immune evasion.
Analysis of the immune response of a new malaria vaccine based on the modification of cryptic epitopes.
Molecular Analysis of Pfs47-Mediated Plasmodium Evasion of Mosquito Immunity.
Recombination events among virulence genes in malaria parasites are associated with G-quadruplex-forming DNA motifs.
Plasmodium falciparum protein 'PfJ23' hosts distinct binding sites for major virulence factor 'PfEMP1' and Maurer's cleft marker 'PfSBP1'.
Antigenic variation in Plasmodium falciparum is associated with movement of var loci between subnuclear locations.
Molecular mechanisms of hematological and biochemical alterations in malaria: A review.
Immune effector mechanisms in malaria: An update focusing on human immunity.
The human malaria parasite Pfs47 gene mediates evasion of the mosquito immune system.
The Plasmodium falciparum blood stages acquire factor H family proteins to evade destruction by human complement.
[Genotypic survey of Plasmodium falciparum based on the msp1, msp2 and glurp genes by multiplex PCR].
Characterisation of Plasmodium falciparum populations selected on the human endothelial receptors P-selectin, E-selectin, CD9 and CD151.
The prime role of plasma membrane cholesterol in the pathogenesis of immune evasion and clinical manifestations of falciparum malaria.

Plasmodium falciparum variability and immune evasion proceed from antigenicity of consensus sequences from DBL6e; generalization to all DBL from VAR2CSA.
Genetic analysis of host-parasite coevolution in human malaria.
Plasmodium falciparum RIFIN is a novel ligand for inhibitory immune receptor LILRB2.
Malaria parasite epigenetics: when virulence and romance collide.
A comprehensive evaluation of rodent malaria parasite genomes and gene expression.
A critical role of perinuclear filamentous actin in spatial repositioning and mutually exclusive expression of virulence genes in malaria parasites.
Differential recognition of Plasmodium falciparum merozoite surface protein 2 variants by antibodies from malaria patients in Brazil.
Ten families of variant genes encoded in subtelomeric regions of multiple chromosomes of Plasmodium chabaudi, a malaria species that undergoes antigenic variation in the laboratory mouse.
Dissecting the Gene Expression, Localization, Membrane Topology, and Function of the Plasmodium falciparum STEVOR Protein Family.
The crystal structure of P. knowlesi DBPalpha DBL domain and its implications for immune evasion.
Putative DNA G-quadruplex formation within the promoters of Plasmodium falciparum var genes.
Unravelling a histone code for malaria virulence.
Trans-acting GC-rich non-coding RNA at var expression site modulates gene counting in malaria parasite.
Single-molecule imaging and quantification of the immune-variant adhesin VAR2CSA on knobs of Plasmodium falciparum-infected erythrocytes.
Evasion of Immunity to Plasmodium falciparum: Rosettes of Blood Group A Impair Recognition of PfEMP1.
Seroreactivity to Plasmodium falciparum erythrocyte membrane protein 1 intracellular domain in malaria-exposed children and adults.
IgM, Fc mu Rs, and malarial immune evasion.
The Plasmodium gaboni genome illuminates allelic dimorphism of immunologically important surface antigens in P. falciparum.
Large, rapidly evolving gene families are at the forefront of host-parasite interactions in Apicomplexa.
Antibodies targeting the PFRH1 binding domain inhibit invasion of Plasmodium falciparum merozoites.
Challenges in malaria control in sub-Saharan Africa: the vaccine perspective.
Repetitive sequences in malaria parasite proteins.
Leishmania donovani infection drives the priming of human monocyte-derived dendritic cells during Plasmodium falciparum co-infections.
Transcriptomic Analysis of Chloroquine-Sensitive and Chloroquine-Resistant Strains of Plasmodium falciparum: Toward Malaria Diagnostics and Therapeutics for Global Health.
Investigating a Plasmodium falciparum erythrocyte invasion phenotype switch at the whole transcriptome level.
Leveraging genome editing to functionally evaluate Plasmodium diversity.
Immune evasion strategies of pre-erythrocytic malaria parasites.
Expression dynamics and physiologically relevant functional study of STEVOR in asexual stages of Plasmodium falciparum infection.
Maurer's clefts, the enigma of Plasmodium falciparum.
Plasmodium falciparum has evolved multiple mechanisms to hijack human immunoglobulin M.
A model for the coevolution of immunity and immune evasion in vector-borne diseases with implications for the epidemiology of malaria.
Investigating Pharmacological Targeting of G-Quadruplexes in the Human Malaria Parasite.

Transient cross-reactive immune responses can orchestrate antigenic variation in malaria.
The Impact of Malaria Parasites on Dendritic Cell-T Cell Interaction.
RecQ helicases in the malaria parasite <i>Plasmodium falciparum</i> affect genome stability, gene expression patterns and DNA replication dynamics.
Malaria systems immunology: <i>Plasmodium vivax</i> induces tolerance during primary infection through dysregulation of neutrophils and dendritic cells.
Crystal structure of <i>Plasmodium knowlesi</i> apical membrane antigen 1 and its complex with an invasion-inhibitory monoclonal antibody.
Detection of positive selection acting on protein surfaces at the whole-genome scale in the human malaria parasite <i>Plasmodium falciparum</i> .
Sequence diversity and evolutionary dynamics of the dimorphic antigen merozoite surface protein-6 and other Msp genes of <i>Plasmodium falciparum</i> .
How selection forces dictate the variant surface antigens used by malaria parasites.
The Py235 proteins: glimpses into the versatility of a malaria multigene family.
Effect of infection by <i>Plasmodium falciparum</i> on the melanization immune response of <i>Anopheles gambiae</i> .
Bacterially expressed full-length recombinant <i>Plasmodium falciparum</i> RH5 protein binds erythrocytes and elicits potent strain-transcending parasite-neutralizing antibodies.
Progress in the development of recombinant and synthetic blood-stage malaria vaccines.
Dimorphic <i>Plasmodium falciparum</i> merozoite surface protein-1 epitopes turn off memory T cells and interfere with T cell priming.
A potential novel mechanism for the insertion of a membrane protein revealed by a biochemical analysis of the <i>Plasmodium falciparum</i> cytoadherence molecule PfEMP-1.
Malaria vaccine design: immunological considerations.
Mechanistic within-host models of the asexual <i>Plasmodium falciparum</i> infection: a review and analytical assessment.
[Method for rapid synchronization of different growth cycles of <i>Plasmodium falciparum</i> in vitro and application in differential gene expression profile of 3D7 after dihydroartemisinin treatment].
Biochemical characterization of <i>Plasmodium</i> complement factors binding protein for its role in immune modulation.
Illuminating how malaria parasites export proteins into host erythrocytes.
Nanobody generation and structural characterization of <i>Plasmodium falciparum</i> 6-cysteine protein Pf12p.
A mathematical model for a new mechanism of phenotypic variation in malaria.
Analysis by Flow Cytometry of $\alpha(2)$ -Macroglobulin and Nonimmune IgM-Binding to <i>Plasmodium falciparum</i> -Infected Erythrocytes.
Single-cell RNA-seq reveals hidden transcriptional variation in malaria parasites.
Filarial infection induces protection against <i>P. berghei</i> liver stages in mice.
Direct and indirect immunosuppression by a malaria parasite in its mosquito vector.
Mapping a common interaction site used by <i>Plasmodium falciparum</i> Duffy binding-like domains to bind diverse host receptors.
<i>Plasmodium cynomolgi</i> genome sequences provide insight into <i>Plasmodium vivax</i> and the monkey malaria clade.
Exposure of the <i>Plasmodium falciparum</i> clonally variant STEVOR proteins on the merozoite surface.
Short report: Positive correlation between rosetting and parasitemia in <i>Plasmodium falciparum</i> clinical isolates.

The Impact of Recruitment on the Dynamics of an Immune-Suppressed Within-Human-Host Model of the Plasmodium falciparum Parasite.
Defining species-specific and conserved interactions of apical membrane protein 1 during erythrocyte invasion in malaria to inform multi-species vaccines.
Polymorphism in liver-stage malaria vaccine candidate proteins: immune evasion and implications for vaccine design.
Plasmodium 6-Cysteine Proteins: Functional Diversity, Transmission-Blocking Antibodies and Structural Scaffolds.
Towards functional antibody-based vaccines to prevent pre-erythrocytic malaria infection.
Conflicting immune responses can prolong the length of infection in Plasmodium falciparum malaria.
Does malaria suffer from lack of memory?
Structural conservation despite huge sequence diversity allows EPCR binding by the PfEMP1 family implicated in severe childhood malaria.
An essential dual-function complex mediates erythrocyte invasion and channel-mediated nutrient uptake in malaria parasites.
The Plasmodium falciparum STEVOR multigene family mediates antigenic variation of the infected erythrocyte.
In vitro adaptation of Plasmodium falciparum reveal variations in cultivability.
Trans-kingdom small RNA transfer during host-pathogen interactions: The case of P. falciparum and erythrocytes.
Structural insights into diverse modes of ICAM-1 binding by Plasmodium falciparum-infected erythrocytes.
Nuclear repositioning precedes promoter accessibility and is linked to the switching frequency of a Plasmodium falciparum invasion gene.
Antibodies to a merozoite surface protein promote multiple invasion of red blood cells by malaria parasites.
CD47 regulates the phagocytic clearance and replication of the Plasmodium yoelii malaria parasite.
Parasite genomics: current status and future prospects.
Genome-Wide Scans for Ghanaian Plasmodium falciparum Genes Under Selection From Local and Chinese Host Populations.
An immunologically cryptic epitope of Plasmodium falciparum circumsporozoite protein facilitates liver cell recognition and induces protective antibodies that block liver cell invasion.
Immune Modulation and Prevention of Autoimmune Disease by Repeated Sequences from Parasites Linked to Self Antigens.
A Model to Study the Impact of Polymorphism Driven Liver-Stage Immune Evasion by Malaria Parasites, to Help Design Effective Cross-Reactive Vaccines.
A statistically rigorous method for determining antigenic switching networks.
The origin of malaria: mixed messages from genetic diversity.
Genotyping of MSP3 $\beta$ gene in Indian Plasmodium vivax.
The 235 kDa rhoptry protein of Plasmodium (yoelii) yoelii: function at the junction.
Genome plasticity in Plasmodium.
Analysis of pir gene expression across the Plasmodium life cycle.
Assessing the genetic diversity of the vir genes in Indian Plasmodium vivax population.
Dimorphism and intergenic recombination within the microneme protein (MP-1) gene family of Plasmodium knowlesi.
Surface Plasmon Resonance Analysis of PfEMP1 Interaction with Receptors.



Suppression of adaptive immunity to heterologous antigens during Plasmodium infection through hemozoin-induced failure of dendritic cell function.
High non-protective, long-lasting antibody levels in malaria are associated with haplotype shifting in MHC-peptide-TCR complex formation: a new mechanism for immune evasion.
Bioinformatics analysis of the structure and function of NADPH-cytochrome p450 reductase of Plasmodium vivax.
Differential antibody recognition of four allelic variants of the merozoite surface protein-2 (MSP-2) of Plasmodium falciparum.
Malaria in pregnancy: getting to grips with a sticky problem.
Varia: a tool for prediction, analysis and visualisation of variable genes.
Artemisinin-Ginkgo biloba extract combination therapy for Plasmodium yoelii.
Extensive genetic diversity of Plasmodium vivax dbp-II in Rio de Janeiro Atlantic Forest and Brazilian Amazon Basin: evidence of positive selection.
Plasmodium TatD-Like DNase Antibodies Blocked Parasite Development in the Mosquito Gut.
Plasmodium falciparum: Rosettes do not protect merozoites from invasion-inhibitory antibodies.
Hijacking the human complement inhibitor C4b-binding protein by the sporozoite stage of the Plasmodium falciparum parasite.
Identification and Immune Assessment of T Cell Epitopes in Five Plasmodium falciparum Blood Stage Antigens to Facilitate Vaccine Candidate Selection and Optimization.
Dimerization of Plasmodium vivax DBP is induced upon receptor binding and drives recognition of DARC.
Live attenuated rubella vectors expressing Plasmodium falciparum circumsporozoite protein (Pf-CSP) provide a novel malaria vaccine platform in the rhesus macaque.
Meta-analysis of immune epitope data for all Plasmodia: overview and applications for malarial immunobiology and vaccine-related issues.
Human immune recognition of recombinant proteins representing discrete domains of the Plasmodium falciparum gamete surface protein, Pfs230.
Plasmodium falciparum: red blood cell binding studies of peptides derived from histidine-rich KAHRP-I, HRP-II and HRP-III proteins.
Adapting immunity with subunit vaccines: case studies with group A Streptococcus and malaria.
Maurer's clefts: a novel multi-functional organelle in the cytoplasm of Plasmodium falciparum-infected erythrocytes.
Amplification of Duffy binding protein-encoding gene allows Plasmodium vivax to evade host anti-DBP humoral immunity.
Bioinformatics analysis and prediction for structure and function of nitric oxide synthase and similar proteins from Plasmodium berghei.
Malaria oocysts require circumsporozoite protein to evade mosquito immunity.
Proteomic analysis reveals novel proteins associated with the Plasmodium protein exporter PTEX and a loss of complex stability upon truncation of the core PTEX component, PTEX150.
Distinct genetic difference between the Duffy binding protein (PkDBPall) of Plasmodium knowlesi clinical isolates from North Borneo and Peninsular Malaysia.
Polymorphism patterns in Duffy-binding protein among Thai Plasmodium vivax isolates.
Size polymorphism and low sequence diversity in the locus encoding the Plasmodium vivax rhoptry neck protein 4 (PvRON4) in Colombian isolates.
Rosettes integrity protects Plasmodium vivax of being phagocytized.
Placental Malaria is Associated with Higher LILRB2 Expression in Monocyte Subsets and Lower Anti-Malarial IgG Antibodies During Infancy.

FT-GPI, a highly sensitive and accurate predictor of GPI-anchored proteins, reveals the composition and evolution of the GPI proteome in Plasmodium species.
Hemozoin-mediated inflammasome activation limits long-lived anti-malarial immunity.
Quantification of human complement factor H binding to asexual malaria blood stages by an enzyme-linked immunosorbent assay.
Widespread distribution of antisense transcripts in the Plasmodium falciparum genome.
Modified MHC Class II-Associated Invariant Chain Induces Increased Antibody Responses against Plasmodium falciparum Antigens after Adenoviral Vaccination.
The N-terminus of EXP2 forms the membrane-associated pore of the protein exporting translocon PTEX in Plasmodium falciparum.
Genomic Analysis of Plasmodium vivax in Southern Ethiopia Reveals Selective Pressures in Multiple Parasite Mechanisms.
Cryo-EM reveals the architecture of placental malaria VAR2CSA and provides molecular insight into chondroitin sulfate binding.
Functional analysis of epigenetic regulation of tandem RhopH1/clag genes reveals a role in Plasmodium falciparum growth.
Vaccination in a humanized mouse model elicits highly protective PfCSP-targeting anti-malarial antibodies.
Leptospirosis: Molecular trial path and immunopathogenesis correlated with dengue, malaria and mimetic hemorrhagic infections.
Genetic and epigenetic mechanisms underlying cell-surface variability in protozoa and fungi.
Genetically attenuated parasite vaccines induce contact-dependent CD8+ T cell killing of Plasmodium yoelii liver stage-infected hepatocytes.
Decoding the impact of nuclear organization on antigenic variation in parasites.
Identification of the Recombinant Plasmodium vivax Surface-Related Antigen as a Possible Immune Evasion Factor Against Human Splenic Fibroblasts by Targeting ITGB1.
Force Spectroscopy of the Plasmodium falciparum Vaccine Candidate Circumsporozoite Protein Suggests a Mechanically Pliable Repeat Region.
The effects of symmetry on the dynamics of antigenic variation.
Structural basis of antigenic escape of a malaria vaccine candidate.
Profoundly Reduced CD1c+ Myeloid Dendritic Cell HLA-DR and CD86 Expression and Increased Tumor Necrosis Factor Production in Experimental Human Blood-Stage Malaria Infection.
Independent regulation of Plasmodium falciparum rif gene promoters.
Neutralizing and interfering human antibodies define the structural and mechanistic basis for antigenic diversion.
Alanine mutagenesis of the primary antigenic escape residue cluster, c1, of apical membrane antigen 1.
Morphological and proteomic characterization of midgut of the malaria vector Anopheles albimanus at early time after a blood feeding.
Blood feast: Exploring the erythrocyte-feeding behaviour of the myxozoan Sphaerospora molnari.
Pathogenomics of Culex quinquefasciatus and meta-analysis of infection responses to diverse pathogens.
Plasmodium falciparum Sir2A preferentially hydrolyzes medium and long chain fatty acyl lysine.
Tim-3 signaling blockade with $\alpha$ -lactose induces compensatory TIGIT expression in Plasmodium berghei ANKA-infected mice.
What is the function of MSP-I on the malaria merozoite?
Incorporating genetic selection into individual-based models of malaria and other infectious diseases.

PfPRH2b specific monoclonal antibodies inhibit merozoite invasion.
Lipid hijacking: a unifying theme in vector-borne diseases.
Characterization and gene expression analysis of the cir multi-gene family of Plasmodium chabaudi chabaudi (AS).
Red blood cell invasion by Plasmodium vivax: structural basis for DBP engagement of DARC.
Effect of low complexity regions within the PvMSP3 $\alpha$ block II on the tertiary structure of the protein and implications to immune escape mechanisms.
Expression and strain variation of the novel "small open reading frame" (smorf) multigene family in Babesia bovis.
Revisiting the diffusion approximation to estimate evolutionary rates of gene family diversification.
Anopheles midgut epithelium evades human complement activity by capturing factor H from the blood meal.
Merozoite surface antigen 2 proteins of Babesia bovis vaccine breakthrough isolates contain a unique hypervariable region composed of degenerate repeats.
Rab GTPase regulation of bacteria and protozoa phagocytosis occurs through the modulation of phagocytic receptor surface expression.
Selection for antigenic diversity of Tams1, the major merozoite antigen of Theileria annulata.
Subdominance in Antibody Responses: Implications for Vaccine Development.
Pathogen escape from host immunity by a genome program for antigenic variation.
Vaccine design: emerging concepts and renewed optimism.
Malaria circumsporozoite protein inhibits protein synthesis in mammalian cells.
Selection of Babesia bovis-infected erythrocytes for adhesion to endothelial cells coselects for altered variant erythrocyte surface antigen isoforms.
Genetic polymorphism of vir genes of Plasmodium vivax in Myanmar.
Localisation of laminin within Plasmodium berghei oocysts and the midgut epithelial cells of Anopheles stephensi.
On the possible role of robustness in the evolution of infectious diseases.
Demonstration of the genetic stability and temporal expression of select members of the Lyme disease spirochete OspF protein family during infection in mice.
How do antigenically varying pathogens avoid cross-reactive responses to invariant antigens?
Molecular cloning and genetic polymorphism of Babesia capreoli gene Bcp37/41, an ortholog of Babesia divergens merozoite surface antigen Bd37.
Dynamics and stability: epigenetic conversions in position effect variegation.
Neuroinflammation and brain infections: historical context and current perspectives.
Molecular characterization of a 28 kDa surface antigen gene family of the tribe Ehrlichiae.
Ehrlichia ruminantium major antigenic protein gene (map1) variants are not geographically constrained and show no evidence of having evolved under positive selection pressure.
Variable and Variant Protein Multigene Families in Babesia bovis Persistence.
Recent advance of erythrocyte-mimicking nanovehicles: From bench to bedside.
Single-stranded DNA aptamers mask RhD antigenic epitopes on human RhD+ red blood cells to escape alloanti-RhD immunological recognition.
[Optimization of formulation of paclitaxel nanosuspension encapsulated by erythrocyte membrane based on Box-Behnken method].
Corrigendum: Mechanism of Rhinovirus Immunity and Asthma.

Table A2-33, Cluster 32

Cluster 32 focuses on breast cancer, emphasizing mechanisms of immune evasion and immunotherapy resistance (425)
Immune System Effects on Breast Cancer.
The Role of Immune Escape and Immune Cell Infiltration in Breast Cancer.
Circular RNA circWWC3 augments breast cancer progression through promoting M2 macrophage polarization and tumor immune escape via regulating the expression and secretion of IL-4.
Mechanisms of immune evasion in breast cancer.
The Immune Landscape of Breast Cancer: Strategies for Overcoming Immunotherapy Resistance.
The genomic landscape of breast cancer and its interaction with host immunity.
Targeted immunotherapy for HER2-low breast cancer with 17p loss.
The Complex Interaction between the Tumor Micro-Environment and Immune Checkpoints in Breast Cancer.
Immunotherapy: New insights in breast cancer treatment.
Icariin-induced inhibition of SIRT6/NF- $\kappa$ B triggers redox mediated apoptosis and enhances anti-tumor immunity in triple-negative breast cancer.
LncRNA TINCR impairs the efficacy of immunotherapy against breast cancer by recruiting DNMT1 and downregulating MiR-199a-5p via the STAT1-TINCR-USP20-PD-L1 axis.
Understanding the Role of Innate Immune Cells and Identifying Genes in Breast Cancer Microenvironment.
The immune system and inflammation in breast cancer.
Propensity for Early Metastatic Spread in Breast Cancer: Role of Tumor Vascularization Features and Tumor Immune Infiltrate.
The Great Immune Escape: Understanding the Divergent Immune Response in Breast Cancer Subtypes.
Impact of the Tumor Microenvironment on Tumor-Infiltrating Lymphocytes: Focus on Breast Cancer.
Insight into the Crosstalk between Photodynamic Therapy and Immunotherapy in Breast Cancer.
miRNA-1260b Promotes Breast Cancer Cell Migration and Invasion by Downregulating CCDC134.
Expression of toll-like receptors on breast tumors: taking a toll on tumor microenvironment.
PD-L1/PD-1 axis as a potent therapeutic target in breast cancer.
Cancer stem cell-mediated drug resistance: A comprehensive gene expression profile analysis in breast cancer.
Checkpoint inhibitors in triple-negative breast cancer (TNBC): Where to go from here.
Review of 10 years of research on breast cancer patients: Focus on indoleamine 2,3-dioxygenase.
DNA damage induces STING mediated IL-6-STAT3 survival pathway in triple-negative breast cancer cells and decreased survival of breast cancer patients.
Breast Cancer Immunity: It is TIME for the Next Chapter.
Immunotherapeutics for breast cancer.
Metabolism and immunity in breast cancer.
LncRNA TYMSOS facilitates breast cancer metastasis and immune escape through downregulating ULBP3.
Nanomedicine as a Promising Tool to Overcome Immune Escape in Breast Cancer.
Immunotherapy in Breast Cancer.
Semaphorins and Their Roles in Breast Cancer: Implications for Therapy Resistance.
Mechanisms of tumor immune escape in triple-negative breast cancers (TNBC) with and without mutated BRCA 1.
Sympathetic activity in breast cancer and metastasis: partners in crime.

Lactate from glycolysis regulates inflammatory macrophage polarization in breast cancer.
Laboratory Models for Investigating Breast Cancer Therapy Resistance and Metastasis.
Key Factor Regulating Inflammatory Microenvironment, Metastasis, and Resistance in Breast Cancer: Interleukin-1 Signaling.
Subgrouping breast cancer patients based on immune evasion mechanisms unravels a high involvement of transforming growth factor-beta and decoy receptor 3.
Macrophage balance fraction determines the degree of immunosuppression and metastatic ability of breast cancer.
Bone serves as a transfer station for secondary dissemination of breast cancer.
Identification of MicroRNAs as Diagnostic Biomarkers for Breast Cancer Based on the Cancer Genome Atlas.
The Extracellular Matrix and Vesicles Modulate the Breast Tumor Microenvironment.
The Biology of Exosomes in Breast Cancer Progression: Dissemination, Immune Evasion and Metastatic Colonization.
Therapeutic targeting of B7-H1 in breast cancer.
Immunology and breast cancer: therapeutic cancer vaccines.
Immunotherapy of breast cancer.
Alterations of Signaling Pathways Related to the Immune System in Breast Cancer: New Perspectives in Patient Management.
Neurotrophin-3 modulates breast cancer cells and the microenvironment to promote the growth of breast cancer brain metastasis.
Genetic variation in the immunosuppression pathway genes and breast cancer susceptibility: a pooled analysis of 42,510 cases and 40,577 controls from the Breast Cancer Association Consortium.
CD47 expression and tumor-associated immune cells in breast cancer and their correlation with molecular subtypes and prognostic factors.
A Rosetta Stone for Breast Cancer: Prognostic Value and Dynamic Regulation of Neutrophil in Tumor Microenvironment.
Single-cell RNA sequencing reveals cell heterogeneity and transcriptome profile of breast cancer lymph node metastasis.
Novel classes of immunotherapy for breast cancer.
Epigenetic Regulation of Immunotherapy Response in Triple-Negative Breast Cancer.
The onco-immunological implications of Fusobacterium nucleatum in breast cancer.
The biology of breast cancer.
Immune cell infiltration-based signature for prognosis and immunogenomic analysis in breast cancer.
The role of cytokines in breast cancer development and progression.
STAT family of transcription factors in breast cancer: Pathogenesis and therapeutic opportunities and challenges.
Polyloid giant cancer cells, cytokines and cytomegalovirus in breast cancer progression.
Molecular basis of invasion in breast cancer.
Therapeutic Potential of Lymphoid Infiltrates in Breast Cancer.
PPP2R2B downregulation is associated with immune evasion and predicts poor clinical outcomes in triple-negative breast cancer.
Genetic markers of immunoglobulin G and susceptibility to breast cancer.
Adipose Tissue-Derived Mesenchymal Stromal/Stem Cells, Obesity and the Tumor Microenvironment of Breast Cancer.
miR-195/miR-497 Regulate CD274 Expression of Immune Regulatory Ligands in Triple-Negative Breast Cancer.

Extracellular vesicle-mediated transport: Reprogramming a tumor microenvironment conducive with breast cancer progression and metastasis.
Insights into Immune Escape During Tumor Evolution and Response to Immunotherapy Using a Rat Model of Breast Cancer.
Triple negative breast cancer: Pitfalls and progress.
Immune Escape during Breast Tumor Progression.
The prognostic significance of human ovarian aging-related signature in breast cancer after surgery: A multicohort study.
The role of tumor-associated macrophage in breast cancer biology.
Regional Variation in the Tumor Microenvironment, Immune Escape and Prognostic Factors in Breast Cancer in Sub-Saharan Africa.
Counteracting Chemoresistance with Metformin in Breast Cancers: Targeting Cancer Stem Cells.
Tumor vaccines for breast cancer.
CD95 ligand expression as a mechanism of immune escape in breast cancer.
Immunizing against breast cancer: a new swing for an old sword.
T cell coinhibition and immunotherapy in human breast cancer.
Molecular mechanisms of immunotherapy resistance in triple-negative breast cancer.
Ly6E/K Signaling to TGF $\beta$ Promotes Breast Cancer Progression, Immune Escape, and Drug Resistance.
The long non-coding RNA GHSROS facilitates breast cancer cell migration and orthotopic xenograft tumour growth.
Chlorogenic acid induces apoptosis, inhibits metastasis and improves antitumor immunity in breast cancer via the NF- $\kappa$ B signaling pathway.
Integrative single-cell sequencing analysis distinguishes survival-associated cells from the breast cancer microenvironment.
[The anti-tumor immune response in breast cancer: Update and therapeutic perspectives].
Integrative Dissection of Novel Lactate Metabolism-Related Signature in the Tumor Immune Microenvironment and Prognostic Prediction in Breast Cancer.
Deregulated molecules and pathways in the predisposition and dissemination of breast cancer cells to bone.
Serum-derived extracellular vesicles from breast cancer patients contribute to differential regulation of T-cell-mediated immune-escape mechanisms in breast cancer subtypes.
Immunotherapy and targeted therapy combinations in metastatic breast cancer.
Targeting Stat3 induces senescence in tumor cells and elicits prophylactic and therapeutic immune responses against breast cancer growth mediated by NK cells and CD4+ T cells.
The role of CPT1A as a biomarker of breast cancer progression: a bioinformatic approach.
Exploring the Potential of Breast Microbiota as Biomarker for Breast Cancer and Therapeutic Response.
Beyond Immunosuppression: The Multifaceted Functions of Tumor-Promoting Myeloid Cells in Breast Cancers.
Standard of Care and Promising New Agents for the Treatment of Mesenchymal Triple-Negative Breast Cancer.
New horizons in breast cancer: the promise of immunotherapy.
Myeloid cells in circulation and tumor microenvironment of breast cancer patients.
Rationale for immunological approaches to breast cancer therapy.
Yin-yang effect of tumor infiltrating B cells in breast cancer: From mechanism to immunotherapy.
STAT3 Signaling in Breast Cancer: Multicellular Actions and Therapeutic Potential.

ILT4 reprograms glucose metabolism to promote tumor progression in triple negative breast cancer.
CIBERSORT analysis of TCGA and METABRIC identifies subgroups with better outcomes in triple negative breast cancer.
Developing an effective breast cancer vaccine.
Targeting adipocyte-immune cell crosstalk to control breast cancer progression.
Progesterone Receptor Attenuates STAT1-Mediated IFN Signaling in Breast Cancer.
Breast cancer stem cells and the immune system: promotion, evasion and therapy.
The ShcA adaptor protein is a critical regulator of breast cancer progression.
Lactate Metabolism and Immune Modulation in Breast Cancer: A Focused Review on Triple Negative Breast Tumors.
Immunoglobulin G expression and its potential role in primary and metastatic breast cancers.
RCAS1 is associated with ductal breast cancer progression.
Consequences of EMT-Driven Changes in the Immune Microenvironment of Breast Cancer and Therapeutic Response of Cancer Cells.
BET proteins in abnormal metabolism, inflammation, and the breast cancer microenvironment.
Host-Related Factors in the Interplay among Inflammation, Immunity and Dormancy in Breast Cancer Recurrence and Prognosis: An Overview for Clinicians.
STAT3 as a potential therapeutic target in triple negative breast cancer: a systematic review.
LncRNA GATA3-AS1 facilitates tumour progression and immune escape in triple-negative breast cancer through destabilization of GATA3 but stabilization of PD-L1.
Systemic Immune Dysregulation in Early Breast Cancer Is Associated With Decreased Plasma Levels of Both Soluble Co-Inhibitory and Co-Stimulatory Immune Checkpoint Molecules.
Peptide vaccines in early breast cancer.
Mechanisms of resistance of chemotherapy in early-stage triple negative breast cancer (TNBC).
Entinostat for the treatment of breast cancer.
Reduced expression of Toll-like receptor 4 inhibits human breast cancer cells proliferation and inflammatory cytokines secretion.
Microparticles shed from multidrug resistant breast cancer cells provide a parallel survival pathway through immune evasion.
Bioinformatic profiling of prognosis-related genes in the breast cancer immune microenvironment.
The Multifaceted Role of Regulatory T Cells in Breast Cancer.
Targeting hypoxia-inducible factor-1alpha: A new strategy for triple-negative breast cancer therapy.
Cognate Nonlytic Interactions between CD8(+) T Cells and Breast Cancer Cells Induce Cancer Stem Cell-like Properties.
Genetic Heterogeneity, Tumor Microenvironment and Immunotherapy in Triple-Negative Breast Cancer.
Mechanism of immune evasion in breast cancer.
Thwarting galectin-induced immunosuppression in breast cancer.
Chemotherapy induces enrichment of CD47(+)/CD73(+)/PDL1(+) immune evasive triple-negative breast cancer cells.
Endoplasmic reticulum stress-induced exosomal miR-27a-3p promotes immune escape in breast cancer via regulating PD-L1 expression in macrophages.
Targeting Notch-Driven Cytokine Secretion: Novel Therapies for Triple Negative Breast Cancer.
microRNAs in the Antitumor Immune Response and in Bone Metastasis of Breast Cancer: From Biological Mechanisms to Therapeutics.
Expression of Treg-associated lncRNAs in breast cancer.

Advancing Immunotherapy in Metastatic Breast Cancer.
Tumor microenvironment of human breast cancer, and feline mammary carcinoma as a potential study model.
Novel combinatorial strategies for boosting the efficacy of immune checkpoint inhibitors in advanced breast cancers.
LCOR mediates interferon-independent tumor immunogenicity and responsiveness to immune-checkpoint blockade in triple-negative breast cancer.
Kaiso (ZBTB33) subcellular partitioning functionally links LC3A/B, the tumor microenvironment, and breast cancer survival.
Differential distribution of immune cells in breast invasive carcinoma vs. breast carcinoma in situ and its significance in interpretation of immune surveillance.
Interleukin-17 Could Promote Breast Cancer Progression at Several Stages of the Disease.
The ectopic expression of IFN regulatory factor 4-binding protein is correlated with the malignant behavior of human breast cancer cells.
Comprehensive molecular profiling of the B7 family of immune-regulatory ligands in breast cancer.
Challenges to the development of antigen-specific breast cancer vaccines.
Triple-negative breast cancer cells rely on kinase-independent functions of CDK8 to evade NK-cell-mediated tumor surveillance.
The STAT3 inhibitor pyrimethamine displays anti-cancer and immune stimulatory effects in murine models of breast cancer.
Revisiting the hallmarks of cancer: A new look at long noncoding RNAs in breast cancer.
Hypoxia: syndicating triple negative breast cancer against various therapeutic regimens.
Intercellular hif1 $\alpha$ reprograms mammary progenitors and myeloid immune evasion to drive high-risk breast lesions.
Misaligned Chromosomes are a Major Source of Chromosomal Instability in Breast Cancer.
Hypoxia-driven ncRNAs in breast cancer.
MicroRNAs delivered by extracellular vesicles: an emerging resistance mechanism for breast cancer.
Lysine-Specific Demethylase 1 (LSD1)-Mediated Epigenetic Modification of Immunogenicity and Immunomodulatory Effects in Breast Cancers.
The BRCA1 Pseudogene Negatively Regulates Antitumor Responses through Inhibition of Innate Immune Defense Mechanisms.
Pharmaceutical targeting Th2-mediated immunity enhances immunotherapy response in breast cancer.
The application of exosomes in the treatment of triple-negative breast cancer.
Breast Cancer Tumor Microenvironment and Molecular Aberrations Hijack Tumoricidal Immunity.
Activation of Host-NLRP3 Inflammasome in Myeloid Cells Dictates Response to Anti-PD-1 Therapy in Metastatic Breast Cancers.
NLRX1 regulates TNF- $\alpha$ -induced mitochondria-lysosomal crosstalk to maintain the invasive and metastatic potential of breast cancer cells.
Therapeutic breast cancer vaccines: a new strategy for early-stage disease.
Tumor infiltrating lymphocytes in early breast cancer.
Identification of a centrosome-related prognostic signature for breast cancer.
Protein Profiling of Breast Carcinomas Reveals Expression of Immune-Suppressive Factors and Signatures Relevant for Patient Outcome.
Association between Cyclooxygenase-2 and Indoleamine 2,3-Dioxygenase Expression in Breast Cancer Patients from Pakistan.
The Multifaceted Effects of Breast Cancer on Tumor-Draining Lymph Nodes.



A Bioinformatics Study of the Involved Mechanisms in Relapse and Drug Resistance of Tamoxifen-Treated Breast Cancer.
The heterogeneous immune microenvironment in breast cancer is affected by hypoxia-related genes.
Analysis of circulating breast cancer cell heterogeneity and interactions with peripheral blood mononuclear cells.
CTLA-4 Expression and Its Clinical Significance in Breast Cancer.
Cancer-associated Fibroblasts Communicate with Breast Tumor Cells Through Extracellular Vesicles in Tumor Development.
Fusobacterium nucleatum: a novel immune modulator in breast cancer?
MicroRNAs Regulating Tumor Immune Response in the Prediction of the Outcome in Patients With Breast Cancer.
Suppression of indoleamine-2,3-dioxygenase 1 expression by promoter hypermethylation in ER-positive breast cancer.
HIF-1 regulates CD47 expression in breast cancer cells to promote evasion of phagocytosis and maintenance of cancer stem cells.
microRNA-155 deficiency impairs dendritic cell function in breast cancer.
The Single-Cell Landscape of Intratumoral Heterogeneity and The Immunosuppressive Microenvironment in Liver and Brain Metastases of Breast Cancer.
High IL-1R8 expression in breast tumors promotes tumor growth and contributes to impaired antitumor immunity.
RAS/MAPK Activation Is Associated with Reduced Tumor-Infiltrating Lymphocytes in Triple-Negative Breast Cancer: Therapeutic Cooperation Between MEK and PD-1/PD-L1 Immune Checkpoint Inhibitors.
Differences in Tumor Immune Microenvironment in Metastatic Sites of Breast Cancer.
Dendritic cell defects in patients with cancer: mechanisms and significance.
Hypoxia inducible factor-1 $\alpha$ (HIF-1 $\alpha$ ) in breast cancer: The crosstalk with oncogenic and onco-suppressor factors in regulation of cancer hallmarks.
Interferon-Stimulated Genes Are Transcriptionally Repressed by PR in Breast Cancer.
Inflammatory breast cancer: High incidence of GCC haplotypes (-1082A/G, -819T/C, and -592A/C) in the interleukin-10 gene promoter correlates with over-expression of interleukin-10 in patients' carcinoma tissues.
Injecting Hope--A Review of Breast Cancer Vaccines.
BTF3 promotes stemness and inhibits Type I Interferon signaling pathway in triple-negative breast cancer.
Tumor cell intrinsic RON signaling suppresses innate immune responses in breast cancer through inhibition of IRAK4 signaling.
A Multi-Source Data Fusion Framework for Revealing the Regulatory Mechanism of Breast Cancer Immune Evasion.
In silico investigation of heparanase-correlated genes in breast cancer subtypes.
The Prognostic Significance of MACC1 Expression in Breast Cancer and Its Relationship to Immune Cells in the Tumor Microenvironment and Patient Survival.
Macrophage Cell Membrane Coating on Piperine-Loaded MIL-100(Fe) Nanoparticles for Breast Cancer Treatment.
Immunogenomic Landscape in Breast Cancer Reveals Immunotherapeutically Relevant Gene Signatures.
Aptamer-Based Strategies to Boost Immunotherapy in TNBC.

Immunobiological characterization of N -nitrosomethylurea-induced rat breast carcinomas: tumoral IL-10 expression as a possible immune escape mechanism.
Lysophosphatidic Acid Receptor Signaling in the Human Breast Cancer Tumor Microenvironment Elicits Receptor-Dependent Effects on Tumor Progression.
c-MYC mediates the crosstalk between breast cancer cells and tumor microenvironment.
Integrating Single-Cell RNA-Seq and Bulk RNA-Seq Data to Explore the Key Role of Fatty Acid Metabolism in Breast Cancer.
In vivo multidimensional CRISPR screens identify Lgals2 as an immunotherapy target in triple-negative breast cancer.
Proteomic analysis of arylamine N-acetyltransferase 1 knockout breast cancer cells: Implications in immune evasion and mitochondrial biogenesis.
COX-2 and PGE2-dependent immunomodulation in breast cancer.
Gemcitabine and doxorubicin in immunostimulatory monophosphoryl lipid A liposomes for treating breast cancer.
FBXW7 in breast cancer: mechanism of action and therapeutic potential.
EGFR Signaling Enhances Aerobic Glycolysis in Triple-Negative Breast Cancer Cells to Promote Tumor Growth and Immune Escape.
Synergistic association of FOXP3+ tumor infiltrating lymphocytes with CCL20 expressions with poor prognosis of primary breast cancer: A retrospective cohort study.
Effect of metabolism on the immune microenvironment of breast cancer.
Silencing of Irf7 pathways in breast cancer cells promotes bone metastasis through immune escape.
Effects of Cancer Stem Cells in Triple-Negative Breast Cancer and Brain Metastasis: Challenges and Solutions.
Interleukin-10 promoter polymorphism is associated with decreased breast cancer risk.
Monocyte chemotactic protein-1 secreted by primary breast tumors stimulates migration of mesenchymal stem cells.
Biological characteristics and genetic heterogeneity between carcinoma-associated fibroblasts and their paired normal fibroblasts in human breast cancer.
The ubiquitin-specific peptidase 22 is a deubiquitinase of CD73 in breast cancer cells.
Cross-Platform in-silico Analyses Exploring Tumor Immune Microenvironment with Prognostic Value in Triple-Negative Breast Cancer.
Analyzing the influence of IL18 in regulation of YAP1 in breast oncogenesis using cBioportal.
Cellular Immune Responses and Immune Escape Mechanisms in Breast Cancer: Determinants of Immunotherapy.
MYC promotes immune-suppression in triple-negative breast cancer via inhibition of interferon signaling.
Resistance to HER2-targeted anti-cancer drugs is associated with immune evasion in cancer cells and their derived extracellular vesicles.
Identification of Immune-Related Therapeutically Relevant Biomarkers in Breast Cancer and Breast Cancer Stem Cells by Transcriptome-Wide Analysis: A Clinical Prospective Study.
Anti-CD47 immunotherapy as a therapeutic strategy for the treatment of breast cancer brain metastasis.
Mucin 1-specific immunotherapy in a mouse model of spontaneous breast cancer.
B7-H3 augments the pro-angiogenic function of tumor-associated macrophages and acts as a novel adjuvant target for triple-negative breast cancer therapy.
Targeting of interleukin (IL)-17A inhibits PDL1 expression in tumor cells and induces anticancer immunity in an estrogen receptor-negative murine model of breast cancer.

Characterization of Apoptosis in a Breast Cancer Cell Line after IL-10 Silencing.
Obesity Influences the Expression of Key Immunomodulators in Normal Human Breast Tissue, Basal-like Breast Cancer Patients, and Cell Lines.
DNA barcoding reveals ongoing immunoediting of clonal cancer populations during metastatic progression and immunotherapy response.
Integrin $\alpha\beta6$ -TGF $\beta$ -SOX4 Pathway Drives Immune Evasion in Triple-Negative Breast Cancer.
Investigation of the effects of overexpression of jumping translocation breakpoint (JTB) protein in MCF7 cells for potential use as a biomarker in breast cancer.
TNBC-derived Gal3BP/Gal3 complex induces immunosuppression through CD45 receptor.
Deciphering Common Traits of Breast and Ovarian Cancer Stem Cells and Possible Therapeutic Approaches.
Decreased Lipid Phosphate Phosphatase 1/3 and Increased Lipid Phosphate Phosphatase 2 Expression in the Human Breast Cancer Tumor Microenvironment Promotes Tumor Progression and Immune System Evasion.
Gap Junctions and Breast Cancer Dormancy.
$\beta$ , $\beta$ -Dimethylacrylshikonin potentiates paclitaxel activity, suppresses immune evasion and triple negative breast cancer progression via STAT3Y705 phosphorylation inhibition based on network pharmacology and transcriptomics analysis.
Comprehensive analysis of the association between tumor glycolysis and immune/inflammation function in breast cancer.
Correlation between the Warburg effect and progression of triple-negative breast cancer.
Doxorubicin downregulates cell surface B7-H1 expression and upregulates its nuclear expression in breast cancer cells: role of B7-H1 as an anti-apoptotic molecule.
Dormant Tumor Cell Vaccination: A Mathematical Model of Immunological Dormancy in Triple-Negative Breast Cancer.
Multiple functions of sushi domain containing 2 (SUSD2) in breast tumorigenesis.
Sequentially-targeted biomimetic nano drug system for triple-negative breast cancer ablation and lung metastasis inhibition.
Tradeoff between metabolic i-proteasome addiction and immune evasion in triple-negative breast cancer.
Tumor-derived Jagged1 promotes cancer progression through immune evasion.
Peripheral blood cells inform on the presence of breast cancer: a population-based case-control study.
A novel immune subtype classification of ER-positive, PR-negative and HER2-negative breast cancer based on the genomic and transcriptomic landscape.
Changes in peripheral immune cells after intraoperative radiation therapy in low-risk breast cancer.
Mesenchymal stem/stromal cells in breast cancer development and management.
Therapy of advanced established murine breast cancer with a recombinant adenoviral ErbB-2/neu vaccine.
The Generation of Dual-Targeting Fusion Protein PD-L1/CD47 for the Inhibition of Triple-Negative Breast Cancer.
LXR-inverse agonism stimulates immune-mediated tumor destruction by enhancing CD8 T-cell activity in triple negative breast cancer.
Unraveling Triple-Negative Breast Cancer Tumor Microenvironment Heterogeneity: Towards an Optimized Treatment Approach.
MYC suppresses STING-dependent innate immunity by transcriptionally upregulating DNMT1 in triple-negative breast cancer.

CD46 Expression is an unfavorable prognostic factor in breast cancer cases.
Triple-negative and HER2-overexpressing breast cancer cell sialylation impacts tumor microenvironment T-lymphocyte subset recruitment: a possible mechanism of tumor escape.
Extracellular Vesicles in Breast Cancer: From Biology and Function to Clinical Diagnosis and Therapeutic Management.
The mutation and low expression of ARID1A are predictive of a poor prognosis and high immune infiltration in triple-negative breast cancer.
STAT3 in Breast Cancer Onset and Progression: A Matter of Time and Context.
Breast Cancer Immunology and Immunotherapy: Current Status and Future Perspectives.
Metabolic crosstalk in the breast cancer microenvironment.
Identification of molecular subtypes and a six-gene risk model related to cuproptosis for triple negative breast cancer.
Histone deacetylase 6 regulates the immunosuppressive properties of cancer-associated fibroblasts in breast cancer through the STAT3-COX2-dependent pathway.
The impact of lipid metabolism on breast cancer: a review about its role in tumorigenesis and immune escape.
Cellular and molecular insights into the roles of visfatin in breast cancer cells plasticity programs.
The synthetic histone-binding regulator protein PcTF activates interferon genes in breast cancer cells.
B7-H3 on circulating epithelial tumor cells correlates with the proliferation marker, Ki-67, and may be associated with the aggressiveness of tumors in breast cancer patients.
Spatial immunophenotypes predict response to anti-PD1 treatment and capture distinct paths of T cell evasion in triple negative breast cancer.
Glycosyltransferases as marker genes for the quantitative polymerase chain reaction-based detection of circulating tumour cells from blood samples of patients with breast cancer undergoing adjuvant therapy.
ABCG2 is a potential marker of tumor-initiating cells in breast cancer.
Progesterone receptor promotes degradation of STAT2 to inhibit the interferon response in breast cancer.
Increased Expression of INHBA Is Correlated With Poor Prognosis and High Immune Infiltrating Level in Breast Cancer.
The Multi-Faced Role of PAPP-A in Post-Partum Breast Cancer: IGF-Signaling is Only the Beginning.
Hybrid-cell membrane-coated nanocomplex-loaded chikusetsusaponin IVa methyl ester for a combinational therapy against breast cancer assisted by Ce6.
Targeting DCLK1 attenuates tumor stemness and evokes antitumor immunity in triple-negative breast cancer by inhibiting IL-6/STAT3 signaling.
Hypoxia induces HIF1 $\alpha$ -dependent epigenetic vulnerability in triple negative breast cancer to confer immune effector dysfunction and resistance to anti-PD-1 immunotherapy.
Multi-Omics Profiling Reveals Distinct Microenvironment Characterization and Suggests Immune Escape Mechanisms of Triple-Negative Breast Cancer.
Clinical significance of CD73 in triple-negative breast cancer: multiplex analysis of a phase III clinical trial.
Tumor-infiltrating lymphocytes are associated with $\beta$ -catenin overexpression in breast cancer.
Expression and Clinical Significance of Herpes Virus Entry Mediator (HVEM) in Breast Cancer.
Drug-repositioning screening identified piperlongumine as a direct STAT3 inhibitor with potent activity against breast cancer.
Strong Correlation of Indoleamine 2,3-Dioxygenase 1 Expression with Basal-Like Phenotype and Increased Lymphocytic Infiltration in Triple-Negative Breast Cancer.

Combined Radionuclide Therapy and Immunotherapy for Treatment of Triple Negative Breast Cancer.
Metastatic breast cancers have reduced immune cell recruitment but harbor increased macrophages relative to their matched primary tumors.
Treatment of Metastatic or High-Risk Solid Cancer Patients by Targeting the Immune System and/or Tumor Burden: Six Cases Reports.
ER-localized JmjC domain-containing protein JMJD8 targets STING to promote immune evasion and tumor growth in breast cancer.
2-Methoxy-1,4-Naphthoquinone (MNQ) Inhibits Glucose Uptake and Lactate Production in Triple-Negative Breast Cancer Cells.
Integrative analysis of copy number and gene expression data identifies potential oncogenic drivers that promote mammary tumor recurrence.
Type I Interferon Regulates a Coordinated Gene Network to Enhance Cytotoxic T Cell-Mediated Tumor Killing.
Expression of death decoy receptor-3 (DcR3) in human breast cancer and its functional effects on breast cancer cells in vitro.
Epstein-Barr virus as a promoter of tumorigenesis in the tumor microenvironment of breast cancer (Review).
The Tim-3-Galectin-9 Pathway and Its Regulatory Mechanisms in Human Breast Cancer.
Pattern Analysis of Serum Galectins-1, -3, and -9 in Breast Cancer.
Targeting galectin-1 overcomes breast cancer-associated immunosuppression and prevents metastatic disease.
Stromal cell diversity associated with immune evasion in human triple-negative breast cancer.
Extracellular vesicles as a novel approach for breast cancer therapeutics.
Bioinformatics identification of CCL8/21 as potential prognostic biomarkers in breast cancer microenvironment.
Targeting mTOR to overcome resistance to hormone and CDK4/6 inhibitors in ER-positive breast cancer models.
The Combination of Immune Checkpoint Blockade with Tumor Vessel Normalization as a Promising Therapeutic Strategy for Breast Cancer: An Overview of Preclinical and Clinical Studies.
OLR1 is a prognostic factor and correlated with immune infiltration in breast cancer.
MUC1-C integrates activation of the IFN- $\gamma$ pathway with suppression of the tumor immune microenvironment in triple-negative breast cancer.
The Immunosuppressive Microenvironment in BRCA1-IRIS-Overexpressing TNBC Tumors Is Induced by Bidirectional Interaction with Tumor-Associated Macrophages.
Breast cancer cells survive chemotherapy by activating targetable immune-modulatory programs characterized by PD-L1 or CD80.
Effects of Jaeumkanghwa-tang on tamoxifen responsiveness in preclinical ER+ breast cancer model.
HER2 + breast cancers evade anti-HER2 therapy via a switch in driver pathway.
Tumor-B-cell interactions promote isotype switching to an immunosuppressive IgG4 antibody response through upregulation of IL-10 in triple negative breast cancers.
Free fatty acid release from human breast cancer tissue inhibits cytotoxic T-lymphocyte-mediated killing.
Corrigendum: Immunogenomic landscape in breast cancer reveals immunotherapeutically relevant gene signatures.
Breaking through to the Other Side: Microenvironment Contributions to DCIS Initiation and Progression.

Triptolide inhibits interferon-gamma-induced programmed death-1-ligand 1 surface expression in breast cancer cells.
Epigenetic oncogenesis, biomarkers and emerging chemotherapeutics for breast cancer.
B7-H3 expression in ductal and lobular breast cancer and its association with IL-10.
Increased B Regulatory Phenotype in Non-Metastatic Lymph Nodes of Node-Positive Breast Cancer Patients.
The role and application of small extracellular vesicles in breast cancer.
Immune signatures of murine and human cancers reveal unique mechanisms of tumor escape and new targets for cancer immunotherapy.
Cell- and subcellular organelle-targeting nanoparticle-mediated breast cancer therapy.
Tumor escape and progression of HER-2/neu negative breast cancer under immune pressure.
The Role of Hypoxia-Inducible Factor Isoforms in Breast Cancer and Perspectives on Their Inhibition in Therapy.
The modulatory role of dendritic cell-T cell cross-talk in breast cancer: Challenges and prospects.
Effect of antiprogestins and tamoxifen on growth inhibition of MCF-7 human breast cancer cells in nude mice.
Apigenin inhibits TNF $\alpha$ /IL-1 $\alpha$ -induced CCL2 release through IKK $\epsilon$ -signaling in MDA-MB-231 human breast cancer cells.
Tumor promoting effect of circ_002172 associates with induced immune escape in breast cancer via the miR-296-5p/CXCL12 axis.
Deciphering antihormone-induced compensatory mechanisms in breast cancer and their therapeutic implications.
Addition of Cancer Stem Cells to MUC1-C in Triple-Negative Breast Cancer Progression.
Relationship between circulating tumor cells and peripheral T-cells in patients with primary breast cancer.
Tumor-derived CCL-2 and CXCL-8 as possible prognostic markers of breast cancer: correlation with estrogen and progesterone receptor phenotyping.
Erythrocyte-cancer hybrid membrane-coated reduction-sensitive nanoparticles for enhancing chemotherapy efficacy in breast cancer.
Glycosylation Modulates Plasma Membrane Trafficking of CD24 in Breast Cancer Cells.
The C-X-C Motif Chemokine Ligand 1 Sustains Breast Cancer Stem Cell Self-Renewal and Promotes Tumor Progression and Immune Escape Programs.
Surface Expression of TGF $\beta$ Docking Receptor GARP Promotes Oncogenesis and Immune Tolerance in Breast Cancer.
Heat shock protein 27 differentiates tolerogenic macrophages that may support human breast cancer progression.
Are aggressive epithelial cancers 'a disease' of Eutherian mammals?
MicroRNAs Regulating Tumor and Immune Cell Interactions in the Prediction of Relapse in Early Stage Breast Cancer.
Toll-like receptors gene polymorphisms may confer increased susceptibility to breast cancer development.
MACC1 as a Potential Target for the Treatment and Prevention of Breast Cancer.
HER3 PET Imaging Identifies Dynamic Changes in HER3 in Response to HER2 Inhibition with Lapatinib.
Cancer-produced metabolites of 5-lipoxygenase induce tumor-evoked regulatory B cells via peroxisome proliferator-activated receptor $\alpha$ .
P-Cadherin Linking Breast Cancer Stem Cells and Invasion: A Promising Marker to Identify an "Intermediate/Metastable" EMT State.

Expression of B7-H1 in breast cancer patients is strongly associated with high proliferative Ki-67-expressing tumor cells.
Spontaneous apoptosis of blood dendritic cells in patients with breast cancer.
Sialylation of CD55 by ST3GAL1 Facilitates Immune Evasion in Cancer.
In Vivo Imaging to Measure Spontaneous Lung Metastasis of Orthotopically-injected Breast Tumor Cells.
RNF31 represses cell progression and immune evasion via YAP/PD-L1 suppression in triple negative breast Cancer.
Single-Cell Analysis Reveals Fibroblast Clusters Linked to Immunotherapy Resistance in Cancer.
Blocking soluble TNF $\alpha$ sensitizes HER2-positive breast cancer to trastuzumab through MUC4 downregulation and subverts immunosuppression.
Enhanced glutamine uptake influences composition of immune cell infiltrates in breast cancer.
Inflammasomes in breast cancer: the ignition spark of progression and resistance?
Breast cancer detection using targeted plasma metabolomics.
Integrated multiomics analyses unveil the implication of a costimulatory molecule score on tumor aggressiveness and immune evasion in breast cancer: A large-scale study through over 8,000 patients.
miR-18a Mediates Immune Evasion in ER-Positive Breast Cancer through Wnt Signaling.
Cancer Cell-Erythrocyte Hybrid Membrane Coated Gold Nanocages for Near Infrared Light-Activated Photothermal/Radio/Chemotherapy of Breast Cancer.
Breast tumor microenvironment structures are associated with genomic features and clinical outcome.
A map of human breast cancer: new players in stromal-immune crosstalk.
Selectively hampered activation of lymph node-resident dendritic cells precedes profound T cell suppression and metastatic spread in the breast cancer sentinel lymph node.
Overrepresentation of the EBAG9 gene at 8q23 associated with early-stage breast cancers.
Confined migration promotes cancer metastasis through resistance to anoikis and increased invasiveness.
Anti-KL-6/MUC1 monoclonal antibody reverses resistance to trastuzumab-mediated antibody-dependent cell-mediated cytotoxicity by capping MUC1.
FGFR blockade boosts T cell infiltration into triple-negative breast cancer by regulating cancer-associated fibroblasts.
Epigenetic Switch-Induced Viral Mimicry Evasion in Chemotherapy-Resistant Breast Cancer.
Evaluation of glucocorticoid-induced TNF receptor (GITR) expression in breast cancer and across multiple tumor types.
A highly polymorphic CA repeat marker at the EBAG9/RCA51 locus on 8q23 that detected frequent multiplication in breast cancer.
Immunotherapy for inflammatory breast cancer: current evidences and future perspectives.
Tumor-derived microparticles promote the progression of triple-negative breast cancer via PD-L1-associated immune suppression.
Does hCG or hCG $\beta$ play a role in cancer cell biology?
TLR4 and pSTAT3 Expression on Circulating Tumor Cells (CTCs) and Immune Cells in the Peripheral Blood of Breast Cancer Patients: Prognostic Implications.
Placental immune editing switch (PIES): learning about immunomodulatory pathways from a unique case report.
Naturally Equipped Urinary Exosomes Coated Poly (2-ethyl-2-oxazoline)-Poly (D, L-lactide) Nanocarriers for the Pre-Clinical Translation of Breast Cancer.
Autophagy-deficient breast cancer shows early tumor recurrence and escape from dormancy.

Understanding the role of the kynurenine pathway in human breast cancer immunobiology.
Tumor-Associated Fibroblasts Promote HER2-Targeted Therapy Resistance through FGFR2 Activation.
Carbonic Anhydrase-IX Guided Albumin Nanoparticles for Hypoxia-mediated Triple-Negative Breast Cancer Cell Killing and Imaging of Patient-derived Tumor.
Interaction Between MUC1 and STAT1 Drives IFITM1 Overexpression in Aromatase Inhibitor-Resistant Breast Cancer Cells and Mediates Estrogen-Induced Apoptosis.
Specific N-cadherin-dependent pathways drive human breast cancer dormancy in bone marrow.
The PI3K-Akt-mTOR Pathway and New Tools to Prevent Acquired Hormone Resistance in Breast Cancer.
Interferon regulatory factor 1 (IRF-1) and IRF-2 expression in breast cancer tissue microarrays.
"Lazarus Response" to Olaparib in a Virtually Chemonaive Breast Cancer Patient Carrying Gross BRCA2 Gene Deletion.
Proteomics analysis of E-cadherin knockdown in epithelial breast cancer cells.
Improved Autophagic Flux in Escapers from Doxorubicin-Induced Senescence/Polyplody of Breast Cancer Cells.
Anti-CD20 antibody promotes cancer escape via enrichment of tumor-evoked regulatory B cells expressing low levels of CD20 and CD137L.
Progressive loss of anti-HER2 CD4(+) T-helper type 1 response in breast tumorigenesis and the potential for immune restoration.
Novel Mechanisms of Tumor Promotion by the Insulin Receptor Isoform A in Triple-Negative Breast Cancer Cells.
MUC1 and MUC4: switching the emphasis from large to small.
The immunogram of inflammatory breast cancer.
Circular RNA hsa_circ_0067842 facilitates tumor metastasis and immune escape in breast cancer through HuR/CMTM6/PD-L1 axis.
HNRNPC promotes collagen fiber alignment and immune evasion in breast cancer via activation of the VIRMA-mediated TFAP2A/DDR1 axis.
Development of a High-Affinity Antibody against the Tumor-Specific and Hyperactive 611-p95HER2 Isoform.
Principles of immunotherapy.
The breast pre-cancer atlas illustrates the molecular and micro-environmental diversity of ductal carcinoma in situ.
The efficacy of indoximod upon stimulation with pro-inflammatory cytokines in triple-negative breast cancer cells.
A simple and rapid protocol to non-enzymatically dissociate fresh human tissues for the analysis of infiltrating lymphocytes.
Tumor-secreted products repress B-cell lymphopoiesis in a murine model of breast cancer.
Indoleamine 2,3-dioxygenase activity and L-tryptophan transport in human breast cancer cells.
Oncogenic lncRNA downregulates cancer cell antigen presentation and intrinsic tumor suppression.
MicroRNA signature in the chemoprevention of functionally-enriched stem and progenitor pools (FESPP) by Active Hexose Correlated Compound (AHCC).
Hypoxia-Induced Intracellular and Extracellular Heat Shock Protein gp96 Increases Paclitaxel-Resistance and Facilitates Immune Evasion in Breast Cancer.
Pattern of hormone receptor status of secondary contralateral breast cancers in patients receiving adjuvant tamoxifen.
Increased Circulating Levels of Galectin Proteins in Patients with Breast, Colon, and Lung Cancer.
Chemokines in tumor progression and metastasis.



Cognitive-behavioral stress management reverses anxiety-related leukocyte transcriptional dynamics.
Generation and characterisation of two D2A1 mammary cancer sublines to model spontaneous and experimental metastasis in a syngeneic BALB/c host.
MUC1-C Dictates PBRM1-Mediated Chronic Induction of Interferon Signaling, DNA Damage Resistance, and Immunosuppression in Triple-Negative Breast Cancer.
Metabolomic and Mitochondrial Fingerprinting of the Epithelial-to-Mesenchymal Transition (EMT) in Non-Tumorigenic and Tumorigenic Human Breast Cells.
PI(3) kinase is associated with a mechanism of immunoresistance in breast and prostate cancer.
Biological Role and Aberrant Overexpression of Syntenin-1 in Cancer: Potential Role as a Biomarker and Therapeutic Target.
EBAG9/RCAS1 in human breast carcinoma: a possible factor in endocrine-immune interactions.
piTracer - Automatic reconstruction of molecular cascades for the identification of synergistic drug targets.
The complement regulatory proteins CD46 and CD59, but not CD55, are highly expressed by glandular epithelium of human breast and colorectal tumour tissues.
Transcriptomic and Functional Evidence for Differential Effects of MCF-7 Breast Cancer Cell-Secretome on Vascular and Lymphatic Endothelial Cell Growth.
Deciphering the role of interferon alpha signaling and microenvironment crosstalk in inflammatory breast cancer.
The dynamics of the inflammatory response during BBN-induced bladder carcinogenesis in mice.
Leptin decreases BC cell susceptibility to NK lysis via PGC1A pathway.
High expression levels of egfl7 correlate with low endothelial cell activation in peritumoral vessels of human breast cancer.
Tanapoxvirus: From discovery towards oncolytic immunovirotherapy.
Retraction statement: Ding, J., Wang, X., Gao, J. and Song, T. (2021), Silencing of cystatin SN abrogates cancer progression and stem cell properties in papillary thyroid carcinoma. FEBS Open Bio, 11: 2186-2197.
Plasmacytoid dendritic cells (PDC) are the major DC subset innately producing cytokines in human lymph nodes.
Epithelial-to-mesenchymal transition (EMT) induced by inflammatory priming elicits mesenchymal stromal cell-like immune-modulatory properties in cancer cells.
Circulating Tumor Cells Develop Resistance to TRAIL-Induced Apoptosis Through Autophagic Removal of Death Receptor 5: Evidence from an In Vitro Model.
Estrogen matters in metastasis.
EBAG9/RCAS1 expression and its prognostic significance in prostatic cancer.
Thymic stromal lymphopoietin is a key mediator of breast cancer progression.
Integrating somatic CNV and gene expression in breast cancers from women with PTEN hamartoma tumor syndrome.
Upregulation of PD-L1 expression by resveratrol and piceatannol in breast and colorectal cancer cells occurs via HDAC3/p300-mediated NF-κB signaling.
The microRNA target site landscape is a novel molecular feature associating alternative polyadenylation with immune evasion activity in breast cancer.
Relationship Between Hematogenous Tumor Cell Dissemination and Cellular Immunity in DCIS Patients.
SUSD2 Proteolytic Cleavage Requires the GDPH Sequence and Inter-Fragment Disulfide Bonds for Surface Presentation of Galectin-1 on Breast Cancer Cells.

A targeted isotope dilution mass spectrometry assay for osteopontin quantification in plasma of metastatic breast cancer patients.
If You Cannot Win Them, Join Them: Understanding New Ways to Target STAT3 by Small Molecules.
Glycan elongation beyond the mucin associated Tn antigen protects tumor cells from immune-mediated killing.
Differential kynurenine pathway metabolism in highly metastatic aggressive breast cancer subtypes: beyond IDO1-induced immunosuppression.
Invasive lobular and ductal breast carcinoma differ in immune response, protein translation efficiency and metabolism.
Promoter analysis and chromosomal mapping of human EBAG9 gene.
Altered Glycosylation in Progression and Management of Bladder Cancer.
Genetic and epigenetic changes in mammary epithelial cells identify a subpopulation of cells involved in early carcinogenesis.
Immunosuppressive Traits of the Hybrid Epithelial/Mesenchymal Phenotype.
Learning to detect lymphocytes in immunohistochemistry with deep learning.
Rapalog-Mediated Repression of Tribbles Pseudokinase 3 Regulates Pre-mRNA Splicing.

Table A2-34, Cluster 33

Cluster 33 focuses on MHC class I, especially the viral interference with the MHC Class I antigen processing and presentation pathway and the subsequent viral immune system evasion (297)
A Flow Cytometry-Based Approach to Unravel Viral Interference with the MHC Class I Antigen Processing and Presentation Pathway.
The MHC class I antigen presentation pathway: strategies for viral immune evasion.
Involvement of autophagy in MHC class I antigen presentation.
A comparison of viral immune escape strategies targeting the MHC class I assembly pathway.
Control of MHC class I traffic from the endoplasmic reticulum by cellular chaperones and viral anti-chaperones.
Inhibition of MHC class I is a virulence factor in herpes simplex virus infection of mice.
MHC class I antigen presentation: learning from viral evasion strategies.
Viral immune evasion: Lessons in MHC class I antigen presentation.
Equine herpesvirus type 4 UL56 and UL49.5 proteins downregulate cell surface major histocompatibility complex class I expression independently of each other.
A peptide filtering relation quantifies MHC class I peptide optimization.
Cytokines restore MHC class I complex formation and control antigen presentation in human cytomegalovirus-infected cells.
Herpesviruses and immunity: the art of evasion.
New insights into the structure of the MHC class I peptide-loading complex and mechanisms of TAP inhibition by viral immune evasion proteins.
Molecular mechanisms of viral immune evasion proteins to inhibit MHC class I antigen processing and presentation.
In vivo Downregulation of MHC Class I Molecules by HCMV Occurs During All Phases of Viral Replication but Is Not Always Complete.
Processing and MHC class I presentation of human cytomegalovirus pp65-derived peptides persist despite gpUS2-11-mediated immune evasion.

Human cytomegalovirus-encoded immune modulators partner to downregulate major histocompatibility complex class I molecules.
Varicella-zoster virus retains major histocompatibility complex class I proteins in the Golgi compartment of infected cells.
Major histocompatibility complex class I allele-specific cooperative and competitive interactions between immune evasion proteins of cytomegalovirus.
MHC class I-subversive gene functions of cytomegalovirus and their regulation by interferons-an intricate balance.
The influence of TAP1 and TAP2 gene polymorphisms on TAP function and its inhibition by viral immune evasion proteins.
The macromolecular peptide-loading complex in MHC class I-dependent antigen presentation.
Cutting edge: adenovirus E19 has two mechanisms for affecting class I MHC expression.
An MHC class I immune evasion gene of Marek's disease virus.
The class I MHC homologue of human cytomegalovirus inhibits attack by natural killer cells.
Kaposi's sarcoma-associated herpesvirus encodes two proteins that block cell surface display of MHC class I chains by enhancing their endocytosis.
Immune evasion mechanisms of varicella-zoster virus.
Dynamics of the major histocompatibility complex class I processing and presentation pathway in the course of malaria parasite development in human hepatocytes: implications for vaccine development.
Herpes B virus, macacine herpesvirus 1, breaks simplex virus tradition via major histocompatibility complex class I expression in cells from human and macaque hosts.
The ABCs of immunology: structure and function of TAP, the transporter associated with antigen processing.
Human cytomegalovirus disrupts the major histocompatibility complex class I peptide-loading complex and inhibits tapasin gene transcription.
Epstein Barr virus-encoded EBNA1 interference with MHC class I antigen presentation reveals a close correlation between mRNA translation initiation and antigen presentation.
The MHC class I homolog of human cytomegalovirus is resistant to down-regulation mediated by the unique short region protein (US)2, US3, US6, and US11 gene products.
KSHV-K5 inhibits phosphorylation of the major histocompatibility complex class I cytoplasmic tail.
Adaptor protein 1 promotes cross-presentation through the same tyrosine signal in major histocompatibility complex class I as that targeted by HIV-1.
Antigen translocation machineries in adaptive immunity and viral immune evasion.
Identifying the ERAD ubiquitin E3 ligases for viral and cellular targeting of MHC class I.
Human cytomegalovirus inhibits tapasin-dependent peptide loading and optimization of the MHC class I peptide cargo for immune evasion.
Downregulation of MHC Class I Expression by Influenza A and B Viruses.
Two mechanistically distinct immune evasion proteins of cowpox virus combine to avoid antiviral CD8 T cells.
Characterization of a murine cytomegalovirus class I major histocompatibility complex (MHC) homolog: comparison to MHC molecules and to the human cytomegalovirus MHC homolog.
Restriction of self-antigen presentation to cytolytic T lymphocytes by mouse peptide pumps.
MHC class I up-regulation by flaviviruses: Immune interaction with unknown advantage to host or pathogen.
Requirements for the selective degradation of endoplasmic reticulum-resident major histocompatibility complex class I proteins by the viral immune evasion molecule mK3.
Peptide transport in antigen presentation.

Downregulation of class I major histocompatibility complex surface expression by varicella-zoster virus involves open reading frame 66 protein kinase-dependent and -independent mechanisms.
Cytomegalovirus gp40/m152 Uses TMED10 as ER Anchor to Retain MHC Class I.
Identification of key amino acid residues that determine the ability of high risk HPV16-E7 to dysregulate major histocompatibility complex class I expression.
Giving CD4+ T cells the slip: viral interference with MHC class II-restricted antigen processing and presentation.
Distinct Roles of Extracellular Domains in the Epstein-Barr Virus-Encoded BILF1 Receptor for Signaling and Major Histocompatibility Complex Class I Downregulation.
HIV-1 Nef Interacts with LMP7 To Attenuate Immunoproteasome Formation and Major Histocompatibility Complex Class I Antigen Presentation.
Effects of messenger RNA structure and other translational control mechanisms on major histocompatibility complex-I mediated antigen presentation.
Infection with foot-and-mouth disease virus results in a rapid reduction of MHC class I surface expression.
A Novel MHC-I Surface Targeted for Binding by the MCMV m06 Immuno-evasin Revealed by Solution NMR.
Inhibition of natural killer cells by a cytomegalovirus MHC class I homologue in vivo.
Recent advances in viral evasion of the MHC Class I processing pathway.
Human pathogen subversion of antigen presentation.
Viral immune evasion molecules attack the ER peptide-loading complex and exploit ER-associated degradation pathways.
Inhibition of the MHC class II antigen presentation pathway by human cytomegalovirus.
Downregulation of Cell Surface Major Histocompatibility Complex Class I Expression Is Mediated by the Left-End Transcription Unit of Fowl Adenovirus 9.
Down-regulation of MHC class I expression by equine herpesvirus-1.
Murine cytomegalovirus degrades MHC class II to colonize the salivary glands.
Structural basis of mouse cytomegalovirus m152/gp40 interaction with RAE1 $\gamma$ reveals a paradigm for MHC/MHC interaction in immune evasion.
Varicella-zoster virus immune evasion.
Inhibition of MHC class I-restricted antigen presentation by gamma 2-herpesviruses.
Autophagy mediates transporter associated with antigen processing-independent presentation of viral epitopes through MHC class I pathway.
Trypanosoma cruzi infection down-modulates the immunoproteasome biosynthesis and the MHC class I cell surface expression in HeLa cells.
Bovine herpesvirus-1 infection affects the peptide transport activity in bovine cells.
Function of the cargo sorting dileucine motif in a cytomegalovirus immune evasion protein.
Human Cytomegalovirus Decreases Major Histocompatibility Complex Class II by Regulating Class II Transactivator Transcript Levels in a Myeloid Cell Line.
Structural determinants of peptide-dependent TAP1-TAP2 transit passage targeted by viral proteins and altered by cancer-associated mutations.
A viral inhibitor of peptide transporters for antigen presentation.
Modulation of MHC and MHC-Like Molecules by Varicella Zoster Virus.
Toxoplasma gondii glycosylphosphatidylinositols up-regulate major histocompatibility complex (MHC) molecule expression on primary murine macrophages.
Immune evasion by a novel family of viral PHD/LAP-finger proteins of gamma-2 herpesviruses and poxviruses.

Antigen presentation subverted: Structure of the human cytomegalovirus protein US2 bound to the class I molecule HLA-A2.
A mechanism of viral immune evasion revealed by cryo-EM analysis of the TAP transporter.
HIV-1 Nef disrupts antigen presentation early in the secretory pathway.
Highly pathogenic simian immunodeficiency virus mne variants that emerge during the course of infection evolve enhanced infectivity and the ability to downregulate CD4 but not class I major histocompatibility complex antigens.
HIV-1 Nef protein binds to the cellular protein PACS-1 to downregulate class I major histocompatibility complexes.
Viral inhibition of the transporter associated with antigen processing (TAP): a striking example of functional convergent evolution.
The ER-luminal domain of the HCMV glycoprotein US6 inhibits peptide translocation by TAP.
Down-regulation of MHC class I by the Marek's disease virus (MDV) UL49.5 gene product mildly affects virulence in a haplotype-specific fashion.
Inhibition of HLA-DR assembly, transport, and loading by human cytomegalovirus glycoprotein US3: a novel mechanism for evading major histocompatibility complex class II antigen presentation.
Cowpox virus inhibits the transporter associated with antigen processing to evade T cell recognition.
Autophagy in innate and adaptive immunity against intracellular pathogens.
Equine Herpesvirus 1 Multiply Inserted Transmembrane Protein pUL43 Cooperates with pUL56 in Downregulation of Cell Surface Major Histocompatibility Complex Class I.
The murine cytomegalovirus immune evasion protein m4/gp34 forms biochemically distinct complexes with class I MHC at the cell surface and in a pre-Golgi compartment.
The murine cytomegalovirus immunoevasin gp40 binds MHC class I molecules to retain them in the early secretory pathway.
Functional analysis of the human cytomegalovirus immune evasion protein, pUS3(22kDa).
Downregulation of major histocompatibility complex class I in bovine papillomas.
Suppression of CD8+ T-cell recognition in the immediate-early phase of human cytomegalovirus infection.
Major histocompatibility complex class I downregulation induced by equine herpesvirus type 1 pUL56 is through dynamin-dependent endocytosis.
Evidence that a mechanism for efficient flavivirus budding upregulates MHC class I.
The C-terminal amino acid of the MHC-I heavy chain is critical for binding to Derlin-1 in human cytomegalovirus US11-induced MHC-I degradation.
A survival game of hide and seek: cytomegaloviruses and MHC class I antigen presentation pathways.
Herpesviruses placating the unwilling host: manipulation of the MHC class II antigen presentation pathway.
The p36 isoform of murine cytomegalovirus m152 protein suffices for mediating innate and adaptive immune evasion.
Mouse Norovirus Infection Reduces the Surface Expression of Major Histocompatibility Complex Class I Proteins and Inhibits CD8(+) T Cell Recognition and Activation.
T cell immunity in HSV-1- and VZV-infected neural ganglia.
How the virus outsmarts the host: function and structure of cytomegalovirus MHC-I-like molecules in the evasion of natural killer cell surveillance.
Coxsackievirus B3 proteins directionally complement each other to downregulate surface major histocompatibility complex class I.
The putative natural killer decoy early gene m04 (gp34) of murine cytomegalovirus encodes an antigenic peptide recognized by protective antiviral CD8 T cells.

The Epstein-Barr virus-encoded BILF1 protein modulates immune recognition of endogenously processed antigen by targeting major histocompatibility complex class I molecules trafficking on both the exocytic and endocytic pathways.
Immune evasion proteins of murine cytomegalovirus preferentially affect cell surface display of recently generated peptide presentation complexes.
The TAP translocation machinery in adaptive immunity and viral escape mechanisms.
Degradation of transcription factor RFX5 during the inhibition of both constitutive and interferon gamma-inducible major histocompatibility complex class I expression in chlamydia-infected cells.
Downregulation of major histocompatibility complex class I by human ubiquitin ligases related to viral immune evasion proteins.
The trafficking and regulation of membrane receptors by the RING-CH ubiquitin E3 ligases.
Chlamydia trachomatis-infected epithelial cells and fibroblasts retain the ability to express surface-presented major histocompatibility complex class I molecules.
MHC class I cross-presentation by dendritic cells counteracts viral immune evasion.
Downregulation of major histocompatibility complex class I molecules by Kaposi's sarcoma-associated herpesvirus K3 and K5 proteins.
Inhibition of mouse TAP by immune evasion molecules encoded by non-murine herpesviruses.
Chlamydia trachomatis Infection Impairs MHC-I Intracellular Trafficking and Antigen Cross-Presentation by Dendritic Cells.
Cowpox virus evades CTL recognition and inhibits the intracellular transport of MHC class I molecules.
Viral degradation of the MHC class I peptide loading complex.
In vivo virulence of MHC-adapted AIDS virus serially-passaged through MHC-mismatched hosts.
Cyclophilin C Participates in the US2-Mediated Degradation of Major Histocompatibility Complex Class I Molecules.
Viral evasion of T cell immunity: ancient mechanisms offering new applications.
Model for the interaction of gammaherpesvirus 68 RING-CH finger protein mK3 with major histocompatibility complex class I and the peptide-loading complex.
Legionella pneumophila down-regulates MHC class I expression of human monocytic host cells and thereby inhibits T cell activation.
Cooperative binding of the class I major histocompatibility complex cytoplasmic domain and human immunodeficiency virus type 1 Nef to the endosomal AP-1 complex via its mu subunit.
Epitope-specific in vivo protection against cytomegalovirus disease by CD8 T cells in the murine model of preemptive immunotherapy.
Moving the Cellular Peptidome by Transporters.
Investigation of endogenous antigen processing by delivery of an intact protein into cells.
The murine gamma-herpesvirus-68 MK3 protein causes TAP degradation independent of MHC class I heavy chain degradation.
Varicelloviruses avoid T cell recognition by UL49.5-mediated inactivation of the transporter associated with antigen processing.
Marek's disease virus up-regulates major histocompatibility complex class II cell surface expression in infected cells.
Salmonella modulates B cell biology to evade CD8(+) T cell-mediated immune responses.
Varicellovirus UL 49.5 proteins differentially affect the function of the transporter associated with antigen processing, TAP.
HIV-1 Nef sequesters MHC-I intracellularly by targeting early stages of endocytosis and recycling.
Substrate-specific presentation of MHC class I-restricted antigens via autophagy pathway.

Herpes viral proteins blocking the transporter associated with antigen processing TAP--from genes to function and structure.
Immune evasion during varicella zoster virus infection of keratinocytes.
Applications of major histocompatibility complex class I molecules expressed as single chains.
Immune evasion proteins of human cytomegalovirus do not prevent a diverse CD8+ cytotoxic T-cell response in natural infection.
MHC class I immune evasion in MCMV infection.
Forced interaction of cell surface proteins with Derlin-1 in the endoplasmic reticulum is sufficient to induce their dislocation into the cytosol for degradation.
The PHD/LAP-domain protein M153R of myxomavirus is a ubiquitin ligase that induces the rapid internalization and lysosomal destruction of CD4.
A viral ER-resident glycoprotein inactivates the MHC-encoded peptide transporter.
Viral interference with antigen presentation does not alter acute or chronic CD8 T cell immunodominance in murine cytomegalovirus infection.
Heterologous expression of adenovirus E3-gp19K in an E1a-deleted adenovirus vector inhibits MHC I expression in vitro, but does not prolong transgene expression in vivo.
IK induced by coxsackievirus B3 infection transiently downregulates expression of MHC class II through increasing cAMP.
The downregulation of CD4 and MHC-I by primate lentiviruses: a paradigm for the modulation of cell surface receptors.
Molluscum contagiosum virus MC80 sabotages MHC-I antigen presentation by targeting tapasin for ER-associated degradation.
Processing and presentation of murine cytomegalovirus pORFm164-derived peptide in fibroblasts in the face of all viral immunosubversive early gene functions.
Salmonella regulates polyubiquitination and surface expression of MHC class II antigens.
Viral interference with antigen presentation: trapping TAP.
COVID-19: Attacks Immune Cells and Interferes With Antigen Presentation Through MHC-Like Decoy System.
Chlamydia trachomatis immune evasion via downregulation of MHC class I surface expression involves direct and indirect mechanisms.
Viruses, cancer and non-self recognition.
A human herpesvirus 7 glycoprotein, U21, diverts major histocompatibility complex class I molecules to lysosomes.
Identification of a K(d)-restricted antigenic peptide encoded by murine cytomegalovirus early gene M84.
Viral immune evasion: a masterpiece of evolution.
The multiple immune-evasion genes of murine cytomegalovirus are not redundant: m4 and m152 inhibit antigen presentation in a complementary and cooperative fashion.
Antigens and immunoevasins: opponents in cytomegalovirus immune surveillance.
B. abortus RNA is the component involved in the down-modulation of MHC-I expression on human monocytes via TLR8 and the EGFR pathway.
Down-regulation of MHC class I is a property common to papillomavirus E5 proteins.
Cowpox virus protein CPXV012 eludes CTLs by blocking ATP binding to TAP.
Herpes simplex virus type I infection of mature dendritic cells leads to reduced LMP7-mRNA-expression levels.
Cytomegalovirus misleads its host by priming of CD8 T cells specific for an epitope not presented in infected tissues.

Genetic variability of the major histocompatibility complex class I homologue encoded by human cytomegalovirus leads to differential binding to the inhibitory receptor ILT2.
Vaccinia virus decreases major histocompatibility complex (MHC) class II antigen presentation, T-cell priming, and peptide association with MHC class II.
HIV-1 Nef binds a subpopulation of MHC-I throughout its trafficking itinerary and down-regulates MHC-I by perturbing both anterograde and retrograde trafficking.
Exploiting human herpesvirus immune evasion for therapeutic gain: potential and pitfalls.
Absence of intercellular adhesion molecule 1 expression in varicella zoster virus-infected keratinocytes during herpes zoster: another immune evasion strategy?
The capacity of UL49.5 proteins to inhibit TAP is widely distributed among members of the genus Varicellovirus.
MHC class I ubiquitination by a viral PHD/LAP finger protein.
Bovine herpesvirus 1 interferes with TAP-dependent peptide transport and intracellular trafficking of MHC class I molecules in human cells.
Molecular mechanism and species specificity of TAP inhibition by herpes simplex virus ICP47.
Priming of CD8+ T cells against cytomegalovirus-encoded antigens is dominated by cross-presentation.
Immunological evasion of immediate-early varicella zoster virus proteins.
Molecular mimicry by cytomegaloviruses. Function of cytomegalovirus-encoded homologues of G protein-coupled receptors, MHC class I heavy chains and chemokines.
Tumor-associated calreticulin variants functionally compromise the peptide loading complex and impair its recruitment of MHC-I.
IFN- $\alpha$ secretion by type 2 predendritic cells up-regulates MHC class I in the HIV-1-infected thymus.
A Coevolutionary Arms Race between Hosts and Viruses Drives Polymorphism and Polygenicity of NK Cell Receptors.
Evasion of CD8+ T cells is critical for superinfection by cytomegalovirus.
Inhibition of MHC-I by Brucella abortus is an early event during infection and involves EGFR pathway.
The Nature of Selection on the Major Histocompatibility Complex.
Signaling of a varicelloviral factor across the endoplasmic reticulum membrane induces destruction of the peptide-loading complex and immune evasion.
HIV-1 Nef disrupts intracellular trafficking of major histocompatibility complex class I, CD4, CD8, and CD28 by distinct pathways that share common elements.
Fluorescent TAP as a Platform for Virus-Induced Degradation of the Antigenic Peptide Transporter.
Chlamydia inhibits interferon gamma-inducible major histocompatibility complex class II expression by degradation of upstream stimulatory factor 1.
Absence of cross-presenting cells in the salivary gland and viral immune evasion confine cytomegalovirus immune control to effector CD4 T cells.
A negative feedback modulator of antigen processing evolved from a frameshift in the cowpox virus genome.
Nowhere to hide: unconventional translation yields cryptic peptides for immune surveillance.
Cutting edge: FcR-like 5 on innate B cells is targeted by a poxvirus MHC class I-like immunoevasin.
Downregulation of HLA-I by the molluscum contagiosum virus mc080 impacts NK-cell recognition and promotes CD8(+) T-cell evasion.
HLA-B locus products resist degradation by the human cytomegalovirus immunoevasin US11.
Structural mechanism of ER retrieval of MHC class I by cowpox.



The role of cytomegalovirus-encoded homologs of G protein-coupled receptors and chemokines in manipulation of and evasion from the immune system.
Interference with major histocompatibility complex class II-restricted antigen presentation in the brain by herpes simplex virus type 1: a possible mechanism of evasion of the immune response.
Plasma membrane profiling defines an expanded class of cell surface proteins selectively targeted for degradation by HCMV US2 in cooperation with UL141.
The ER-luminal domain of the HHV-7 immunoevasin U21 directs class I MHC molecules to lysosomes.
MHC class II gene expression is not induced in HPIV3-infected respiratory epithelial cells.
The immune evasion paradox: immunoevasins of murine cytomegalovirus enhance priming of CD8 T cells by preventing negative feedback regulation.
Advances in the Evolutionary Understanding of MHC Polymorphism.
Cytomegalovirus immune evasion sets the functional avidity threshold for protection by CD8 T cells.
Rinderpest virus isolates of different virulence vary in their capacity to infect bovine monocytes and macrophages.
Allele- and locus-specific recognition of class I MHC molecules by the immunomodulatory E3-19K protein from adenovirus.
Intracellular retention of the MHC class I-related chain B ligand of NKG2D by the human cytomegalovirus UL16 glycoprotein.
N Protein of Viral Hemorrhagic Septicemia Virus Suppresses STAT1-Mediated MHC Class II Transcription to Impair Antigen Presentation in Sea Perch, <i>Lateolabrax japonicus</i> .
TMEM129 is a Derlin-1 associated ERAD E3 ligase essential for virus-induced degradation of MHC-I.
Varicella-zoster virus productively infects mature dendritic cells and alters their immune function.
The viral E3 ubiquitin ligase mk3 uses the Derlin/p97 endoplasmic reticulum-associated degradation pathway to mediate down-regulation of major histocompatibility complex class I proteins.
Consequences of cytotoxic T lymphocyte interaction with major histocompatibility complex class I-expressing neurons in vivo.
VZV: immunobiology and host response.
Structure of the active domain of the herpes simplex virus protein ICP47 in water/sodium dodecyl sulfate solution determined by nuclear magnetic resonance spectroscopy.
Nuclear processing of nascent transcripts determines synthesis of full-length proteins and antigenic peptides.
Structure and function of the porcine TAP protein and its inhibition by the viral immune evasion protein ICP47.
Structure of the transporter associated with antigen processing trapped by herpes simplex virus.
Murine cytomegalovirus downregulates ERAAP and induces an unconventional T cell response to self.
Herpes simplex virus type 1 targets the MHC class II processing pathway for immune evasion.
Analysis of human cytomegalovirus US3 gene products.
Regulation of protein translation through mRNA structure influences MHC class I loading and T cell recognition.
ABC transporters in adaptive immunity.
Control of MHC II antigen presentation by ubiquitination.
Antigen presentation assays to investigate uncharacterized immunoregulatory genes.
Different serotypes of dengue viruses differently regulate the expression of the host cell antigen processing machinery.
An endocytic YXXΦ (YRRF) cargo sorting motif in the cytoplasmic tail of murine cytomegalovirus AP2 'adapter adapter' protein m04/gp34 antagonizes virus evasion of natural killer cells.

Insufficient Antigen Presentation Due to Viral Immune Evasion Explains Lethal Cytomegalovirus Organ Disease After Allogeneic Hematopoietic Cell Transplantation.
Bovine herpesvirus type 1 (BHV-1) mutant lacking U(L)49.5 luminal domain residues 30-32 and cytoplasmic tail residues 80-96 induces more rapid onset of virus neutralizing antibody and cellular immune responses in calves than the wild-type strain Cooper.
Enhancement of Antigen Presentation by Deletion of Viral Immune Evasion Genes Prevents Lethal Cytomegalovirus Disease in Minor Histocompatibility Antigen-Mismatched Hematopoietic Cell Transplantation.
Interaction of flaviviruses with cells of the vertebrate host and decoy of the immune response.
The synthesis of truncated polypeptides for immune surveillance and viral evasion.
Molecular mechanisms of human herpes viruses inferring with host immune surveillance.
Discerning regulation of cis- and trans-presentation of CD8+ T-cell epitopes by EBV-encoded oncogene LMP-1 through self-aggregation.
Early gene m18, a novel player in the immune response to murine cytomegalovirus.
A dual inhibition mechanism of herpesviral ICP47 arresting a conformationally thermostable TAP complex.
Brucella abortus down-regulates MHC class II by the IL-6-dependent inhibition of CIITA through the downmodulation of IFN regulatory factor-1 (IRF-1).
Simplexviruses Successfully Adapt to Their Host by Fine-Tuning Immune Responses.
The efficacy of antigen processing is critical for protection against cytomegalovirus disease in the presence of viral immune evasion proteins.
Bovine herpesvirus type 1 (BHV-1) UL49.5 luminal domain residues 30 to 32 are critical for MHC-I down-regulation in virus-infected cells.
TNF-induced target cell killing by CTL activated through cross-presentation.
Antibody-induced endocytosis of viral glycoproteins and major histocompatibility complex class I on pseudorabies virus-infected monocytes.
Identification of class I MHC regions which bind to the adenovirus E3-19k protein.
Experimental viral evolution to specific host MHC genotypes reveals fitness and virulence trade-offs in alternative MHC types.
Cytomegalovirus immune evasion by perturbation of endosomal trafficking.
Shigella effector IpaH4.5 targets 19S regulatory particle subunit RPN13 in the 26S proteasome to dampen cytotoxic T lymphocyte activation.
Spatial separation of HLA-DM/HLA-DR interactions within MIIC and phagosome-induced immune escape.
The N terminus of adenovirus type 12 E1A inhibits major histocompatibility complex class I expression by preventing phosphorylation of NF-kappaB p65 Ser276 through direct binding.
Human CD8+ CTL recognition and in vitro lysis of herpes simplex virus-infected cells by a non-MHC restricted mechanism.
Direct Priming of CD8(+) T Cells Persists in the Face of Cowpox Virus Inhibitors of Antigen Presentation.
The human herpesvirus-7 (HHV-7) U21 immunoevasin subverts NK-mediated cytotoxicity through modulation of MICA and MICB.
Salivary gland resident APCs are Flt3L- and CCR2-independent macrophage-like cells incapable of cross-presentation.
Effective inhibition of K(b)- and D(b)-restricted antigen presentation in primary macrophages by murine cytomegalovirus.
[Varicella-zoster virus (VZV)].

A battle for survival: immune control and immune evasion in murine gamma-herpesvirus-68 infection.
In Vivo Killing Capacity of Cytotoxic T Cells Is Limited and Involves Dynamic Interactions and T Cell Cooperativity.
Herpesviral capture of immunomodulatory host genes.
Costimulation provided by DNA immunization enhances antitumor immunity.
Decoupling the role of ubiquitination for the dislocation versus degradation of major histocompatibility complex (MHC) class I proteins during endoplasmic reticulum-associated degradation (ERAD).
Structure and dynamics of membrane-associated ICP47, a viral inhibitor of the MHC I antigen-processing machinery.
Adenovirus early region 3 antiapoptotic 10.4K, 14.5K, and 14.7K genes decrease the incidence of autoimmune diabetes in NOD mice.
Dendritic cells as Achilles' heel and Trojan horse during varicella zoster virus infection.
The structure of the cytomegalovirus-encoded m04 glycoprotein, a prototypical member of the m02 family of immunoevasins.
Herpes simplex inhibits the capacity of lymphoblastoid B cell lines to stimulate CD4+ T cells.
Immune evasion by adenoviruses: a window into host-virus adaptation.
Manipulation of the Innate Immune Response by Varicella Zoster Virus.
Structural Models for Roseolovirus U20 And U21: Non-Classical MHC-I Like Proteins From HHV-6A, HHV-6B, and HHV-7.
The effect of bovine viral diarrhea virus on bovine monocyte phenotype.
The antigen processing-associated transporter gene polymorphism: Role on gene and protein expression in HPV-infected pre-cancerous cervical lesion.
The herpes simplex virus-1 encoded glycoprotein B diverts HLA-DR into the exosome pathway.
Vaccine-induced memory CD8+ T cells cannot prevent central nervous system virus reactivation.
Infection and Functional Modulation of Human Monocytes and Macrophages by Varicella-Zoster Virus.
Immune interactions at the maternal-fetal interface: a focus on antigen presentation.
Mouse Cytomegalovirus m153 Protein Stabilizes Expression of the Inhibitory NKR-P1B Ligand Clr-b.
Human NKG2D-ligands: cell biology strategies to ensure immune recognition.
Insights into the pathogenesis of varicella viruses.
A novel transmembrane domain mediating retention of a highly motile herpesvirus glycoprotein in the endoplasmic reticulum.
Combined single-cell quantitation of host and SIV genes and proteins ex vivo reveals host-pathogen interactions in individual cells.
Herpes simplex virus 1 glycoprotein B and US3 collaborate to inhibit CD1d antigen presentation and NKT cell function.
Non-classical HLA class I molecules and their potential role in viral infections.
Immune evasion as a pathogenic mechanism of varicella zoster virus.
Varicella-zoster virus inhibits autophagosome-lysosome fusion and the degradation stage of mTOR-mediated autophagic flux.
Proteasome function shapes innate and adaptive immune responses.
Cowpox virus employs a two-pronged strategy to outflank MHCI antigen presentation.
Brucella Peptide Cross-Reactive Major Histocompatibility Complex Class I Presentation Activates SIINFEKL-Specific T Cell Receptor-Expressing T Cells.
Human DUX4 and mouse Dux interact with STAT1 and broadly inhibit interferon-stimulated gene induction.

Superantigens of mouse mammary tumor virus.
Subversion of immunoproteasome subunit expression in dengue virus serotype 2-infected HepG2 cells.
Bordetella pertussis infection of primary human monocytes alters HLA-DR expression.
Insights into the function of tegument proteins from the varicella zoster virus.
Accumulating evidence suggests that several AB-toxins subvert the endoplasmic reticulum-associated protein degradation pathway to enter target cells.
Immunity to bovine herpesvirus 1: II. Adaptive immunity and vaccinology.
Viral strategies of immune evasion.
Poxvirus-encoded serpins do not prevent cytolytic T cell-mediated recovery from primary infections.
Structural Conservation and Functional Diversity of the Poxvirus Immune Evasion (PIE) Domain Superfamily.
Characterization of the HHV-6B U20 Immuno-evasin.
Intracellular Sequestration of the NKG2D Ligand MIC B by Species F Adenovirus.
Structures of the four Ig-like domain LILRB2 and the four-domain LILRB1 and HLA-G1 complex.
Sorting Motifs in the Cytoplasmic Tail of the Immunomodulatory E3/49K Protein of Species D Adenoviruses Modulate Cell Surface Expression and Ectodomain Shedding.
A GPI anchor explains the unique biological features of the common NKG2D-ligand allele MICA*008.
Genotyping of human killer-cell immunoglobulin-like receptor genes by polymerase chain reaction with sequence-specific primers: an update.
Cytotoxic T lymphocytes: all roads lead to death.
Varicella-zoster virus gH:gL contains a structure reactive with the anti-human gamma chain of IgG near the glycosylation site.
Immune and cytokine/chemokine responses of PBMCs in rotavirus-infected rhesus infants and their significance in viral pathogenesis.

Table A2-35, Cluster 34

Cluster 34 focuses on B-cell lymphoma, emphasizing mechanisms of immune evasion by lymphoma cells (271)
Targeting the tumor microenvironment in B-cell lymphoma: challenges and opportunities.
Mechanisms of Immune Evasion and Immune Modulation by Lymphoma Cells.
The immune landscape and response to immune checkpoint blockade therapy in lymphoma.
A single-cell atlas of diffuse large B cell lymphoma.
Molecular profiling of EBV associated diffuse large B-cell lymphoma.
High frequency of inactivating tetraspanin C D37 mutations in diffuse large B-cell lymphoma at immune-privileged sites.
Genomic rearrangements involving programmed death ligands are recurrent in primary mediastinal large B-cell lymphoma.
Dissecting aggressive B-cell lymphoma through genomic analysis - What is clinically relevant?
Immune evasion-related extranodal large B-cell lymphoma: A report of six patients with neoplastic PD-L1-positive extranodal diffuse large B-cell lymphoma.
Spatial signatures identify immune escape via PD-1 as a defining feature of T-cell/histiocyte-rich large B-cell lymphoma.
Profiling Immune Escape in Hodgkin's and Diffuse large B-Cell Lymphomas Using the Transcriptome and Immunostaining.

Frequent structural variations involving programmed death ligands in Epstein-Barr virus-associated lymphomas.
Mutational dynamics and immune evasion in diffuse large B-cell lymphoma explored in a relapse-enriched patient series.
The biology of Hodgkin's lymphoma.
The molecular pathogenesis of primary mediastinal large B-cell lymphoma.
Combined multiomics analysis reveals the mechanism of CENPF overexpression-mediated immune dysfunction in diffuse large B-cell lymphoma in vitro.
Tumor Microenvironment in Diffuse Large B-Cell Lymphoma: Role and Prognosis.
The tumour microenvironment in B cell lymphomas.
Evaluation of the diagnostic and prognostic value of PDL1 expression in Hodgkin and B-cell lymphomas.
Checkpoint Inhibition in Non-Hodgkin's Lymphoma.
Adoptive immunotherapy for Hodgkin's lymphoma.
Microbial dysbiosis is associated with aggressive histology and adverse clinical outcome in B-cell non-Hodgkin lymphoma.
Pembrolizumab in newly diagnosed EBV-negative extranodal natural killer/T-cell lymphoma: A case report.
Epstein-Barr Virus-Positive Lymphomas Exploit Ectonucleotidase Activity To Limit Immune Responses and Prevent Cell Death.
The pathobiology of primary testicular diffuse large B-cell lymphoma: Implications for novel therapies.
lncRNA SNHG14 promotes oncogenesis and immune evasion in diffuse large-B-cell lymphoma by sequestering miR-152-3p.
Newly identified mechanisms in B-cell non-Hodgkin lymphomas uncovered by next-generation sequencing.
EBV-positive B-cell lymphomas and lymphoproliferative disorders: Review from the perspective of immune escape and immunodeficiency.
Inhibition of STAT3 activity re-activates anti-tumor immunity but fails to restore the immunogenicity of tumor cells in a B-cell lymphoma model.
Therapy Strategy of CD47 in Diffuse Large B-Cell Lymphoma (DLBCL).
Primary mediastinal large B-cell lymphoma.
Anti-CD47 immunotherapy in combination with BCL-2 inhibitor to enhance anti-tumor activity in B-cell lymphoma.
Regulatory myeloid cells: an underexplored continent in B-cell lymphomas.
Mutational landscape of immune surveillance genes in diffuse large B-cell lymphoma.
PD-1/PD-L1 immune checkpoint and p53 loss facilitate tumor progression in activated B-cell diffuse large B-cell lymphomas.
Genetic lesions and targeted therapy in Hodgkin lymphoma.
PD-1 Blockade Can Restore Functions of T-Cells in Epstein-Barr Virus-Positive Diffuse Large B-Cell Lymphoma In Vitro.
Impact of a Faulty Germinal Center Reaction on the Pathogenesis of Primary Diffuse Large B Cell Lymphoma of the Central Nervous System.
[The story about Hodgkin's lymphoma].
Primary Mediastinal B-Cell Lymphoma in Children and Young Adults.
Epstein-Barr virus-negative diffuse large B-cell lymphoma hosts intra- and peritumoral B-cells with activated Epstein-Barr virus.

B cell lymphoma progression promotes the accumulation of circulating Ly6Clo monocytes with immunosuppressive activity.
Genomic Analyses Identify Recurrent Alterations in Immune Evasion Genes in Diffuse Large B-Cell Lymphoma, Leg Type.
HIV-associated Hodgkin lymphoma.
Immuno-oncology for B-cell lymphomas.
EZH2 Inhibitors: Take It EZy, It Is All About Context.
The PD-1/PD-L1 Checkpoint in Normal Germinal Centers and Diffuse Large B-Cell Lymphomas.
CD2-negative lymphoma-associated T-cells: a potential mechanism of immune-evasion in diffuse large B-cell lymphoma.
Lymphoma endothelium preferentially expresses Tim-3 and facilitates the progression of lymphoma by mediating immune evasion.
EBNA3B-deficient EBV promotes B cell lymphomagenesis in humanized mice and is found in human tumors.
Impact of MYC on Anti-Tumor Immune Responses in Aggressive B Cell Non-Hodgkin Lymphomas: Consequences for Cancer Immunotherapy.
Characterization of DLBCL with a PMBL gene expression signature.
Epigenetic, Metabolic, and Immune Crosstalk in Germinal-Center-Derived B-Cell Lymphomas: Unveiling New Vulnerabilities for Rational Combination Therapies.
[Genetic Alterations Involving PD-L1/PD-L2 in Human Malignancies].
B cell lymphomas of C57L/J mice; the role of natural killer cells and T helper cells in lymphoma development and growth.
Mass cytometry defines distinct immune profile in germinal center B-cell lymphomas.
Host immune response in B-cell lymphomas: friend or foe?
Engaging the Innate and Adaptive Antitumor Immune Response in Lymphoma.
The genomic and transcriptional landscape of primary central nervous system lymphoma.
Biology of classical Hodgkin lymphoma: implications for prognosis and novel therapies.
Murine models of B-cell lymphomas: promising tools for designing cancer therapies.
Toll-like receptors in the pathogenesis of human B cell malignancies.
Immune reactions in classical Hodgkin's lymphoma.
Integrative genomic analysis identifies key pathogenic mechanisms in primary mediastinal large B-cell lymphoma.
[Circulating Tumor DNA Analysis in Lymphomas].
Clinical and molecular characterization of diffuse large B-cell lymphomas with 13q14.3 deletion.
Molecular biology of Hodgkin lymphoma.
Tumor-activated lymph node fibroblasts suppress T cell function in diffuse large B cell lymphoma.
HHV8-positive, EBV-positive Hodgkin lymphoma-like large B cell lymphoma: expanding the spectrum of HHV8 and EBV-associated lymphoproliferative disorders.
Classic Hodgkin Lymphoma - Old Disease, New Directions: An Update on Pathology, Molecular Features and Biological Prognostic Markers.
Tumor cell-derived IL-10 promotes cell-autonomous growth and immune escape in diffuse large B-cell lymphoma.
Novel Approaches in Molecular Characterization of Classical Hodgkin Lymphoma.
Detection of Aberrant CD58 Expression in a Wide Spectrum of Lymphoma Subtypes: Implications for Treatment Resistance.
Tumor B cells from non-Hodgkin's lymphoma are resistant to CD95 (Fas/Apo-1)-mediated apoptosis.

MicroRNA signature in classical Hodgkin lymphoma.
The Immunology of DLBCL.
Deep sequencing reveals clonal evolution patterns and mutation events associated with relapse in B-cell lymphomas.
Gray-zone Lymphoma Between cHL and Large B-Cell Lymphoma: A Histopathologic Series From the LYSA.
Single-Cell Transcriptome Analysis Reveals Disease-Defining T-cell Subsets in the Tumor Microenvironment of Classic Hodgkin Lymphoma.
Molecular pathogenesis of germinal center-derived B cell lymphomas.
Induction of antigen-specific effector-phase tolerance following vaccination against a previously ignored B-cell lymphoma.
Genomic landscape of mature B-cell non-Hodgkin lymphomas - an appraisal from lymphomagenesis to drug resistance.
Immunotherapy in aggressive B-cell lymphomas.
An Update on the Pathology and Molecular Features of Hodgkin Lymphoma.
Epstein-Barr virus-positive diffuse large B-cell lymphoma features disrupted antigen capture/presentation and hijacked T-cell suppression.
Role of the tumor microenvironment in mature B-cell lymphoid malignancies.
Pathogenesis of human B cell lymphomas.
Primary and acquired resistance mechanisms to immune checkpoint inhibition in Hodgkin lymphoma.
From genetics to the clinic: a translational perspective on follicular lymphoma.
Large-scale microarray profiling reveals four stages of immune escape in non-Hodgkin lymphomas.
Molecular Pathogenesis of Hodgkin Lymphoma: Past, Present, Future.
Immune Microenvironment Features and Dynamics in Hodgkin Lymphoma.
Current status of prognostication in classical Hodgkin lymphoma.
A miR-129-5P/ARID3A Negative Feedback Loop Modulates Diffuse Large B Cell Lymphoma Progression and Immune Evasion Through Regulating the PD-1/PD-L1 Checkpoint.
PD-1-PD-1 ligand interaction contributes to immunosuppressive microenvironment of Hodgkin lymphoma.
Stromal interferon- $\gamma$ signaling and cross-presentation are required to eliminate antigen-loss variants of B cell lymphomas in mice.
Berberine exerts anti-tumor activity in diffuse large B-cell lymphoma by modulating c-myc/CD47 axis.
Frequent genetic alterations in immune checkpoint-related genes in intravascular large B-cell lymphoma.
T-cell lymphoma secondary to checkpoint inhibitor therapy.
Challenges and perspectives in the immunotherapy of Hodgkin lymphoma.
CRISPR/Cas9-Mediated Foxp1 Silencing Restores Immune Surveillance in an Immunocompetent A20 Lymphoma Model.
Phase I/Ib Study of the Efficacy and Safety of Buparlisib and Ibrutinib Therapy in MCL, FL, and DLBCL with Serial Cell-Free DNA Monitoring.
Follicular lymphoma: The long and winding road leading to your cure?
CD58 mutations are common in Hodgkin lymphoma cell lines and loss of CD58 expression in tumor cells occurs in Hodgkin lymphoma patients who relapse.
T-Cell Therapy for Lymphoma Using Nonengineered Multiantigen-Targeted T Cells Is Safe and Produces Durable Clinical Effects.
Targetable genetic features of primary testicular and primary central nervous system lymphomas.

Biology and therapy of primary mediastinal B-cell lymphoma: current status and future directions.
Mutational Profile and Clonal Evolution of Relapsed/Refractory Diffuse Large B-Cell Lymphoma.
Diffuse large B-cell lymphomas have spatially defined, tumor immune microenvironments revealed by high-parameter imaging.
T-cell inflamed tumor microenvironment predicts favorable prognosis in primary testicular lymphoma.
METTL3 enhances the effect of YTHDF1 on NEDD1 mRNA stability by m6A modification in diffuse large B-cell lymphoma cells.
Checkpoint Inhibitors for the Treatment of Hodgkin Lymphoma.
Lymphomagenesis in Hodgkin lymphoma.
Immune priming with avelumab and rituximab prior to R-CHOP in diffuse large B-cell lymphoma: the phase II AvR-CHOP study.
Interleukin-10 counteracts T-helper type 1 responses in B-cell lymphoma and is a target for tumor immunotherapy.
Ratios of T-cell immune effectors and checkpoint molecules as prognostic biomarkers in diffuse large B-cell lymphoma: a population-based study.
Germinal center-derived lymphomas: The darkest side of humoral immunity.
Signaling through CD80: an approach for treating lymphomas.
Immune escape mechanisms for TCRLBCL.
The microenvironment in classical Hodgkin lymphoma: an actively shaped and essential tumor component.
Genetically Engineered Mouse Models Support a Major Role of Immune Checkpoint-Dependent Immunosurveillance Escape in B-Cell Lymphomas.
MYD88 L265P mutation promoted malignant B cell resistance against T cell-mediated cytotoxicity via upregulating the IL-10/STAT3 cascade.
Gene expression predicts overall survival in paraffin-embedded tissues of diffuse large B-cell lymphoma treated with R-CHOP.
Immunotherapies for Hodgkin's lymphoma.
The dangers of déjà vu: memory B cells as the cells of origin of ABC-DLBCLs.
Evidence for IL-35 Expression in Diffuse Large B-Cell Lymphoma and Impact on the Patient's Prognosis.
Follicular lymphoma cells induce T-cell immunologic synapse dysfunction that can be repaired with lenalidomide: implications for the tumor microenvironment and immunotherapy.
mRNA-Based Anti-TCR CDR3 Tumour Vaccine for T-Cell Lymphoma.
The role of trogocytosis in immune surveillance of Hodgkin lymphoma.
Genomic analyses of flow-sorted Hodgkin Reed-Sternberg cells reveal complementary mechanisms of immune evasion.
Double-edge Role of B Cells in Tumor Immunity: Potential Molecular Mechanism.
NFkB Pathway and Hodgkin Lymphoma.
[Immune evasion mechanism and its clinical application value in Hodgkin's lymphoma].
Formation of the Immunosuppressive Microenvironment of Classic Hodgkin Lymphoma and Therapeutic Approaches to Counter It.
Distinct roles for PARP-1 and PARP-2 in c-Myc-driven B-cell lymphoma in mice.
T-Cell Traffic Jam in Hodgkin's Lymphoma: Pathogenetic and Therapeutic Implications.
STAT3 Activation and Oncogenesis in Lymphoma.
Heat Shock Proteins in Lymphoma Immunotherapy.
Novel therapeutics for aggressive non-Hodgkin's lymphoma.



Low T-cell proportion in the tumor microenvironment is associated with immune escape and poor survival in diffuse large B-cell lymphoma.
Epstein-Barr virus-positive pyothorax-associated lymphoma expresses CCL17 and CCL22 chemokines that attract CCR4-expressing regulatory T cells.
Gene expression meta-analysis reveals immune response convergence on the IFN $\gamma$ -STAT1-IRF1 axis and adaptive immune resistance mechanisms in lymphoma.
Optimizing therapy for nodal marginal zone lymphoma.
T Cell Histiocyte Rich Large B Cell Lymphoma Presenting as Hemophagocytic Lymphohistiocytosis: An Uncommon Presentation of a Rare Disease.
Aberrant expression of the Th2 cytokine IL-21 in Hodgkin lymphoma cells regulates STAT3 signaling and attracts Treg cells via regulation of MIP-3 $\alpha$ .
The dual role of CD70 in B-cell lymphomagenesis.
Ibrutinib.
Expression of programmed death-1 (CD279) in primary cutaneous B-cell lymphomas with correlation to lymphoma entities and biological behaviour.
Leukemic and meningeal relapse of CD5+ intravascular large B-cell lymphoma with down-modulation of CD20 after rituximab therapy.
Intratumor heterogeneity and T cell exhaustion in primary CNS lymphoma.
Immune evasion of mantle cell lymphoma: expression of B7-H1 leads to inhibited T-cell response to and killing of tumor cells.
Microenvironment Cell Contribution to Lymphoma Immunity.
Therapeutic Strategies against Epstein-Barr Virus-Associated Cancers Using Proteasome Inhibitors.
Genomic Landscape of Hodgkin Lymphoma.
CC chemokine receptor 2 (CCR2) expression promotes diffuse large B-Cell lymphoma survival and invasion.
Transcriptomic Abnormalities in Epstein Barr Virus Associated T/NK Lymphoproliferative Disorders.
RNA fingerprints provide direct evidence for the inhibitory role of TGF $\beta$ and PD-1 on CD4+ T cells in Hodgkin lymphoma.
Tumor-associated antigen PRAME exhibits dualistic functions that are targetable in diffuse large B cell lymphoma.
Relapsed Primary Central Nervous System Lymphoma: Current Advances.
Drug-induced aneuploidy and polyploidy is a mechanism of disease relapse in MYC/BCL2-addicted diffuse large B-cell lymphoma.
Adoptive T-cell therapy for Epstein-Barr virus-positive Hodgkin's disease.
Novel agents for the treatment of Hodgkin lymphoma.
Inactivation of CREBBP expands the germinal center B cell compartment, down-regulates MHCII expression and promotes DLBCL growth.
Nivolumab for classical Hodgkin's lymphoma after failure of both autologous stem-cell transplantation and brentuximab vedotin: a multicentre, multicohort, single-arm phase 2 trial.
The interplay between Epstein-Barr virus and the immune system: a rationale for adoptive cell therapy of EBV-related disorders.
Mesenchymal stromal cells promote B-cell lymphoma in lacrimal glands by inducing immunosuppressive microenvironment.
Molecularly Stratified Treatment Options in Primary Refractory DLBCL/HGBL with MYC and BCL2 or BCL6 Rearrangements (HGBL, NOS with MYC/BCL6).
Epstein Barr Virus Associated Lymphomas and Epithelia Cancers in Humans.
Novel agents and immune invasion in Hodgkin lymphoma.

Alterations of the CD58 gene in classical Hodgkin lymphoma.
Beclin1-armed oncolytic Vaccinia virus enhances the therapeutic efficacy of R-CHOP against lymphoma in vitro and in vivo.
Immune subtyping of extranodal NK/T-cell lymphoma: a new biomarker and an immune shift during disease progression.
High Expression of IKZF2 in Malignant T Cells Promotes Disease Progression in Cutaneous T Cell Lymphoma.
Concordant bone marrow involvement of diffuse large B-cell lymphoma represents a distinct clinical and biological entity in the era of immunotherapy.
The circuitry of the tumor microenvironment in adult and pediatric Hodgkin lymphoma: cellular composition, cytokine profile, EBV, and exosomes.
The aggressiveness of murine lymphomas selected in vivo by growth rate correlates with galectin-1 expression and response to cyclophosphamide.
Genomic and Single-Cell Landscape Reveals Novel Drivers and Therapeutic Vulnerabilities of Transformed Cutaneous T-cell Lymphoma.
Expression of human tumor-associated antigen RCAS1 in adult T-cell leukemia/lymphoma.
Collapse of the CD27+ B-cell compartment associated with systemic plasmacytosis in patients with advanced melanoma and other cancers.
Control of B-cell lymphoma by therapeutic vaccination and acquisition of immune resistance is independent of direct tumour IFN-gamma signalling.
IDO, TDO, and AHR overexpression is associated with poor outcome in diffuse large B-cell lymphoma patients in the rituximab era.
Expression of CD137 on Hodgkin and Reed-Sternberg cells inhibits T-cell activation by eliminating CD137 ligand expression.
Galectins in hematological malignancies.
Mutations in early follicular lymphoma progenitors are associated with suppressed antigen presentation.
Central Nervous System Lymphomas.
Large B-cell Lymphomas of Immune-Privileged Sites Relapse via Parallel Clonal Evolution from a Common Progenitor B Cell.
Epigenetic reprogramming sensitizes immunologically silent EBV+ lymphomas to virus-directed immunotherapy.
EBV infection determines the immune hallmarks of plasmablastic lymphoma.
PD-1 and LAG-3 expression in EBV-associated pediatric Hodgkin lymphoma has influence on survival.
Novel cell enrichment technique for robust genetic analysis of archival classical Hodgkin lymphoma tissues.
Treatment of Epstein-Barr virus-associated malignancies with specific T cells.
Screening for adenoviruses in haematological neoplasia: High prevalence in mantle cell lymphoma.
Cytotoxic T lymphocyte therapy for Epstein-Barr virus+ Hodgkin's disease.
Cell growth in aggregates determines gene expression, proliferation, survival, chemoresistance, and sensitivity to immune effectors in follicular lymphoma.
Identification of FAT4 as a positive prognostic biomarker in DLBCL by comprehensive genomic analysis.
Spatial and molecular profiling of the mononuclear phagocyte network in classic Hodgkin lymphoma.
Combination of cyclophosphamide, rituximab, and intratumoral CpG oligodeoxynucleotide successfully eradicates established B cell lymphoma.

Filling the Gap: The Immune Therapeutic Armamentarium for Relapsed/Refractory Hodgkin Lymphoma.
Checkpoint protein expression in the tumor microenvironment defines the outcome of classical Hodgkin lymphoma patients.
PIM Kinases Promote Survival and Immune Escape in Primary Mediastinal Large B-Cell Lymphoma through Modulation of JAK-STAT and NF- $\kappa$ B Activity.
Efficacy of an adapted granzyme B-based anti-CD30 cytolytic fusion protein against PI-9-positive classical Hodgkin lymphoma cells in a murine model.
Ectopic CD137 expression facilitates the escape of Hodgkin and Reed-Sternberg cells from immunosurveillance.
Heterogeneous expression of interleukin-18 and its receptor in B-cell lymphoproliferative disorders deriving from naive, germinal center, and memory B lymphocytes.
Comprehensive genomic analysis identifying heterogeneity in peripheral T-cell lymphoma.
AMD3100-Mediated CXCR4 Inhibition Impairs Development of Primary Lymphoma of the Central Nervous System.
EZH2 inhibitors restore epigenetically silenced CD58 expression in B-cell lymphomas.
Nivolumab for Relapsed/Refractory Diffuse Large B-Cell Lymphoma in Patients Ineligible for or Having Failed Autologous Transplantation: A Single-Arm, Phase II Study.
Preclinical Development of a Bispecific Antibody that Safely and Effectively Targets CD19 and CD47 for the Treatment of B-Cell Lymphoma and Leukemia.
CREBBP and STAT6 co-mutation and 16p13 and 1p36 loss define the t(14;18)-negative diffuse variant of follicular lymphoma.
Immunotherapy for Epstein-Barr virus-associated cancers in children.
Mechanisms of Immune Tolerance in Leukemia and Lymphoma.
Targeting the JAK-STAT pathway in lymphoma: a focus on pacritinib.
Cutaneous T-cell lymphoma: Biologic targets for therapy.
Adoptive immunotherapy for EBV-associated malignancies.
Update on Novel Therapeutics for Primary CNS Lymphoma.
Galectin-1 serum levels reflect tumor burden and adverse clinical features in classical Hodgkin lymphoma.
Safety and efficacy of immune checkpoint inhibitors after allogeneic hematopoietic cell transplantation.
B cells, plasma cells and antibody repertoires in the tumour microenvironment.
Concepts in immuno-oncology: tackling B cell malignancies with CD19-directed bispecific T cell engager therapies.
Identification of granulocytic myeloid-derived suppressor cells (G-MDSCs) in the peripheral blood of Hodgkin and non-Hodgkin lymphoma patients.
CD137 signaling in Hodgkin and Reed-Sternberg cell lines induces IL-13 secretion, immune deviation and enhanced growth.
Clinical application of genomic aberrations in adult T-cell leukemia/lymphoma.
Viral transduction of primary human lymphoma B cells reveals mechanisms of NOTCH-mediated immune escape.
Immunogenetic therapy for B-cell malignancies.
Immunotherapy against EBV-lymphoma in recipients of HSCT.
Landscape of B cell immunity and related immune evasion in human cancers.
Distinct immune signatures in chronic lymphocytic leukemia and Richter syndrome.
The role of T cells in the microenvironment of Hodgkin lymphoma.

Novel approach to identify putative Epstein-Barr-virus microRNAs regulating host cell genes with relevance in tumor biology and immunology.
Temporal multiomic modeling reveals a B-cell receptor proliferative program in chronic lymphocytic leukemia.
Immunotherapy for Hodgkin's disease.
Toll-like receptor-4 signaling in mantle cell lymphoma: effects on tumor growth and immune evasion.
Innate predisposition to immune escape in follicular lymphoma cells.
Hodgkin disease and the role of the immune system.
Exhausted Markers in Cutaneous T-Cell Lymphoma: The Face that Launched a Thousand Ships.
[Treatment approach for relapse of adult T-cell leukemia/lymphoma after allogeneic hematopoietic stem cell transplantation].
Molecular heterogeneity in peripheral T-cell lymphoma, not otherwise specified revealed by comprehensive genetic profiling.
Facts and Hopes in the Relationship of EBV with Cancer Immunity and Immunotherapy.
Proteomic changes in cerebrospinal fluid from primary central nervous system lymphoma patients are associated with protein ectodomain shedding.
Vaccine and antibody-directed T cell tumour immunotherapy.
Expression of PIM kinases in Reed-Sternberg cells fosters immune privilege and tumor cell survival in Hodgkin lymphoma.
Expression of the signal transduction molecule zeta in peripheral and tumour-associated lymphocytes in Hodgkin's disease in relation to the Epstein-Barr virus status of the tumour cells.
[Genetic landscape of adult T-cell leukemia/lymphoma].
PD-L1 mediated the differentiation of tumor-infiltrating CD19(+) B lymphocytes and T cells in Invasive breast cancer.
The BLIMP1-EZH2 nexus in a non-Hodgkin lymphoma.
CD200 expression in hematopoietic neoplasms: Beyond a marker for diagnosis of B-cell neoplasms.
Differential expression of thymus and activation regulated chemokine and its receptor CCR4 in nodal and cutaneous anaplastic large-cell lymphomas and Hodgkin's disease.
Perivascular microenvironment in primary central nervous system lymphomas: the role of chemokines and the endothelin B receptor.
Expression of soluble CD27 in extranodal natural killer/T-cell lymphoma, nasal type: potential as a biomarker for diagnosis and CD27/CD70-targeted therapy.
Siglec-15 Promotes Evasion of Adaptive Immunity in B-cell Acute Lymphoblastic Leukemia.
Methylation of the IL-12Rbeta2 gene as novel tumor escape mechanism for pediatric B-acute lymphoblastic leukemia cells.
Nucleophosmin-anaplastic lymphoma kinase: the ultimate oncogene and therapeutic target.
Macrophages and dendritic cells as actors in the immune reaction of classical Hodgkin lymphoma.
Mutation Analysis for TP53 in Chronic-Type Adult T-Cell Leukemia/Lymphoma.
Targeted deep sequencing reveals clonal and subclonal mutational signatures in Adult T-cell leukemia/lymphoma and defines an unfavorable indolent subtype.
Immune Response against ALK in Children with ALK-Positive Anaplastic Large Cell Lymphoma.
Overexpression of interleukin-13 in a murine T-cell lymphoma: a possible factor of DL-induced immunosuppression and tumor progression.
Immunophenotypic Spectrum and Genomic Landscape of Refractory Celiac Disease Type II.
Modeling the proteome of a Marek's disease transformed cell line: a natural animal model for CD30 overexpressing lymphomas.

Viral induction and targeted inhibition of galectin-1 in EBV+ posttransplant lymphoproliferative disorders.
Survival and Immunosuppression Induced by Hepatocyte Growth Factor in Chronic Lymphocytic Leukemia.
Clinical Applications of Genomic Alterations in ATLL: Predictive Markers and Therapeutic Targets.
Gastric carcinoma: monoclonal epithelial malignant cells expressing Epstein-Barr virus latent infection protein.
Blocking "don't eat me" signal of CD47-SIRPα in hematological malignancies, an in-depth review.
The potential evasion of immune surveillance in mucosa associated lymphoid tissue lymphoma by DcR2-mediated up-regulation of nuclear factor-κB.
Metabolic activity of extranodal NK/T cell lymphoma on (18)F-FDG PET/CT according to immune subtyping.
Lenalidomide in the treatment of chronic lymphocytic leukemia.
EMAP II Expression Is Increased on Peripheral Blood Cells from Non-Hodgkin Lymphoma.
Treatment of AML Relapse After Allo-HCT.
Emergence of clone with PHF6 nonsense mutation in chronic myelomonocytic leukemia at relapse after allogeneic HCT.
Clonal cytogenetic changes and myeloma relapse after reduced intensity conditioning allogeneic transplantation.

Table A2-36, Cluster 35

Cluster 35 focuses on HIV-1, emphasizing 1) generation of vaccine-based broadly neutralizing antibodies against HIV-1 and 2) HIV-1 mutations and escape (456)
Broadly Neutralizing Antibodies against HIV-1 As a Novel Aspect of the Immune Response.
The Antibody Response against HIV-1.
Viral evolution and escape during acute HIV-1 infection.
The Hard Way towards an Antibody-Based HIV-1 Env Vaccine: Lessons from Other Viruses.
Candida albicans delays HIV-1 replication in macrophages.
HIV-1: the confounding variables of virus neutralization.
HIV-1 escapes from N332-directed antibody neutralization in an elite neutralizer by envelope glycoprotein elongation and introduction of unusual disulfide bonds.
Strategies for HIV-1 vaccines that induce broadly neutralizing antibodies.
Envelope-specific antibodies and antibody-derived molecules for treating and curing HIV infection.
Structural Constraints at the Trimer Apex Stabilize the HIV-1 Envelope in a Closed, Antibody-Protected Conformation.
In vivo functions of CPSF6 for HIV-1 as revealed by HIV-1 capsid evolution in HLA-B27-positive subjects.
Passive sexual transmission of human immunodeficiency virus type 1 variants and adaptation in new hosts.
Antibody polyspecificity and neutralization of HIV-1: a hypothesis.
Antibody-virus co-evolution in HIV infection: paths for HIV vaccine development.
HIV-1 mutates to adapt in fluxing environments.
Long-Acting BMS-378806 Analogues Stabilize the State-1 Conformation of the Human Immunodeficiency Virus Type 1 Envelope Glycoproteins.
Research progress on HIV-1 immune escape mechanisms.

[Role of the HIV-1 gp120 V1/V2 domains in the induction of neutralizing antibodies].
Therapeutic Vaccination Refocuses T-cell Responses Towards Conserved Regions of HIV-1 in Early Treated Individuals (BCN 01 study).
Neutralizing and other antiviral antibodies in HIV-1 infection and vaccination.
Phenotypic deficits in the HIV-1 envelope are associated with the maturation of a V2-directed broadly neutralizing antibody lineage.
The autophagy protein ATG9A promotes HIV-1 infectivity.
Structural Survey of Broadly Neutralizing Antibodies Targeting the HIV-1 Env Trimer Delineates Epitope Categories and Characteristics of Recognition.
Early low-titer neutralizing antibodies impede HIV-1 replication and select for virus escape.
Conserved Role of an N-Linked Glycan on the Surface Antigen of Human Immunodeficiency Virus Type 1 Modulating Virus Sensitivity to Broadly Neutralizing Antibodies against the Receptor and Coreceptor Binding Sites.
Feline immunodeficiency virus (FIV) neutralization: a review.
The Conformational States of the HIV-1 Envelope Glycoproteins.
Evasion from NK cell-mediated immune responses by HIV-1.
HIV-1 continues to replicate and evolve in patients with natural control of HIV infection.
Discovery of O-Linked Carbohydrate on HIV-1 Envelope and Its Role in Shielding against One Category of Broadly Neutralizing Antibodies.
Driving HIV-1 into a Vulnerable Corner by Taking Advantage of Viral Adaptation and Evolution.
Gp41-targeted antibodies restore infectivity of a fusion-deficient HIV-1 envelope glycoprotein.
Role of Viral Protein U (Vpu) in HIV-1 Infection and Pathogenesis.
HIV-1 blocks the signaling adaptor MAVS to evade antiviral host defense after sensing of abortive HIV-1 RNA by the host helicase DDX3.
Development of vaccination strategies that elicit broadly neutralizing antibodies against human immunodeficiency virus type 1 in both the mucosal and systemic immune compartments.
Human immunodeficiency virus type-1 (HIV-1) continues to evolve in presence of broadly neutralizing antibodies more than ten years after infection.
The HIV-1 envelope glycoproteins: fusogens, antigens, and immunogens.
HIV-1 mutates to evade IFITM1 restriction.
Stimulation of HIV-1 replication in immature dendritic cells in contact with primary CD4 T or B lymphocytes.
Subtle Longitudinal Alterations in Env Sequence Potentiate Differences in Sensitivity to Broadly Neutralizing Antibodies following Acute HIV-1 Subtype C Infection.
HIV-1 cell-to-cell transmission and broadly neutralizing antibodies.
A mechanistic understanding of allosteric immune escape pathways in the HIV-1 envelope glycoprotein.
Glycosylation of the core of the HIV-1 envelope subunit protein gp120 is not required for native trimer formation or viral infectivity.
4-phenylquinoline-8-amine induces HIV-1 reactivation and apoptosis in latently HIV-1 infected cells.
Survivors Remorse: antibody-mediated protection against HIV-1.
Short Communication: HIV-1 Infection Suppresses Circulating Viral Restriction microRNAs.
Migration of antigen-specific T cells away from CXCR4-binding human immunodeficiency virus type 1 gp120.
The V3 Loop of HIV-1 Env Determines Viral Susceptibility to IFITM3 Impairment of Viral Infectivity.
Unraveling the structural basis of HIV-1 neutralization.

Autophagy: an overlooked mechanism of HIV-1 pathogenesis and neuroAIDS?
Conformational dynamics of single HIV-1 envelope trimers on the surface of native virions.
Cell-to-cell spread of HIV-1 and evasion of neutralizing antibodies.
A Rare Mutation in an Infant-Derived HIV-1 Envelope Glycoprotein Alters Interprotomer Stability and Susceptibility to Broadly Neutralizing Antibodies Targeting the Trimer Apex.
High-multiplicity HIV-1 infection and neutralizing antibody evasion mediated by the macrophage-T cell virological synapse.
Neutralization Sensitivity of HIV-1 CRF07_BC From an Untreated Patient With a Focus on Evolution Over Time.
The antigenic structure of the HIV gp120 envelope glycoprotein.
Role of Escape Mutant-Specific T Cells in Suppression of HIV-1 Replication and Coevolution with HIV-1.
In vivo emergence of HIV-1 highly sensitive to neutralizing antibodies.
Neutralizing Antibody Responses and Evolution of the Viral Envelope in the Course of HIV-1 Korean Clade B Infection.
Evolution of B cell analysis and Env trimer redesign.
Murine monoclonal antibodies biologically active against the amino region of HIV-1 gp120: isolation and characterization.
Conformational Epitope-Specific Broadly Neutralizing Plasma Antibodies Obtained from an HIV-1 Clade C-Infected Elite Neutralizer Mediate Autologous Virus Escape through Mutations in the V1 Loop.
Polyclonal Broadly Neutralizing Antibody Activity Characterized by CD4 Binding Site and V3-Glycan Antibodies in a Subset of HIV-1 Virus Controllers.
SNAPPy: A snakemake pipeline for scalable HIV-1 subtyping by phylogenetic pairing.
HIV-1 induced nuclear factor I-B (NF-IB) expression negatively regulates HIV-1 replication through interaction with the long terminal repeat region.
Resistance of human immunodeficiency virus type 1 to neutralization by natural antisera occurs through single amino acid substitutions that cause changes in antibody binding at multiple sites.
Antibody-Induced Internalization of HIV-1 Env Proteins Limits Surface Expression of the Closed Conformation of Env.
Removal of a single N-linked glycan in human immunodeficiency virus type 1 gp120 results in an enhanced ability to induce neutralizing antibody responses.
Viral escape from neutralizing antibodies in early subtype A HIV-1 infection drives an increase in autologous neutralization breadth.
Design and characterization of a germ-line targeting soluble, native-like, trimeric HIV-1 Env lacking key glycans from the V1V2-loop.
The remarkable frequency of human immunodeficiency virus type 1 genetic recombination.
Association of mutations in V3/C3 domain with enhanced sensitivity of HIV-1 clade C primary envelopes to autologous broadly neutralizing plasma antibodies.
Envelope gene evolution and HIV-1 neuropathogenesis.
Solid-phase proteoliposomes containing human immunodeficiency virus envelope glycoproteins.
Antigenic characterization of the human immunodeficiency virus (HIV-1) envelope glycoprotein precursor incorporated into nanodiscs.
Stoichiometry of envelope glycoprotein trimers in the entry of human immunodeficiency virus type 1.
T-cell evasion and invasion during HIV-1 infection: The role of HIV-1 Tat protein.
Aiming to induce broadly reactive neutralizing antibody responses with HIV-1 vaccine candidates.
Recognition of a highly conserved region of human immunodeficiency virus type 1 gp120 by an HLA-Cw4-restricted cytotoxic T-lymphocyte clone.

Structure and immune recognition of trimeric pre-fusion HIV-1 Env.
Fusion proteins of HIV-1 envelope glycoprotein gp120 with CD4-induced antibodies showed enhanced binding to CD4 and CD4 binding site antibodies.
Limited Evidence for a Relationship between HIV-1 Glycan Shield Features in Early Infection and the Development of Neutralization Breadth.
The presence of the Trim5alpha escape mutation H87Q in the capsid of late stage HIV-1 variants is preceded by a prolonged asymptomatic infection phase.
Immunoescape of HIV-1 in Env-EL9 CD8 + T cell response restricted by HLA-B*14:02 in a Non progressor who lost twenty-seven years of HIV-1 control.
HIV-1 gp120 as a therapeutic target: navigating a moving labyrinth.
Structure and Recognition of a Novel HIV-1 gp120-gp41 Interface Antibody that Caused MPER Exposure through Viral Escape.
Non-synonymous Substitutions in HIV-1 GAG Are Frequent in Epitopes Outside the Functionally Conserved Regions and Associated With Subtype Differences.
Functional Anatomy of the Trimer Apex Reveals Key Hydrophobic Constraints That Maintain the HIV-1 Envelope Spike in a Closed State.
HIV-1, lipid rafts, and antibodies to liposomes: implications for anti-viral-neutralizing antibodies.
Straightforward selection of broadly neutralizing single-domain antibodies targeting the conserved CD4 and coreceptor binding sites of HIV-1 gp120.
Structure of Simian Immunodeficiency Virus Envelope Spikes Bound with CD4 and Monoclonal Antibody 36D5.
Mechanisms and Factors That Drive Extensive Human Immunodeficiency Virus Type-1 Hypervariability: An Overview.
Broadly Neutralizing Antibodies Against HIV: New Insights to Inform Vaccine Design.
Control of HIV-1 Replication by CD8(+) T Cells Specific for Two Novel Pol Protective Epitopes in HIV-1 Subtype A/E Infection.
Complement and HIV-1 infection/HIV-associated neurocognitive disorders.
Greater diversity of HIV-1 quasispecies in HIV-infected individuals with active tuberculosis.
HIV-1 fitness cost associated with escape from the VRC01 class of CD4 binding site neutralizing antibodies.
The Interplay of HIV-1 and Macrophages in Viral Persistence.
HIV-1 mutants that escape the cytotoxic T-lymphocytes are defective in viral DNA integration.
Viral escape from HIV-1 neutralizing antibodies drives increased plasma neutralization breadth through sequential recognition of multiple epitopes and immunotypes.
High-mannose-specific deglycosylation of HIV-1 gp120 induced by resistance to cyanovirin-N and the impact on antibody neutralization.
Chemical Cross-Linking Stabilizes Native-Like HIV-1 Envelope Glycoprotein Trimer Antigens.
Reduced evolutionary rates in HIV-1 reveal extensive latency periods among replicating lineages.
Neutralization function affected by single amino acid replacement in the HIV-1 antibody targets.
Candidate vaccine sequences to represent intra- and inter-clade HIV-1 variation.
CD4 binding site broadly neutralizing antibody selection of HIV-1 escape mutants.
HIV-1 evolution in gag and env is highly correlated but exhibits different relationships with viral load and the immune response.
Vpu-Mediated Counteraction of Tetherin Is a Major Determinant of HIV-1 Interferon Resistance.
Predicting in vivo escape dynamics of HIV-1 from a broadly neutralizing antibody.
Structure dynamics of HIV-1 Env trimers on native virions engaged with living T cells.
Rational modifications of HIV-1 envelope glycoproteins for immunogen design.



Roles of HIV-1 capsid in viral replication and immune evasion.
A Single Substitution in gp41 Modulates the Neutralization Profile of SHIV during In Vivo Adaptation.
A Highly Unusual V1 Region of Env in an Elite Controller of HIV Infection.
Human immunodeficiency virus-1 inhibition of immunoamphisomes in dendritic cells impairs early innate and adaptive immune responses.
Maturation-induced cloaking of neutralization epitopes on HIV-1 particles.
V4 region of the HIV-1 envelope gene mediates immune escape and may not promote the development of broadly neutralizing antibodies.
Structural basis of immune evasion at the site of CD4 attachment on HIV-1 gp120.
Glycosylation Benchmark Profile for HIV-1 Envelope Glycoprotein Production Based on Eleven Env Trimers.
A multi-scale mathematical modeling framework to investigate anti-viral therapeutic opportunities in targeting HIV-1 accessory proteins.
HIV-1 subverts the complement system in semen to enhance viral transmission.
DNA vaccine molecular adjuvants SP-D-BAFF and SP-D-APRIL enhance anti-gp120 immune response and increase HIV-1 neutralizing antibody titers.
Endocytosed HIV-1 Envelope Glycoprotein Traffics to Rab14(+) Late Endosomes and Lysosomes to Regulate Surface Levels in T-Cell Lines.
Neutralization sensitivity of HIV-1 subtype B' clinical isolates from former plasma donors in China.
The selection of low envelope glycoprotein reactivity to soluble CD4 and cold during simian-human immunodeficiency virus infection of rhesus macaques.
Development of Antibodies with Broad Neutralization Specificities against HIV-1 after Long Term SHIV Infection in Macaques.
The Oncolytic Virus MG1 Targets and Eliminates Cells Latently Infected With HIV-1: Implications for an HIV Cure.
HIV-1 adaptation to HLA: a window into virus-host immune interactions.
Identifying possible sites for antibody neutralization escape: Implications for unique functional properties of the C-terminal tail of Human Immunodeficiency Virus Type 1 gp41.
An Antigenic Atlas of HIV-1 Escape from Broadly Neutralizing Antibodies Distinguishes Functional and Structural Epitopes.
Unusual Fusion Proteins of HIV-1.
CD169-mediated trafficking of HIV to plasma membrane invaginations in dendritic cells attenuates efficacy of anti-gp120 broadly neutralizing antibodies.
Comprehensive Mapping of HIV-1 Escape from a Broadly Neutralizing Antibody.
Vertical T cell immunodominance and epitope entropy determine HIV-1 escape.
Dense Array of Spikes on HIV-1 Virion Particles.
The rationale for immunotherapy in HIV-1 infection.
Immunological tolerance as a barrier to protective HIV humoral immunity.
Probability of N332 glycan occupancy on HIV-1 gp120 modulates sensitivity to broadly neutralizing antibodies.
Mechanisms for Cell-to-Cell Transmission of HIV-1.
Rapid HIV-1 Disease Progression in Individuals Infected with a Virus Adapted to Its Host Population.
Primary HIV-1 infection: a review of clinical manifestations, immunologic and virologic changes.
Epitope-vaccine as a new strategy against HIV-1 mutation.
Emerging trends of HIV epidemiology in Asia.

Diverse pathways of escape from all well-characterized VRC01-class broadly neutralizing HIV-1 antibodies.
Neutralizing antibodies and control of HIV: moves and countermoves.
Antibody B cell responses in HIV-1 infection.
An in vitro system to model the establishment and reactivation of HIV-1 latency.
APOBEC3G contributes to HIV-1 variation through sublethal mutagenesis.
Human and rhesus APOBEC3D, APOBEC3F, APOBEC3G, and APOBEC3H demonstrate a conserved capacity to restrict Vif-deficient HIV-1.
Using HIV-1 sequence variability to explore virus biology.
Plasma membrane signaling in HIV-1 infection.
Dendritic Cell Response to HIV-1 Is Controlled by Differentiation Programs in the Cells and Strain-Specific Properties of the Virus.
Selective expansion of HIV-1 envelope glycoprotein-specific B cell subsets recognizing distinct structural elements following immunization.
Germline-like predecessors of broadly neutralizing antibodies lack measurable binding to HIV-1 envelope glycoproteins: implications for evasion of immune responses and design of vaccine immunogens.
Infection and depletion of CD4+ group-1 innate lymphoid cells by HIV-1 via type-I interferon pathway.
Glycan Microheterogeneity at the PGT135 Antibody Recognition Site on HIV-1 gp120 Reveals a Molecular Mechanism for Neutralization Resistance.
Molecular architecture of the uncleaved HIV-1 envelope glycoprotein trimer.
HIV-1 Envelope Mimicry of Host Enzyme Kynureninase Does Not Disrupt Tryptophan Metabolism.
The HIV Env Glycoprotein Conformational States on Cells and Viruses.
Genetic diversity of the highly variable V1 region interferes with Human Immunodeficiency Virus type 1 envelope functionality.
HR-2 mutations in human immunodeficiency virus type 1 gp41 restore fusion kinetics delayed by HR-1 mutations that cause clinical resistance to enfuvirtide.
Immune-mediated attenuation of HIV-1.
HIV-1 gp120 activates the STAT3/interleukin-6 axis in primary human monocyte-derived dendritic cells.
Dual Role of HIV-1 Envelope Signal Peptide in Immune Evasion.
Single-molecule imaging of HIV-1 envelope glycoprotein dynamics and Gag lattice association exposes determinants responsible for virus incorporation.
Human Three Prime Repair Exonuclease 1 Promotes HIV-1 Integration by Preferentially Degrading Unprocessed Viral DNA.
HLA-II-Associated HIV-1 Adaptation Decreases CD4(+) T-Cell Responses in HIV-1 Vaccine Recipients.
How does the humoral response to HIV-2 infection differ from HIV-1 and can this explain the distinct natural history of infection with these two human retroviruses?
The large extracellular loop of CD63 interacts with gp41 of HIV-1 and is essential for establishing the virological synapse.
Research Progress of HIV-1 Nef Inhibitors.
Subtype-specific conservation of isoleucine 309 in the envelope V3 domain is linked to immune evasion in subtype C HIV-1 infection.
HIV-1 resistance to neutralizing antibodies: Determination of antibody concentrations leading to escape mutant evolution.
Catch Me If You Can--The Race Between HIV and Neutralizing Antibodies.
Glycan Positioning Impacts HIV-1 Env Glycan-Shield Density, Function, and Recognition by Antibodies.

A comprehensive analysis of the naturally occurring polymorphisms in HIV-1 Vpr: potential impact on CTL epitopes.
Unliganded HIV-1 gp120 core structures assume the CD4-bound conformation with regulation by quaternary interactions and variable loops.
Loss of Nef-mediated CD3 down-regulation in the HIV-1 lineage increases viral infectivity and spread.
Short communication: HIV type 1 subtype C variants transmitted through the bottleneck of breastfeeding are sensitive to new generation broadly neutralizing antibodies directed against quaternary and CD4-binding site epitopes.
Disruption of helix-capping residues 671 and 674 reveals a role in HIV-1 entry for a specialized hinge segment of the membrane proximal external region of gp41.
Neutralization resistance of virological synapse-mediated HIV-1 Infection is regulated by the gp41 cytoplasmic tail.
4E10-resistant variants in a human immunodeficiency virus type 1 subtype C-infected individual with an anti-membrane-proximal external region-neutralizing antibody response.
HIV-1 Nef CAWLEAQ motif: a regulator of monocytes invasion through ENO1 modulation.
Robust growth of human immunodeficiency virus type 1 (HIV-1).
Neutralizing antibodies directed against the V3 loop select for different escape variants in a virus with mutated reverse transcriptase (M184V) than in wild-type human immunodeficiency virus type 1.
Dissecting the dynamics of HIV-1 protein sequence diversity.
Viral competition assay to assess the role of HIV-1 proteins in immune evasion.
The HIV-1 latent reservoir is largely sensitive to circulating T cells.
[Structural mechanism of immune evasion of HIV-1 gp120 by genomic, computational, and experimental science].
Limited neutralizing antibody specificities drive neutralization escape in early HIV-1 subtype C infection.
HIV-1 Gag and Vpr impair the inflammasome activation and contribute to the establishment of chronic infection in human primary macrophages.
Efficient capture of antibody neutralized HIV-1 by cells expressing DC-SIGN and transfer to CD4+ T lymphocytes.
Human domain antibodies to conserved sterically restricted regions on gp120 as exceptionally potent cross-reactive HIV-1 neutralizers.
Identification and characterization of a new cross-reactive human immunodeficiency virus type 1-neutralizing human monoclonal antibody.
Structures of HIV-1 gp120 envelope glycoproteins from laboratory-adapted and primary isolates.
The HIV-1 envelope transmembrane domain binds TLR2 through a distinct dimerization motif and inhibits TLR2-mediated responses.
Insights into the molecular mechanism underlying CD4-dependency and neutralization sensitivity of HIV-1: a comparative molecular dynamics study on gp120s from isolates with different phenotypes.
Glycans in HIV-1 vaccine design - engaging the shield.
Macrophage internal HIV-1 is protected from neutralizing antibodies.
Glycosylation site-specific analysis of clade C HIV-1 envelope proteins.
Structure of an HIV gp120 envelope glycoprotein in complex with the CD4 receptor and a neutralizing human antibody.
Identification of Human Immunodeficiency Virus Type-1 Subtypes by Heteroduplex Mobility Assay.
Chinks in the armor of the HIV-1 Envelope glycan shield: Implications for immune escape from anti-glycan broadly neutralizing antibodies.

HIV-1-specific CD4+ T-cell responses are not associated with significant viral epitope variation in persons with persistent plasma viremia.
Cryo-electron tomographic structure of an immunodeficiency virus envelope complex in situ.
Highly conserved HIV-1 gp120 glycans proximal to CD4-binding region affect viral infectivity and neutralizing antibody induction.
Detection of Human Immunodeficiency Virus Type 1 (HIV-1) A/AE Circulating Recombinant Form (CRF) in India: Possible Implications.
RNA interference targeted to the conserved dimerization initiation site (DIS) of HIV-1 restricts virus escape mutation.
Brugia malayi Antigen (BmA) Inhibits HIV-1 Trans-Infection but Neither BmA nor ES-62 Alter HIV-1 Infectivity of DC Induced CD4+ Th-Cells.
Trapping the HIV-1 V3 loop in a helical conformation enables broad neutralization.
Drug-Based Lead Discovery: The Novel Ablative Antiretroviral Profile of Deferiprone in HIV-1-Infected Cells and in HIV-Infected Treatment-Naive Subjects of a Double-Blind, Placebo-Controlled, Randomized Exploratory Trial.
Envelope glycoproteins sampling states 2/3 are susceptible to ADCC by sera from HIV-1-infected individuals.
Complex T-cell receptor repertoire dynamics underlie the CD8+ T-cell response to HIV-1.
Emergence of gp120 V3 variants confers neutralization resistance in an R5 simian-human immunodeficiency virus-infected macaque elite neutralizer that targets the N332 glycan of the human immunodeficiency virus type 1 envelope glycoprotein.
[Viral latency of HIV-1].
Human immunodeficiency virus type 1 gp120 envelope characteristics associated with disease progression differ in family members infected with genetically similar viruses.
Natural killer cells in HIV-1 infection: a double-edged sword.
Envelope-constrained neutralization-sensitive HIV-1 after heterosexual transmission.
Effects of CD4 Binding on Conformational Dynamics, Molecular Motions, and Thermodynamics of HIV-1 gp120.
Protease cleavage sites in HIV-1 gp120 recognized by antigen processing enzymes are conserved and located at receptor binding sites.
HIV envelope tail truncation confers resistance to SERINC5 restriction.
The Role of the BCL-2 Family of Proteins in HIV-1 Pathogenesis and Persistence.
Cellular HIV-1 inhibition by truncated old world primate APOBEC3A proteins lacking a complete deaminase domain.
CpG methylation controls reactivation of HIV from latency.
Humanized mouse models of HIV-1 latency.
Population-Level Immune-Mediated Adaptation in HIV-1 Polymerase during the North American Epidemic.
Deciphering human immunodeficiency virus type 1 transmission and early envelope diversification by single-genome amplification and sequencing.
Immunogenetic surveillance of HIV/AIDS.
HIV gp120: double lock strategy foils host defences.
In silico Analysis of HIV-1 Env-gp120 Reveals Structural Bases for Viral Adaptation in Growth-Restrictive Cells.
Cooperation between Strain-Specific and Broadly Neutralizing Responses Limited Viral Escape and Prolonged the Exposure of the Broadly Neutralizing Epitope.

The carbohydrate at asparagine 386 on HIV-1 gp120 is not essential for protein folding and function but is involved in immune evasion.
Sieve analysis in HIV-1 vaccine efficacy trials.
HIV-1 recombination: an experimental assay and a phylogenetic approach.
Detection of orientation-specific anti-gp120 antibodies by a new N-glycanase protection assay.
The spreading of HIV-1 infection in the human organism is caused by fractalkine trafficking of the infected lymphocytes--a review, hypothesis and implications for treatment.
Partial escape of HIV-1 from cytotoxic T lymphocytes during chronic infection.
The effect of CD4 receptor downregulation and its downstream signaling molecules on HIV-1 latency.
Schistosoma mansoni soluble egg antigen (SEA) and recombinant Omega-1 modulate induced CD4+ T-lymphocyte responses and HIV-1 infection in vitro.
The recombinant immunogen with high-density epitopes of ELDKWA and ELDEWA induced antibodies recognizing both epitopes on HIV-1 gp41.
Human immunodeficiency virus type 1 Vpr impairs dendritic cell maturation and T-cell activation: implications for viral immune escape.
Potential for immune-driven viral polymorphisms to compromise antiretroviral-based preexposure prophylaxis for prevention of HIV-1 infection.
Effective Cytotoxic T Lymphocyte Targeting of Persistent HIV-1 during Antiretroviral Therapy Requires Priming of Naive CD8+ T Cells.
Tyrosine-sulfated V2 peptides inhibit HIV-1 infection via coreceptor mimicry.
Broadly neutralizing antibodies: What is needed to move from a rare event in HIV-1 infection to vaccine efficacy?
BST-2 Expression Modulates Small CD4-Mimetic Sensitization of HIV-1-Infected Cells to Antibody-Dependent Cellular Cytotoxicity.
Plasma-Derived HIV-1 Virions Contain Considerable Levels of Defective Genomes.
Phospholipases: at the crossroads of the immune system and the pathogenesis of HIV-1 infection.
Relationship between functional profile of HIV-1 specific CD8 T cells and epitope variability with the selection of escape mutants in acute HIV-1 infection.
Different in vivo effects of HIV-1 immunodominant epitope-specific cytotoxic T lymphocytes on selection of escape mutant viruses.
Antigenicity and predefined specificities of the multi-epitope vaccine in candidate consisting of neutralizing epitope and mutated epitopes suggested a new way against HIV-1 mutation.
HIV-1 mediated immune pathogenesis: spotlight on the role of viral protein R (Vpr).
HIV-1 envelope subregion length variation during disease progression.
Experimental approaches to the study of HIV-1 latency.
Immunological strategies to target HIV persistence.
Effects of neutralizing antibodies on escape from CD8+ T-cell responses in HIV-1 infection.
Human Immunodeficiency Virus-1 Impairs IFN-Alpha Production Induced by TLR-7 Agonist in Plasmacytoid Dendritic Cells.
SAMHD1 restricts HIV-1 infection in resting CD4(+) T cells.
Envelope glycans of immunodeficiency virions are almost entirely oligomannose antigens.
CR4 Signaling Contributes to a DC-Driven Enhanced Immune Response Against Complement-Opsonized HIV-1.
Rare HIV-1 transmitted/founder lineages identified by deep viral sequencing contribute to rapid shifts in dominant quasispecies during acute and early infection.
HIV-1 Vpu Promotes Phagocytosis of Infected CD4(+) T Cells by Macrophages through Downregulation of CD47.

Structure of HIV-1 gp120 V1/V2 domain with broadly neutralizing antibody PG9.
Recurrent signature patterns in HIV-1 B clade envelope glycoproteins associated with either early or chronic infections.
An In Vitro System to Model the Establishment and Reactivation of HIV-1 Latency in Primary Human CD4+ T Cells.
Yeast-elicited cross-reactive antibodies to HIV Env glycans efficiently neutralize virions expressing exclusively high-mannose N-linked glycans.
Immune escape mutations selected by neutralizing antibodies in natural HIV-1 infection can alter coreceptor usage repertoire of the transmitted/founder virus.
Immune responses to HIV Gp120 that facilitate viral escape.
HIV-1 Protease as DNA Immunogen against Drug Resistance in HIV-1 Infection: DNA Immunization with Drug Resistant HIV-1 Protease Protects Mice from Challenge with Protease-Expressing Cells.
HIV evolution in early infection: selection pressures, patterns of insertion and deletion, and the impact of APOBEC.
Therapeutic vaccination for chronic infectious diseases: lessons from HIV-1.
A limited group of class I histone deacetylases acts to repress human immunodeficiency virus type 1 expression.
Large-Scale Arrayed Analysis of Protein Degradation Reveals Cellular Targets for HIV-1 Vpu.
Molecular characterization of HIV-1 subtype C gp-120 regions potentially involved in virus adaptive mechanisms.
Innate Invariant NKT Cell Recognition of HIV-1-Infected Dendritic Cells Is an Early Detection Mechanism Targeted by Viral Immune Evasion.
p21 Restricts HIV-1 in Monocyte-Derived Dendritic Cells through the Reduction of Deoxynucleoside Triphosphate Biosynthesis and Regulation of SAMHD1 Antiviral Activity.
The highly polymorphic cyclophilin A-binding loop in HIV-1 capsid modulates viral resistance to MxB.
Mucosal transmissibility, disease induction and coreceptor switching of R5 SHIVSF162P3N molecular clones in rhesus macaques.
The fitness landscape of HIV-1 gag: advanced modeling approaches and validation of model predictions by in vitro testing.
Broadly Neutralizing Antibodies to HIV and Their Role in Vaccine Design.
CD4 binding determinant mimicry for HIV vaccine design.
Architecture and regulation of the HIV-1 assembly and holding compartment in macrophages.
Generation of multivalent genome-wide T cell responses in HLA-A*0201 transgenic mice by an HIV-1 expression library immunization (ELI) vaccine.
Sequential evolution and escape from neutralization of simian immunodeficiency virus SIVsmE660 clones in rhesus macaques.
Structure of HIV-1 gp120 with gp41-interactive region reveals layered envelope architecture and basis of conformational mobility.
Interactions between HIV-1 and the cell-autonomous innate immune system.
Activities of transmitted/founder and chronic clade B HIV-1 Vpu and a C-terminal polymorphism specifically affecting virion release.
Dendritic cells infected with vpr-positive human immunodeficiency virus type 1 induce CD8+ T-cell apoptosis via upregulation of tumor necrosis factor alpha.
Common HIV-1 peptide variants mediate differential binding of KIR3DL1 to HLA-Bw4 molecules.
HIV-1 adaptation to NK-cell-mediated immune pressure.

A glycoconjugate antigen based on the recognition motif of a broadly neutralizing human immunodeficiency virus antibody, 2G12, is immunogenic but elicits antibodies unable to bind to the self glycans of gp120.
Remodeling of the Host Cell Plasma Membrane by HIV-1 Nef and Vpu: A Strategy to Ensure Viral Fitness and Persistence.
Virus-Host Gene Interactions Define HIV-1 Disease Progression.
Functional constraints on HIV-1 capsid: their impacts on the viral immune escape potency.
A de novo approach to inferring within-host fitness effects during untreated HIV-1 infection.
HIV-1 Hijacking of Host ATPases and GTPases That Control Protein Trafficking.
Host KIR/HLA-C Genotypes Determine HIV-Mediated Changes of the NK Cell Repertoire and Are Associated With Vpu Sequence Variations Impacting Downmodulation of HLA-C.
Turning up the volume on mutational pressure: is more of a good thing always better? (A case study of HIV-1 Vif and APOBEC3).
Human immunodeficiency virus-1 core: The Trojan horse in virus-host interaction.
Latent HIV-1 reservoirs in children: considerations for therapy.
[Virological characteristics of HIV].
A small set of succinct signature patterns distinguishes Chinese and non-Chinese HIV-1 genomes.
Frequent toggling between alternative amino acids is driven by selection in HIV-1.
Escape and compensation from early HLA-B57-mediated cytotoxic T-lymphocyte pressure on human immunodeficiency virus type 1 Gag alter capsid interactions with cyclophilin A.
T cell independent secondary antibody responses to the envelope protein of simian immunodeficiency virus.
Identification of putative, stable binding regions through flexibility analysis of HIV-1 gp120.
Defining HIV-1 Envelope N-Glycan Microdomains through Site-Specific Heterogeneity Profiles.
Immature dendritic cell-derived exosomes can mediate HIV-1 trans infection.
Mechanisms of escape from the PGT128 family of anti-HIV broadly neutralizing antibodies.
Study on molecular mechanisms of CD4 dependency and independency of HIV-1 gp120.
Solution structure, conformational dynamics, and CD4-induced activation in full-length, glycosylated, monomeric HIV gp120.
CRISPR-Cas9 Mediated Exonic Disruption for HIV-1 Elimination.
Pharmacologic Inhibition of Nedd8 Activation Enzyme Exposes CD4-Induced Epitopes within Env on Cells Expressing HIV-1.
HIV-1 conserved-element vaccines: relationship between sequence conservation and replicative capacity.
Immune escape and tropism of HIV.
Specific phospholipid recognition by human immunodeficiency virus type-1 neutralizing anti-gp41 2F5 antibody.
Visualization of the HIV-1 Env glycan shield across scales.
A multivalent HIV-vaccine: development of a plasmid DNA for the expression of HIV envelope glycoproteins with hypervariable V3-loop domains.
Cell-to-cell HIV-1 spread and its implications for immune evasion.
Detection of distinct human immunodeficiency virus type 1 circulating recombinant forms in northeast Brazil.
Enhancement of immunogenicity of an HIV Env DNA vaccine by mutation of the Tyr-based endocytosis motif in the cytoplasmic domain.
Global post-translational modification profiling of HIV-1-infected cells reveals mechanisms of host cellular pathway remodeling.

The B cell response is redundant and highly focused on V1V2 during early subtype C infection in a Zambian seroconverter.
Tracking coreceptor switch of the transmitted/founder HIV-1 identifies co-evolution of HIV-1 antigenicity, coreceptor usage and CD4 subset targeting.
Rapid escape from preserved cross-reactive neutralizing humoral immunity without loss of viral fitness in HIV-1-infected progressors and long-term nonprogressors.
Release of Soluble Ligands for the Activating NKG2D Receptor: One More Immune Evasion Strategy Evolved by HIV-1 ?
Evolution of HIV-1 within untreated individuals and at the population scale in Uganda.
Reliable reconstruction of HIV-1 whole genome haplotypes reveals clonal interference and genetic hitchhiking among immune escape variants.
Comparative glycoprofiling of HIV gp120 immunogens by capillary electrophoresis and MALDI mass spectrometry.
Viral Apoptosis Evasion via the MAPK Pathway by Use of a Host Long Noncoding RNA.
Potent anti-viral activity of a trispecific HIV neutralizing antibody in SHIV-infected monkeys.
Trimeric HIV-1-Env Structures Define Glycan Shields from Clades A, B, and G.
Extensive HIV-1 intra-host recombination is common in tissues with abnormal histopathology.
Antibodies Targeting the Envelope of HIV-1.
HIV Env conserved element DNA vaccine alters immunodominance in macaques.
A strongly selected mutation in the HIV-1 genome is independent of T cell responses and neutralizing antibodies.
Removal of N-linked glycosylation sites in the V1 region of simian immunodeficiency virus gp120 results in redirection of B-cell responses to V3.
Protective efficacy of centralized and polyvalent envelope immunogens in an attenuated equine lentivirus vaccine.
Design of peptide-based inhibitors for human immunodeficiency virus type 1 strains resistant to T-20.
Cooperation of B cell lineages in induction of HIV-1-broadly neutralizing antibodies.
CRISPR/Cas9-Derived Mutations Both Inhibit HIV-1 Replication and Accelerate Viral Escape.
The Clinical Implications of Reduced Viral Fitness.
HIV-1 Vpr hijacks EDD-DYRK2-DDB1(DCAF1) to disrupt centrosome homeostasis.
Variable epitope libraries: new vaccine immunogens capable of inducing broad human immunodeficiency virus type 1-neutralizing antibody response.
Design and construction of a recombinant epitope-peptide gene as a universal epitope-vaccine strategy.
HIV-1 subtype C Envelope function becomes less sensitive to N-glycosylation deletion during disease progression.
Molecular mechanisms of HIV entry.
Human immunodeficiency virus type-1 (HIV-1) evades antibody-dependent phagocytosis.
Allosteric modulation of the HIV-1 gp120-gp41 association site by adjacent gp120 variable region 1 (V1) N-glycans linked to neutralization sensitivity.
Envelope variation as a primary determinant of lentiviral vaccine efficacy.
Endogenous origins of HIV-1 G-to-A hypermutation and restriction in the nonpermissive T cell line CEM2n.
Cryo-EM structures of prefusion SIV envelope trimer.
Cell-to-Cell Spread of HIV and Viral Pathogenesis.
HIV-1 interacts with human endogenous retrovirus K (HML-2) envelopes derived from human primary lymphocytes.



Glycodendrimers prevent HIV transmission via DC-SIGN on dendritic cells.
New insights into the evolutionary rate of HIV-1 at the within-host and epidemiological levels.
Dynamics of viremia in primary HIV-1 infection in Africans: insights from analyses of host and viral correlates.
HIV-1-infected monocyte-derived dendritic cells do not undergo maturation but can elicit IL-10 production and T cell regulation.
Guanylate binding protein 5: Impairing virion infectivity by targeting retroviral envelope glycoproteins.
Macrophage bridging conduit trafficking of HIV-1 through the endoplasmic reticulum and Golgi network.
Novel Compound Inhibitors of HIV-1(NL4-3) Vpu.
A sensitive real-time PCR based assay to estimate the impact of amino acid substitutions on the competitive replication fitness of human immunodeficiency virus type 1 in cell culture.
Efficient BST2 antagonism by Vpu is critical for early HIV-1 dissemination in humanized mice.
Domain Organization of Lentiviral and Betaretroviral Surface Envelope Glycoproteins Modeled with AlphaFold.
The impact of HIV-1 within-host evolution on transmission dynamics.
Temporal Patterns and Drug Resistance in CSF Viral Escape Among ART-Experienced HIV-1 Infected Adults.
ADCC-mediating non-neutralizing antibodies can exert immune pressure in early HIV-1 infection.
A G1-like state allows HIV-1 to bypass SAMHD1 restriction in macrophages.
Human Immunodeficiency Virus Type 1 Vpr Mediates Degradation of APC1, a Scaffolding Component of the Anaphase-Promoting Complex/Cyclosome.
Constraints on HIV-1 diversity from protein structure.
Structural basis of antagonism of human APOBEC3F by HIV-1 Vif.
Modulation of HIV-1 infectivity and cyclophilin A-dependence by Gag sequence and target cell type.
HIV-1 Nef inhibits ASK1-dependent death signalling providing a potential mechanism for protecting the infected host cell.
HIV-1 Nef and Vpu Interfere with L-Selectin (CD62L) Cell Surface Expression To Inhibit Adhesion and Signaling in Infected CD4+ T Lymphocytes.
Change in tropism upon immune escape by human immunodeficiency virus.
Significantly longer envelope V2 loops are characteristic of heterosexually transmitted subtype B HIV-1 in Trinidad.
Impact of HIV-1 infection pathways on susceptibility to antiviral drugs and on virus spread.
Identifying recombination hot spots in the HIV-1 genome.
HIV-1 Vpu mediated downregulation of CD155 requires alanine residues 10, 14 and 18 of the transmembrane domain.
Current Trends of HIV Recombination Worldwide.
The appearance of escape variants in vivo does not account for the failure of recombinant envelope vaccines to protect against simian immunodeficiency virus.
Neutralization of HIV by antibodies.
IL-10-producing B cells are induced early in HIV-1 infection and suppress HIV-1-specific T cell responses.
Prediction and validation of HIV-1 gp41 ecto-transmembrane domain post-fusion trimeric structure using molecular modeling.
Preadaptation of Simian Immunodeficiency Virus SIVsmm Facilitated Env-Mediated Counteraction of Human Tetherin by Human Immunodeficiency Virus Type 2.

Identification of two unique naturally occurring Vpr sequence polymorphisms associated with clinical parameters in HIV-1 chronic infection.
NLRX1 Helps HIV Avoid a STING Operation.
HIV-1 Vpu restricts Fc-mediated effector functions in vivo.
Partial compartmentalisation of HIV-1 subtype C between lymph nodes, peripheral blood mononuclear cells and plasma.
Evolution of neutralizing antibody response against HIV type 1 virions and pseudovirions in multicenter AIDS cohort study participants.
Neisseria gonorrhoeae effectively blocks HIV-1 replication by eliciting a potent TLR9-dependent interferon- $\alpha$ response from plasmacytoid dendritic cells.
Neutralization and infectivity characteristics of envelope glycoproteins from human immunodeficiency virus type 1 infected donors whose sera exhibit broadly cross-reactive neutralizing activity.
HIV-1 subtype C envelope characteristics associated with divergent rates of chronic disease progression.
Structural features of HIV envelope defined by antibody escape mutant analysis.
HAART-persistent HIV-1 latent reservoirs: their origin, mechanisms of stability and potential strategies for eradication.
SIV infection duration largely determines broadening of neutralizing antibody response in macaques.
Lost in translation: implications of HIV-1 codon usage for immune escape and drug resistance.
Genetic divergence of HIV-1 B subtype in Italy over the years 2003-2016 and impact on CTL escape prevalence.
The HIV-1-containing macrophage compartment: a perfect cellular niche?
HIV-1 Vpu Downmodulates ICAM-1 Expression, Resulting in Decreased Killing of Infected CD4(+) T Cells by NK Cells.
Can non-lytic CD8+ T cells drive HIV-1 escape?
The interferon-inducible MxB protein inhibits HIV-1 infection.
The Genetic Diversity of HIV-1 Quasispecies Within Primary Infected Individuals.
Molecular basis for the binding promiscuity of an anti-p24 (HIV-1) monoclonal antibody.
HIV-1 Establishes a Sanctuary Site in the Testis by Permeating the BTB Through Changes in Cytoskeletal Organization.
Virological synapse-mediated spread of human immunodeficiency virus type 1 between T cells is sensitive to entry inhibition.
The degree of HIV-1 amino acid variability is strictly related to different disease progression rates.
Immune evasion activities of accessory proteins Vpu, Nef and Vif are conserved in acute and chronic HIV-1 infection.
Glycoform analysis of recombinant and human immunodeficiency virus envelope protein gp120 via higher energy collisional dissociation and spectral-aligning strategy.
Frequent intrapatient recombination between human immunodeficiency virus type 1 R5 and X4 envelopes: implications for coreceptor switch.
Human immunodeficiency virus type 1 Vpu protein interacts with CD74 and modulates major histocompatibility complex class II presentation.
HIV-1-induced impairment of dendritic cell cross talk with $\gamma\delta$ T lymphocytes.
Identification of the V1 region as a linear neutralizing epitope of the simian immunodeficiency virus SIVmac envelope glycoprotein.
The dynamics of HIV-1 adaptation in early infection.

Envelope glycoprotein determinants of increased fusogenicity in a pathogenic simian-human immunodeficiency virus (SHIV-KB9) passaged in vivo.
Accurately measuring recombination between closely related HIV-1 genomes.
Partial rescue of V1V2 mutant infectivity by HIV-1 cell-cell transmission supports the domain's exceptional capacity for sequence variation.
Dynamic of CSF and serum biomarkers in HIV-1 subtype C encephalitis with CNS genetic compartmentalization-case study.
Dynamics of HIV-1 coinfection in different susceptible target cell populations during cell-free infection.
Dual regulation of L-selectin (CD62L) by HIV-1: Enhanced expression by Vpr in contrast with cell-surface down-modulation by Nef and Vpu.
Immuno-epidemiological modeling of HIV-1 predicts high heritability of the set-point virus load, while selection for CTL escape dominates virulence evolution.
Deciphering gp120 sequence variation and structural dynamics in HIV neutralization phenotype by molecular dynamics simulations and graph machine learning.
Infection of specific dendritic cells by CCR5-tropic human immunodeficiency virus type 1 promotes cell-mediated transmission of virus resistant to broadly neutralizing antibodies.
Antibody-Dependent Cellular Cytotoxicity (ADCC)-Mediating Antibodies Constrain Neutralizing Antibody Escape Pathway.
HIV-1 replication in CD4(+) T cells exploits the down-regulation of antiviral NEAT1 long non-coding RNAs following T cell activation.
Insights into amprenavir resistance in E35D HIV-1 protease mutation from molecular dynamics and binding free-energy calculations.
HIV-1 Vpu blocks recycling and biosynthetic transport of the intrinsic immunity factor CD317/tetherin to overcome the virion release restriction.
APOBEC3G induces a hypermutation gradient: purifying selection at multiple steps during HIV-1 replication results in levels of G-to-A mutations that are high in DNA, intermediate in cellular viral RNA, and low in virion RNA.
Rev variation during persistent lentivirus infection.
Technical advance. Measurement of iNKT cell responses at the single-cell level against rare HIV-1-infected dendritic cells in a mixed culture.
Balance between transmitted HLA preadapted and nonassociated polymorphisms is a major determinant of HIV-1 disease progression.
HIV-1 Vpu affects the anterograde transport and the glycosylation pattern of NTB-A.
The Transmembrane Domain of HIV-1 gp41 Inhibits T-Cell Activation by Targeting Multiple T-Cell Receptor Complex Components through Its GxxxG Motif.
Probing intrinsic dynamics and conformational transition of HIV gp120 by molecular dynamics simulation.
The transmembrane proteins contribute to immunodeficiencies induced by HIV-1 and other retroviruses.
Cutting edge: An antibody recognizing ancestral endogenous virus glycoproteins mediates antibody-dependent cellular cytotoxicity on HIV-1-infected cells.
Detailed analysis of CD4+ Th responses to envelope and Gag proteins of simian immunodeficiency virus reveals an exclusion of broadly reactive Th epitopes from the glycosylated regions of envelope.
Implications of HIV RNA structure for recombination, speciation, and the neutralism-selectionism controversy.
A comparative analysis of viral matrix proteins using disorder predictors.

Simian immunodeficiency virus infection evades vaccine-elicited antibody responses to V2 region.
A combined EM and proteomic analysis places HIV-1 Vpu at the crossroads of retromer and ESCRT complexes: PTPN23 is a Vpu-cofactor.
Expansion of epitope cross-reactivity by anti-idiotypic modulation of the primary humoral response.
Difficulties in eliciting broadly neutralizing anti-HIV antibodies are not explained by cardiolipin autoreactivity.
HIV accessory proteins versus host restriction factors.
Shell disorder, immune evasion and transmission behaviors among human and animal retroviruses.
Decreased infectivity of a neutralization-resistant equine infectious anemia virus variant can be overcome by efficient cell-to-cell spread.
Mutational resilience of antiviral restriction favors primate TRIM5 $\alpha$ in host-virus evolutionary arms races.
T-cell differentiation factor CBF- $\beta$ regulates HIV-1 Vif-mediated evasion of host restriction.
Retroviral infection in vivo requires an immune escape virulence factor encrypted in the envelope protein of oncoretroviruses.
Variation in simian immunodeficiency virus env V1 region in simian AIDS-associated lymphoma.
HIV-2/SIV viral protein X counteracts HUSH repressor complex.
Recombinant feline leukemia virus (FeLV) variants establish a limited infection with altered cell tropism in specific-pathogen-free cats in the absence of FeLV subgroup A helper virus.
Cerebrospinal fluid western Blot profiles in the evolution of HIV-1 pediatric encephalopathy.
Sequence and Functional Variation in the HIV-1 Rev Regulatory Axis.
HIV-1 Vif N-terminal motif is required for recruitment of CUL5 to suppress APOBEC3.
The N-Terminal Region of HIV-1 Tat Protein Binds CD127 in Human CD8 T Cells to Target the Receptor for Down Regulation Through Tat's Basic Region.
Molecular and biological characterization of equine infectious anemia virus Rev.

Table A2-37, Cluster 36

Cluster 36 focuses on metabolic impacts on the immune system, especially on the contribution of metabolic dysfunction to the immunosuppressive tumor microenvironment (256)
Metabolic Hallmarks of Tumor and Immune Cells in the Tumor Microenvironment.
Immunometabolism: A new target for improving cancer immunotherapy.
Emerging Landscapes of Tumor Immunity and Metabolism.
Emerging role of metabolic reprogramming in tumor immune evasion and immunotherapy.
Metabolic reprogramming of immune cells in pancreatic cancer progression.
Lipid Metabolism and Tumor Antigen Presentation.
Immune metabolism in PD-1 blockade-based cancer immunotherapy.
Metabolic Plasticity in Cancers-Distinct Role of Glycolytic Enzymes GPI, LDHs or Membrane Transporters MCTs.
Genetics of enzymatic dysfunctions in metabolic disorders and cancer.
Lipid Metabolism Regulation Based on Nanotechnology for Enhancement of Tumor Immunity.
Harnessing Metabolic Reprogramming to Improve Cancer Immunotherapy.
Metabolism of Dendritic Cells in Tumor Microenvironment: For Immunotherapy.
Targeting immune-onco-metabolism for precision cancer therapy.
Metabolic influence on the differentiation of suppressive myeloid cells in cancer.
[Effect of abnormal lipid metabolism on immune microenvironment in tumors].

CD147-mediated reprogrammed glycolytic metabolism potentially induces immune escape in the tumor microenvironment (Review).
Immunometabolism in cancer at a glance.
Lactate Contribution to the Tumor Microenvironment: Mechanisms, Effects on Immune Cells and Therapeutic Relevance.
Nanodrug regulates lactic acid metabolism to reprogram the immunosuppressive tumor microenvironment for enhanced cancer immunotherapy.
Tumor Microenvironment: A Metabolic Player that Shapes the Immune Response.
[Metabolic Competition in Tumor Microenvironment].
Tumor-associated macrophages: new insights on their metabolic regulation and their influence in cancer immunotherapy.
Targeting Metabolism to Control Immune Responses in Cancer and Improve Checkpoint Blockade Immunotherapy.
Unveiling the veil of lactate in tumor-associated macrophages: a successful strategy for immunometabolic therapy.
Novel strategy for oncogenic alteration-induced lipid metabolism reprogramming in pancreatic cancer.
Aerobic glycolysis and high level of lactate in cancer metabolism and microenvironment.
Cell Intrinsic and Systemic Metabolism in Tumor Immunity and Immunotherapy.
Tumor Microenvironment-Derived Metabolites: A Guide to Find New Metabolic Therapeutic Targets and Biomarkers.
Linking Immuno-evasion and Metabolic Reprogramming in B-Cell-Derived Lymphomas.
Mevalonate metabolism in cancer.
Immunometabolism in the tumor microenvironment and its related research progress.
Lipid Metabolic Regulatory Crosstalk Between Cancer Cells and Tumor-Associated Macrophages.
Targeting Tumor Metabolism: A New Challenge to Improve Immunotherapy.
Adapt and conquer: Metabolic flexibility in cancer growth, invasion and evasion.
Targeted Glucose or Glutamine Metabolic Therapy Combined With PD-1/PD-L1 Checkpoint Blockade Immunotherapy for the Treatment of Tumors - Mechanisms and Strategies.
Lactate in the Regulation of Tumor Microenvironment and Therapeutic Approaches.
Lactate in the tumor microenvironment: A rising star for targeted tumor therapy.
Metabolic Reprogramming and Immune Evasion in Nasopharyngeal Carcinoma.
Fueling the Revolution: Targeting Metabolism to Enhance Immunotherapy.
Cancer Metabolism: Fueling More than Just Growth.
Control of the Antitumor Immune Response by Cancer Metabolism.
Contradictory roles of lipid metabolism in immune response within the tumor microenvironment.
Hypoxia, cancer metabolism and the therapeutic benefit of targeting lactate/H(+) symporters.
Lactate-Lactylation Hands between Metabolic Reprogramming and Immunosuppression.
A novel strategy to fuel cancer immunotherapy: targeting glucose metabolism to remodel the tumor microenvironment.
Influence of the Metabolism on Myeloid Cell Functions in Cancers: Clinical Perspectives.
Lipid metabolic features of T cells in the Tumor Microenvironment.
Sculpting tumor microenvironment with immune system: from immunometabolism to immunoediting.
Aberrant Lipid Metabolism in Cancer: Current Status and Emerging Therapeutic Perspectives.
Even Cancer Cells Watch Their Cholesterol!

Tumor cell metabolism: cancer's Achilles' heel.
Metabolic reprogramming in the tumour microenvironment: a hallmark shared by cancer cells and T lymphocytes.
Targeting L-Lactate Metabolism to Overcome Resistance to Immune Therapy of Melanoma and Other Tumor Entities.
Tumor metabolism as modulator of immune response and tumor progression.
Glutamine blockade induces divergent metabolic programs to overcome tumor immune evasion.
Lipid metabolism reprogramming in tumor-associated macrophages and implications for therapy.
Metabolic programming and immune suppression in the tumor microenvironment.
Reexamining cancer metabolism: lactate production for carcinogenesis could be the purpose and explanation of the Warburg Effect.
Bioenergetic and Metabolic Adaptation in Tumor Progression and Metastasis.
Metabolic reprogramming and epigenetic modifications on the path to cancer.
Mitochondrial immune regulation and anti-tumor immunotherapy strategies targeting mitochondria.
Metabolic Symbiosis and Immunomodulation: How Tumor Cell-Derived Lactate May Disturb Innate and Adaptive Immune Responses.
Tumor-Derived Lactate Creates a Favorable Niche for Tumor via Supplying Energy Source for Tumor and Modulating the Tumor Microenvironment.
Dynamic links between mechanical forces and metabolism shape the tumor milieu.
Reprogramming of glutamine metabolism and its impact on immune response in the tumor microenvironment.
Cancer metabolism within tumor microenvironments.
Lipid Metabolism in Tumor-Associated Macrophages.
Linking tumor glycolysis and immune evasion in cancer: Emerging concepts and therapeutic opportunities.
Revisited Metabolic Control and Reprogramming Cancers by Means of the Warburg Effect in Tumor Cells.
The Role of Metabolism in Tumor Immune Evasion: Novel Approaches to Improve Immunotherapy.
The Development of Single-Cell Metabolism and Its Role in Studying Cancer Emergent Properties.
Overview: Lipid Metabolism in the Tumor Microenvironment.
Reprogramming lipid metabolism as potential strategy for hematological malignancy therapy.
Plant-derived extracts and metabolic modulation in leukemia: a promising approach to overcome treatment resistance.
The roles of metabolic profiles and intracellular signaling pathways of tumor microenvironment cells in angiogenesis of solid tumors.
Microenvironmental Metabolism Regulates Antitumor Immunity.
Functional Genomics In Vivo Reveal Metabolic Dependencies of Pancreatic Cancer Cells.
Effects of tumor metabolic microenvironment on regulatory T cells.
Resveratrol reduces lactate production and modifies the ovarian cancer immune microenvironment.
Metabolic regulation of suppressive myeloid cells in cancer.
Amino acid metabolism in tumor: New shine in the fog?
Lipidomic-Based Advances in Diagnosis and Modulation of Immune Response to Cancer.
Metabolic Reprogramming and Vulnerabilities in Cancer.
Lactate as a signaling molecule: Journey from dead end product of glycolysis to tumor survival.
T Cells and Cancer: How Metabolism Shapes Immunity.
From tumor cell metabolism to tumor immune escape.

The Importance of Cellular Metabolic Pathways in Pathogenesis and Selective Treatments of Hematological Malignancies.
Cancer Metabolism Historical Perspectives: A Chronicle of Controversies and Consensus.
Metabolic adaptations of cancer in extreme tumor microenvironments.
TGFβ-induced metabolic reprogramming during epithelial-to-mesenchymal transition in cancer.
Lactate: a metabolic key player in cancer.
Cancer Cell Metabolism Bolsters Immunotherapy Resistance by Promoting an Immunosuppressive Tumor Microenvironment.
Natural Compounds as Metabolic Modulators of the Tumor Microenvironment.
Role of dendritic cell metabolic reprogramming in tumor immune evasion.
Reprogramming of Metabolism in Kidney Cancer.
Targeting Glucose Metabolism Enzymes in Cancer Treatment: Current and Emerging Strategies.
Early immune pressure makes tumors metabolically stronger.
Lactate as Key Metabolite in Prostate Cancer Progression: What Are the Clinical Implications?
Metabolic communication in tumors: a new layer of immunoregulation for immune evasion.
Pancreatic cancer and fibrosis: Targeting metabolic reprogramming and crosstalk of cancer-associated fibroblasts in the tumor microenvironment.
The crosstalking of lactate-Histone lactylation and tumor.
Chemical metabolic inhibitors for the treatment of blood-borne cancers.
Orchestral role of lipid metabolic reprogramming in T-cell malignancy.
Lactate trafficking inhibition restores sensitivity to proteasome inhibitors and orchestrates immuno-microenvironment in multiple myeloma.
The emerging "hallmarks" of metabolic reprogramming and immune evasion: distinct or linked?
Amino acid metabolism: challenges and opportunities for the therapeutic treatment of leukemia and lymphoma.
Immune Evasion in Tumor's Own Sweet Way.
Enhancing the Efficacy of Glutamine Metabolism Inhibitors in Cancer Therapy.
Nucleotide metabolism: a pan-cancer metabolic dependency.
Immunoediting instructs tumor metabolic reprogramming to support immune evasion.
NAD-Biosynthetic and Consuming Enzymes as Central Players of Metabolic Regulation of Innate and Adaptive Immune Responses in Cancer.
Crosstalk between Macrophages, T Cells, and Iron Metabolism in Tumor Microenvironment.
Sirpigenastat (DRP-104) Induces Antitumor Efficacy through Direct, Broad Antagonism of Glutamine Metabolism and Stimulation of the Innate and Adaptive Immune Systems.
Major fundamental factors hindering immune system in defense against tumor cells: The link between insufficiency of innate immune responses, metabolism, and neurotransmitters with effector immune cells disability.
Metabolic Plasticity of Melanoma Cells and Their Crosstalk With Tumor Microenvironment.
Metabolic barriers in non-small cell lung cancer with LKB1 and/or KEAP1 mutations for immunotherapeutic strategies.
Metabolism, LXR/LXR ligands, and tumor immune escape.
m6A-regulated tumor glycolysis: new advances in epigenetics and metabolism.
Cancer Cell Metabolism Reprogramming and Its Potential Implications on Therapy in Squamous Cell Carcinoma of the Head and Neck: A Review.
Lactate supports a metabolic-epigenetic link in macrophage polarization.
A Forgotten Corner in Cancer Immunotherapy: The Role of Lipids.

Metabolic Remodeling in Glioma Immune Microenvironment: Intercellular Interactions Distinct From Peripheral Tumors.
Metabolic shift underlies tumor progression and immune evasion in S-nitrosoglutathione reductase-deficient cancer.
Metabolism and senescence in the immune microenvironment of osteosarcoma: focus on new therapeutic strategies.
Lactate Exposure Promotes Immunosuppressive Phenotypes in Innate Immune Cells.
Regulating tumor glycometabolism and the immune microenvironment by inhibiting lactate dehydrogenase with platinum(IV) complexes.
The Induction of a Permissive Environment to Promote T Cell Immune Evasion in Acute Myeloid Leukemia: The Metabolic Perspective.
Metabolic regulation of the cancer-immunity cycle.
Mitochondrial Metabolism: A New Dimension of Personalized Oncology.
DNMT1 mediates metabolic reprogramming induced by Epstein-Barr virus latent membrane protein 1 and reversed by grifolin in nasopharyngeal carcinoma.
Lactate/GPR81 signaling and proton motive force in cancer: Role in angiogenesis, immune escape, nutrition, and Warburg phenomenon.
'Warburg effect' controls tumor growth, bacterial, viral infections and immunity - Genetic deconstruction and therapeutic perspectives.
The role of polyamine metabolism in remodeling immune responses and blocking therapy within the tumor immune microenvironment.
TGF- $\beta$ in Cancer: Metabolic Driver of the Tolerogenic Crosstalk in the Tumor Microenvironment.
Cancer Bioenergetics and Tumor Microenvironments-Enhancing Chemotherapeutics and Targeting Resistant Niches through Nanosystems.
Cancer immunoediting: A process driven by metabolic competition as a predator-prey-shared resource type model.
Mutant p53 Gain-of-Function: Role in Cancer Development, Progression, and Therapeutic Approaches.
Immunometabolism at the Nexus of Cancer Therapeutic Efficacy and Resistance.
Lactate modulates microglia polarization via IGFBP6 expression and remodels tumor microenvironment in glioblastoma.
The tortuous path of lactate shuttle discovery: From cinders and boards to the lab and ICU.
Mitochondrial metabolism: Inducer or therapeutic target in tumor immune-resistance?
Mitochondrial and Metabolic Pathways Regulate Nuclear Gene Expression to Control Differentiation, Stem Cell Function, and Immune Response in Leukemia.
Unveiling tumor immune evasion mechanisms: abnormal expression of transporters on immune cells in the tumor microenvironment.
Wnt Signaling in Cancer Metabolism and Immunity.
Unravelling the role of obesity and lipids during tumor progression.
The immunological Warburg effect: Can a metabolic-tumor-stroma score (MeTS) guide cancer immunotherapy?
Dendritic Cell Metabolism and Function in Tumors.
Cancer-generated lactic acid: a regulatory, immunosuppressive metabolite?
Lactate Induces Pro-tumor Reprogramming in Intratumoral Plasmacytoid Dendritic Cells.
Cytokines and metabolic factors regulate tumoricidal T-cell function during cancer immunotherapy.
Targeting nucleotide metabolism: a promising approach to enhance cancer immunotherapy.
UVA, metabolism and melanoma: UVA makes melanoma hungry for metastasis.



UCP2 as a Potential Biomarker for Adjunctive Metabolic Therapies in Tumor Management.
Fighting in a wasteland: deleterious metabolites and antitumor immunity.
Monitoring Lactate Dynamics in Individual Macrophages with a Genetically Encoded Probe.
Out of the cycle: Impact of cell cycle aberrations on cancer metabolism and metastasis.
Targeting lipid metabolism in the treatment of ovarian cancer.
Rewired Metabolism of Amino Acids and Its Roles in Glioma Pathology.
Targeting glutamine metabolism with photodynamic immunotherapy for metastatic tumor eradication.
Targeting Dietary and Microbial Tryptophan-Indole Metabolism as Therapeutic Approaches to Colon Cancer.
The biological implications of Yin Yang 1 in the hallmarks of cancer.
Metastasis is promoted by a bioenergetic switch: new targets for progressive renal cell cancer.
Potential Utility of Synthetic D-Lactate Polymers in Skin Cancer.
Nitric Oxide: The Forgotten Child of Tumor Metabolism.
Recent insights into the implications of metabolism in plasmacytoid dendritic cell innate functions: Potential ways to control these functions.
Mitochondrial reprogramming via ATP5H loss promotes multimodal cancer therapy resistance.
The Double-Edge Sword of Autophagy in Cancer: From Tumor Suppression to Pro-tumor Activity.
Immunomodulatory roles of nitric oxide in cancer: tumor microenvironment says "NO" to antitumor immune response.
MicroRNA-124 Enhances T Cells Functions by Manipulating the Lactic Acid Metabolism of Tumor Cells.
Targeting Oncometabolites in Peritoneal Cancers: Preclinical Insights and Therapeutic Strategies.
Cholesterol Metabolism in Cancer and Cell Death.
Immune-regulated IDO1-dependent tryptophan metabolism is source of one-carbon units for pancreatic cancer and stellate cells.
Lactate enhances motility of tumor cells and inhibits monocyte migration and cytokine release.
Crosstalk between oxidative phosphorylation and immune escape in cancer: a new concept of therapeutic targets selection.
Altered Iron Metabolism and Impact in Cancer Biology, Metastasis, and Immunology.
Tumor aerobic glycolysis confers immune evasion through modulating sensitivity to T cell-mediated bystander killing via TNF- $\alpha$ .
Remodeling "cold" tumor immune microenvironment via epigenetic-based therapy using targeted liposomes with in situ formed albumin corona.
The role and its mechanism of intermittent fasting in tumors: friend or foe?
Transient systemic autophagy ablation irreversibly inhibits lung tumor cell metabolism and promotes T-cell mediated tumor killing.
Transient Systemic Autophagy Inhibition Is Selectively and Irreversibly Deleterious to Lung Cancer.
Notch1/TAZ axis promotes aerobic glycolysis and immune escape in lung cancer.
Iron metabolism: State of the art in hypoxic cancer cell biology.
Metabolic factors contribute to T-cell inhibition in the ovarian cancer ascites.
Exploiting Mitochondrial Vulnerabilities to Trigger Apoptosis Selectively in Cancer Cells.
Glucose-Restricted Diet Regulates the Tumor Immune Microenvironment and Prevents Tumor Growth in Lung Adenocarcinoma.
Glutamine promotes escape from therapy-induced senescence in tumor cells.
lncRNA HITT Inhibits Lactate Production by Repressing PKM2 Oligomerization to Reduce Tumor Growth and Macrophage Polarization.

Mechanism of PKM2 affecting cancer immunity and metabolism in Tumor Microenvironment.
Metabolic inhibitor screening identifies dihydrofolate reductase as an inducer of the tumor immune escape mediator CD24.
Tumor acidity: From hallmark of cancer to target of treatment.
Targeted Anti-Mitochondrial Therapy: The Future of Oncology.
Tryptophan and its metabolites in normal physiology and cancer etiology.
Synthesis and anticancer activity of new coumarin-3-carboxylic acid derivatives as potential lactate transport inhibitors.
Why Warburg Works: Lactate Controls Immune Evasion through GPR81.
Therapeutic targeting of cellular stress responses in cancer.
Lactate and TGF- $\beta$ antagonistically regulate inflammasome activation in the tumor microenvironment.
Cholesterol metabolism: At the cross road between cancer cells and immune environment.
Autophagy Regulates Stress Responses, Metabolism, and Anticancer Immunity.
How nearby nutrients shape tumor growth.
Peroxisomes and cancer: The role of a metabolic specialist in a disease of aberrant metabolism.
Development of a novel lactate dehydrogenase A inhibitor with potent antitumor activity and immune activation.
[The cancer paradigm in pulmonary arterial hypertension: towards anti-remodeling therapies targeting metabolic dysfunction?].
To betray or to fight? The dual identity of the mitochondria in cancer.
The immune regulation in cancer by the amino acid metabolizing enzymes ARG and IDO.
Melanoma and the problem of malignancy.
Dual role of pseudogene TMEM198B in promoting lipid metabolism and immune escape of glioma cells.
Identification of a lipid metabolism-related gene for cancer immunotherapy.
Autophagy in cancer cell remodeling and quality control.
Paracrine Wnt5a- $\beta$ -Catenin Signaling Triggers a Metabolic Program that Drives Dendritic Cell Tolerization.
Hyaluronan in the Tumor Microenvironment.
cROSSing the Line: Between Beneficial and Harmful Effects of Reactive Oxygen Species in B-Cell Malignancies.
Lactic Acid and an Acidic Tumor Microenvironment suppress Anticancer Immunity.
Lactate dehydrogenase 5 (LDH-5) expression in endometrial cancer relates to the activated VEGF/VEGFR2(KDR) pathway and prognosis.
Extracellular-signal-regulated kinase 5 modulates the antioxidant response by transcriptionally controlling Sirtuin 1 expression in leukemic cells.
Tumors exploit FTO-mediated regulation of glycolytic metabolism to evade immune surveillance.
PPAR $\alpha$ Inhibition Overcomes Tumor-Derived Exosomal Lipid-Induced Dendritic Cell Dysfunction.
Targeting Metabolic Bottlenecks in Lung Cancer.
Impact of context-dependent autophagy states on tumor progression.
Monocarboxylate transporter 4 (MCT4) is a high affinity transporter capable of exporting lactate in high-lactate microenvironments.
Indoleamine 2,3-dioxygenase 2 depletion suppresses tumor growth in a mouse model of Lewis lung carcinoma.
High fructose diet: A risk factor for immune system dysregulation.
High-density lipoproteins: A promising tool against cancer.

Kynurenic Acid: The Janus-Faced Role of an Immunomodulatory Tryptophan Metabolite and Its Link to Pathological Conditions.
GRP78 and next generation cancer hallmarks: An underexplored molecular target in cancer chemoprevention research.
Burkitt lymphoma and lactic acidosis: A case report and review of the literature.
Cancer: a de-repression of a default survival program common to all cells?: a life-history perspective on the nature of cancer.
Regulatory roles of copper metabolism and cuproptosis in human cancers.
Glucose-induced toxicity in insulin-producing pituitary cells that coexpress GLUT2 and glucokinase. Implications for metabolic engineering.
Cell cycle on the crossroad of tumorigenesis and cancer therapy.
Tryptophan in health and disease.
Concurrent imaging of vascularization and metabolism in a mouse model of paraganglioma under anti-angiogenic treatment.
Finding your niche: immune evasion in quiescent tumor reservoirs.
Targeting tumor-intrinsic hexosamine biosynthesis sensitizes pancreatic cancer to anti-PD1 therapy.
Targeting tumor endothelial hyperglycolysis enhances immunotherapy through remodeling tumor microenvironment.
Cancer Cachexia and Dysregulated Phosphate Metabolism: Insights from Mutant p53 and Mutant Klotho Mouse Models.
The Transcriptional and Epigenetic Landscape of Cancer Cell Lineage Plasticity.
Cholesterol: An important actor on the cancer immune scene.
Uncovering the interplay between pH receptors and immune cells: Potential drug targets (Review).
Revisiting the hallmarks of cancer: The role of hyaluronan.
O-GlcNAc in cancer: An Oncometabolism-fueled vicious cycle.
Fatty acid oxidation fuels glioblastoma radioresistance with CD47-mediated immune evasion.
The IL411 Enzyme: A New Player in the Immunosuppressive Tumor Microenvironment.
Biological Functions and Analytical Strategies of Sialic Acids in Tumor.
Chinese herbal medicine anticancer cocktail soup activates immune cells to kill colon cancer cells by regulating the gut microbiota-Th17 axis.
Lactate levels with glioblastoma multiforme.
Glucose regulated protein 78: a critical link between tumor microenvironment and cancer hallmarks.
The two faces of the Integrated Stress Response in cancer progression and therapeutic strategies.
Dietary spinach reshapes the gut microbiome in an Apc-mutant genetic background: mechanistic insights from integrated multi-omics.
Cyclic Derivatives of the Chemerin C-Terminus as Metabolically Stable Agonists at the Chemokine-like Receptor 1 for Cancer Treatment.
Methionine and Kynurenine Activate Oncogenic Kinases in Glioblastoma, and Methionine Deprivation Compromises Proliferation.
Bone marrow niches in haematological malignancies.
Efficient Reprogramming of Human Fibroblasts and Blood-Derived Endothelial Progenitor Cells Using Nonmodified RNA for Reprogramming and Immune Evasion.
Excessive intake of sugar: An accomplice of inflammation.
NAD <sup>+</sup> -metabolizing ecto-enzymes shape tumor-host interactions: the chronic lymphocytic leukemia model.
Roles and Modalities of Ectonucleotidases in Remodeling the Multiple Myeloma Niche.

MYCN Drives a Tumor Immunosuppressive Environment Which Impacts Survival in Neuroblastoma.

Table A2-38, Cluster 37

Cluster 36 focuses on metabolic impacts on the immune system, especially on the contribution of metabolic dysfunction to the immunosuppressive tumor microenvironment (256)
Metabolic Hallmarks of Tumor and Immune Cells in the Tumor Microenvironment.
Immunometabolism: A new target for improving cancer immunotherapy.
Emerging Landscapes of Tumor Immunity and Metabolism.
Emerging role of metabolic reprogramming in tumor immune evasion and immunotherapy.
Metabolic reprogramming of immune cells in pancreatic cancer progression.
Lipid Metabolism and Tumor Antigen Presentation.
Immune metabolism in PD-1 blockade-based cancer immunotherapy.
Metabolic Plasticity in Cancers-Distinct Role of Glycolytic Enzymes GPI, LDHs or Membrane Transporters MCTs.
Genetics of enzymatic dysfunctions in metabolic disorders and cancer.
Lipid Metabolism Regulation Based on Nanotechnology for Enhancement of Tumor Immunity.
Harnessing Metabolic Reprogramming to Improve Cancer Immunotherapy.
Metabolism of Dendritic Cells in Tumor Microenvironment: For Immunotherapy.
Targeting immune-onco-metabolism for precision cancer therapy.
Metabolic influence on the differentiation of suppressive myeloid cells in cancer.
[Effect of abnormal lipid metabolism on immune microenvironment in tumors].
CD147-mediated reprogrammed glycolytic metabolism potentially induces immune escape in the tumor microenvironment (Review).
Immunometabolism in cancer at a glance.
Lactate Contribution to the Tumor Microenvironment: Mechanisms, Effects on Immune Cells and Therapeutic Relevance.
Nanodrug regulates lactic acid metabolism to reprogram the immunosuppressive tumor microenvironment for enhanced cancer immunotherapy.
Tumor Microenvironment: A Metabolic Player that Shapes the Immune Response.
[Metabolic Competition in Tumor Microenvironment].
Tumor-associated macrophages: new insights on their metabolic regulation and their influence in cancer immunotherapy.
Targeting Metabolism to Control Immune Responses in Cancer and Improve Checkpoint Blockade Immunotherapy.
Unveiling the veil of lactate in tumor-associated macrophages: a successful strategy for immunometabolic therapy.
Novel strategy for oncogenic alteration-induced lipid metabolism reprogramming in pancreatic cancer.
Aerobic glycolysis and high level of lactate in cancer metabolism and microenvironment.
Cell Intrinsic and Systemic Metabolism in Tumor Immunity and Immunotherapy.
Tumor Microenvironment-Derived Metabolites: A Guide to Find New Metabolic Therapeutic Targets and Biomarkers.
Linking Immune evasion and Metabolic Reprogramming in B-Cell-Derived Lymphomas.
Mevalonate metabolism in cancer.
Immunometabolism in the tumor microenvironment and its related research progress.

Lipid Metabolic Regulatory Crosstalk Between Cancer Cells and Tumor-Associated Macrophages.
Targeting Tumor Metabolism: A New Challenge to Improve Immunotherapy.
Adapt and conquer: Metabolic flexibility in cancer growth, invasion and evasion.
Targeted Glucose or Glutamine Metabolic Therapy Combined With PD-1/PD-L1 Checkpoint Blockade Immunotherapy for the Treatment of Tumors - Mechanisms and Strategies.
Lactate in the Regulation of Tumor Microenvironment and Therapeutic Approaches.
Lactate in the tumor microenvironment: A rising star for targeted tumor therapy.
Metabolic Reprogramming and Immune Evasion in Nasopharyngeal Carcinoma.
Fueling the Revolution: Targeting Metabolism to Enhance Immunotherapy.
Cancer Metabolism: Fueling More than Just Growth.
Control of the Antitumor Immune Response by Cancer Metabolism.
Contradictory roles of lipid metabolism in immune response within the tumor microenvironment.
Hypoxia, cancer metabolism and the therapeutic benefit of targeting lactate/H(+) symporters.
Lactate-Lactylation Hands between Metabolic Reprogramming and Immunosuppression.
A novel strategy to fuel cancer immunotherapy: targeting glucose metabolism to remodel the tumor microenvironment.
Influence of the Metabolism on Myeloid Cell Functions in Cancers: Clinical Perspectives.
Lipid metabolic features of T cells in the Tumor Microenvironment.
Sculpting tumor microenvironment with immune system: from immunometabolism to immunoediting.
Aberrant Lipid Metabolism in Cancer: Current Status and Emerging Therapeutic Perspectives.
Even Cancer Cells Watch Their Cholesterol!
Tumor cell metabolism: cancer's Achilles' heel.
Metabolic reprogramming in the tumour microenvironment: a hallmark shared by cancer cells and T lymphocytes.
Targeting L-Lactate Metabolism to Overcome Resistance to Immune Therapy of Melanoma and Other Tumor Entities.
Tumor metabolism as modulator of immune response and tumor progression.
Glutamine blockade induces divergent metabolic programs to overcome tumor immune evasion.
Lipid metabolism reprogramming in tumor-associated macrophages and implications for therapy.
Metabolic programming and immune suppression in the tumor microenvironment.
Reexamining cancer metabolism: lactate production for carcinogenesis could be the purpose and explanation of the Warburg Effect.
Bioenergetic and Metabolic Adaptation in Tumor Progression and Metastasis.
Metabolic reprogramming and epigenetic modifications on the path to cancer.
Mitochondrial immune regulation and anti-tumor immunotherapy strategies targeting mitochondria.
Metabolic Symbiosis and Immunomodulation: How Tumor Cell-Derived Lactate May Disturb Innate and Adaptive Immune Responses.
Tumor-Derived Lactate Creates a Favorable Niche for Tumor via Supplying Energy Source for Tumor and Modulating the Tumor Microenvironment.
Dynamic links between mechanical forces and metabolism shape the tumor milieu.
Reprogramming of glutamine metabolism and its impact on immune response in the tumor microenvironment.
Cancer metabolism within tumor microenvironments.
Lipid Metabolism in Tumor-Associated Macrophages.

Linking tumor glycolysis and immune evasion in cancer: Emerging concepts and therapeutic opportunities.
Revisited Metabolic Control and Reprogramming Cancers by Means of the Warburg Effect in Tumor Cells.
The Role of Metabolism in Tumor Immune Evasion: Novel Approaches to Improve Immunotherapy.
The Development of Single-Cell Metabolism and Its Role in Studying Cancer Emergent Properties.
Overview: Lipid Metabolism in the Tumor Microenvironment.
Reprogramming lipid metabolism as potential strategy for hematological malignancy therapy.
Plant-derived extracts and metabolic modulation in leukemia: a promising approach to overcome treatment resistance.
The roles of metabolic profiles and intracellular signaling pathways of tumor microenvironment cells in angiogenesis of solid tumors.
Microenvironmental Metabolism Regulates Antitumor Immunity.
Functional Genomics In Vivo Reveal Metabolic Dependencies of Pancreatic Cancer Cells.
Effects of tumor metabolic microenvironment on regulatory T cells.
Resveratrol reduces lactate production and modifies the ovarian cancer immune microenvironment.
Metabolic regulation of suppressive myeloid cells in cancer.
Amino acid metabolism in tumor: New shine in the fog?
Lipidomic-Based Advances in Diagnosis and Modulation of Immune Response to Cancer.
Metabolic Reprogramming and Vulnerabilities in Cancer.
Lactate as a signaling molecule: Journey from dead end product of glycolysis to tumor survival.
T Cells and Cancer: How Metabolism Shapes Immunity.
From tumor cell metabolism to tumor immune escape.
The Importance of Cellular Metabolic Pathways in Pathogenesis and Selective Treatments of Hematological Malignancies.
Cancer Metabolism Historical Perspectives: A Chronicle of Controversies and Consensus.
Metabolic adaptations of cancer in extreme tumor microenvironments.
TGFβ-induced metabolic reprogramming during epithelial-to-mesenchymal transition in cancer.
Lactate: a metabolic key player in cancer.
Cancer Cell Metabolism Bolsters Immunotherapy Resistance by Promoting an Immunosuppressive Tumor Microenvironment.
Natural Compounds as Metabolic Modulators of the Tumor Microenvironment.
Role of dendritic cell metabolic reprogramming in tumor immune evasion.
Reprogramming of Metabolism in Kidney Cancer.
Targeting Glucose Metabolism Enzymes in Cancer Treatment: Current and Emerging Strategies.
Early immune pressure makes tumors metabolically stronger.
Lactate as Key Metabolite in Prostate Cancer Progression: What Are the Clinical Implications?
Metabolic communication in tumors: a new layer of immunoregulation for immune evasion.
Pancreatic cancer and fibrosis: Targeting metabolic reprogramming and crosstalk of cancer-associated fibroblasts in the tumor microenvironment.
The crosstalking of lactate-Histone lactylation and tumor.
Chemical metabolic inhibitors for the treatment of blood-borne cancers.
Orchestral role of lipid metabolic reprogramming in T-cell malignancy.
Lactate trafficking inhibition restores sensitivity to proteasome inhibitors and orchestrates immuno-microenvironment in multiple myeloma.
The emerging "hallmarks" of metabolic reprogramming and immune evasion: distinct or linked?

Amino acid metabolism: challenges and opportunities for the therapeutic treatment of leukemia and lymphoma.
Immune Evasion in Tumor's Own Sweet Way.
Enhancing the Efficacy of Glutamine Metabolism Inhibitors in Cancer Therapy.
Nucleotide metabolism: a pan-cancer metabolic dependency.
Immunoediting instructs tumor metabolic reprogramming to support immune evasion.
NAD-Biosynthetic and Consuming Enzymes as Central Players of Metabolic Regulation of Innate and Adaptive Immune Responses in Cancer.
Crosstalk between Macrophages, T Cells, and Iron Metabolism in Tumor Microenvironment.
Sirpiglenastat (DRP-104) Induces Antitumor Efficacy through Direct, Broad Antagonism of Glutamine Metabolism and Stimulation of the Innate and Adaptive Immune Systems.
Major fundamental factors hindering immune system in defense against tumor cells: The link between insufficiency of innate immune responses, metabolism, and neurotransmitters with effector immune cells disability.
Metabolic Plasticity of Melanoma Cells and Their Crosstalk With Tumor Microenvironment.
Metabolic barriers in non-small cell lung cancer with LKB1 and/or KEAP1 mutations for immunotherapeutic strategies.
Metabolism, LXR/LXR ligands, and tumor immune escape.
m6A-regulated tumor glycolysis: new advances in epigenetics and metabolism.
Cancer Cell Metabolism Reprogramming and Its Potential Implications on Therapy in Squamous Cell Carcinoma of the Head and Neck: A Review.
Lactate supports a metabolic-epigenetic link in macrophage polarization.
A Forgotten Corner in Cancer Immunotherapy: The Role of Lipids.
Metabolic Remodeling in Glioma Immune Microenvironment: Intercellular Interactions Distinct From Peripheral Tumors.
Metabolic shift underlies tumor progression and immune evasion in S-nitrosoglutathione reductase-deficient cancer.
Metabolism and senescence in the immune microenvironment of osteosarcoma: focus on new therapeutic strategies.
Lactate Exposure Promotes Immunosuppressive Phenotypes in Innate Immune Cells.
Regulating tumor glycometabolism and the immune microenvironment by inhibiting lactate dehydrogenase with platinum(IV) complexes.
The Induction of a Permissive Environment to Promote T Cell Immune Evasion in Acute Myeloid Leukemia: The Metabolic Perspective.
Metabolic regulation of the cancer-immunity cycle.
Mitochondrial Metabolism: A New Dimension of Personalized Oncology.
DNMT1 mediates metabolic reprogramming induced by Epstein-Barr virus latent membrane protein 1 and reversed by grifolin in nasopharyngeal carcinoma.
Lactate/GPR81 signaling and proton motive force in cancer: Role in angiogenesis, immune escape, nutrition, and Warburg phenomenon.
'Warburg effect' controls tumor growth, bacterial, viral infections and immunity - Genetic deconstruction and therapeutic perspectives.
The role of polyamine metabolism in remodeling immune responses and blocking therapy within the tumor immune microenvironment.
TGF- $\beta$ in Cancer: Metabolic Driver of the Tolerogenic Crosstalk in the Tumor Microenvironment.
Cancer Bioenergetics and Tumor Microenvironments-Enhancing Chemotherapeutics and Targeting Resistant Niches through Nanosystems.

Cancer immunoediting: A process driven by metabolic competition as a predator-prey-shared resource type model.
Mutant p53 Gain-of-Function: Role in Cancer Development, Progression, and Therapeutic Approaches.
Immunometabolism at the Nexus of Cancer Therapeutic Efficacy and Resistance.
Lactate modulates microglia polarization via IGFBP6 expression and remodels tumor microenvironment in glioblastoma.
The tortuous path of lactate shuttle discovery: From cinders and boards to the lab and ICU.
Mitochondrial metabolism: Inducer or therapeutic target in tumor immune-resistance?
Mitochondrial and Metabolic Pathways Regulate Nuclear Gene Expression to Control Differentiation, Stem Cell Function, and Immune Response in Leukemia.
Unveiling tumor immune evasion mechanisms: abnormal expression of transporters on immune cells in the tumor microenvironment.
Wnt Signaling in Cancer Metabolism and Immunity.
Unravelling the role of obesity and lipids during tumor progression.
The immunological Warburg effect: Can a metabolic-tumor-stroma score (MeTS) guide cancer immunotherapy?
Dendritic Cell Metabolism and Function in Tumors.
Cancer-generated lactic acid: a regulatory, immunosuppressive metabolite?
Lactate Induces Pro-tumor Reprogramming in Intratumoral Plasmacytoid Dendritic Cells.
Cytokines and metabolic factors regulate tumoricidal T-cell function during cancer immunotherapy.
Targeting nucleotide metabolism: a promising approach to enhance cancer immunotherapy.
UVA, metabolism and melanoma: UVA makes melanoma hungry for metastasis.
UCP2 as a Potential Biomarker for Adjunctive Metabolic Therapies in Tumor Management.
Fighting in a wasteland: deleterious metabolites and antitumor immunity.
Monitoring Lactate Dynamics in Individual Macrophages with a Genetically Encoded Probe.
Out of the cycle: Impact of cell cycle aberrations on cancer metabolism and metastasis.
Targeting lipid metabolism in the treatment of ovarian cancer.
Rewired Metabolism of Amino Acids and Its Roles in Glioma Pathology.
Targeting glutamine metabolism with photodynamic immunotherapy for metastatic tumor eradication.
Targeting Dietary and Microbial Tryptophan-Indole Metabolism as Therapeutic Approaches to Colon Cancer.
The biological implications of Yin Yang 1 in the hallmarks of cancer.
Metastasis is promoted by a bioenergetic switch: new targets for progressive renal cell cancer.
Potential Utility of Synthetic D-Lactate Polymers in Skin Cancer.
Nitric Oxide: The Forgotten Child of Tumor Metabolism.
Recent insights into the implications of metabolism in plasmacytoid dendritic cell innate functions: Potential ways to control these functions.
Mitochondrial reprogramming via ATP5H loss promotes multimodal cancer therapy resistance.
The Double-Edge Sword of Autophagy in Cancer: From Tumor Suppression to Pro-tumor Activity.
Immunomodulatory roles of nitric oxide in cancer: tumor microenvironment says "NO" to antitumor immune response.
MicroRNA-124 Enhances T Cells Functions by Manipulating the Lactic Acid Metabolism of Tumor Cells.
Targeting Oncometabolites in Peritoneal Cancers: Preclinical Insights and Therapeutic Strategies.
Cholesterol Metabolism in Cancer and Cell Death.



Immune-regulated IDO1-dependent tryptophan metabolism is source of one-carbon units for pancreatic cancer and stellate cells.
Lactate enhances motility of tumor cells and inhibits monocyte migration and cytokine release.
Crosstalk between oxidative phosphorylation and immune escape in cancer: a new concept of therapeutic targets selection.
Altered Iron Metabolism and Impact in Cancer Biology, Metastasis, and Immunology.
Tumor aerobic glycolysis confers immune evasion through modulating sensitivity to T cell-mediated bystander killing via TNF- $\alpha$ .
Remodeling "cold" tumor immune microenvironment via epigenetic-based therapy using targeted liposomes with in situ formed albumin corona.
The role and its mechanism of intermittent fasting in tumors: friend or foe?
Transient systemic autophagy ablation irreversibly inhibits lung tumor cell metabolism and promotes T-cell mediated tumor killing.
Transient Systemic Autophagy Inhibition Is Selectively and Irreversibly Deleterious to Lung Cancer.
Notch1/TAZ axis promotes aerobic glycolysis and immune escape in lung cancer.
Iron metabolism: State of the art in hypoxic cancer cell biology.
Metabolic factors contribute to T-cell inhibition in the ovarian cancer ascites.
Exploiting Mitochondrial Vulnerabilities to Trigger Apoptosis Selectively in Cancer Cells.
Glucose-Restricted Diet Regulates the Tumor Immune Microenvironment and Prevents Tumor Growth in Lung Adenocarcinoma.
Glutamine promotes escape from therapy-induced senescence in tumor cells.
lncRNA HITT Inhibits Lactate Production by Repressing PKM2 Oligomerization to Reduce Tumor Growth and Macrophage Polarization.
Mechanism of PKM2 affecting cancer immunity and metabolism in Tumor Microenvironment.
Metabolic inhibitor screening identifies dihydrofolate reductase as an inducer of the tumor immune escape mediator CD24.
Tumor acidity: From hallmark of cancer to target of treatment.
Targeted Anti-Mitochondrial Therapy: The Future of Oncology.
Tryptophan and its metabolites in normal physiology and cancer etiology.
Synthesis and anticancer activity of new coumarin-3-carboxylic acid derivatives as potential lactate transport inhibitors.
Why Warburg Works: Lactate Controls Immune Evasion through GPR81.
Therapeutic targeting of cellular stress responses in cancer.
Lactate and TGF- $\beta$ antagonistically regulate inflammasome activation in the tumor microenvironment.
Cholesterol metabolism: At the cross road between cancer cells and immune environment.
Autophagy Regulates Stress Responses, Metabolism, and Anticancer Immunity.
How nearby nutrients shape tumor growth.
Peroxisomes and cancer: The role of a metabolic specialist in a disease of aberrant metabolism.
Development of a novel lactate dehydrogenase A inhibitor with potent antitumor activity and immune activation.
[The cancer paradigm in pulmonary arterial hypertension: towards anti-remodeling therapies targeting metabolic dysfunction?].
To betray or to fight? The dual identity of the mitochondria in cancer.
The immune regulation in cancer by the amino acid metabolizing enzymes ARG and IDO.
Melanoma and the problem of malignancy.

Dual role of pseudogene TMEM198B in promoting lipid metabolism and immune escape of glioma cells.
Identification of a lipid metabolism-related gene for cancer immunotherapy.
Autophagy in cancer cell remodeling and quality control.
Paracrine Wnt5a- $\beta$ -Catenin Signaling Triggers a Metabolic Program that Drives Dendritic Cell Tolerization.
Hyaluronan in the Tumor Microenvironment.
cROSSing the Line: Between Beneficial and Harmful Effects of Reactive Oxygen Species in B-Cell Malignancies.
Lactic Acid and an Acidic Tumor Microenvironment suppress Anticancer Immunity.
Lactate dehydrogenase 5 (LDH-5) expression in endometrial cancer relates to the activated VEGF/VEGFR2(KDR) pathway and prognosis.
Extracellular-signal-regulated kinase 5 modulates the antioxidant response by transcriptionally controlling Sirtuin 1 expression in leukemic cells.
Tumors exploit FTO-mediated regulation of glycolytic metabolism to evade immune surveillance.
PPAR $\alpha$ Inhibition Overcomes Tumor-Derived Exosomal Lipid-Induced Dendritic Cell Dysfunction.
Targeting Metabolic Bottlenecks in Lung Cancer.
Impact of context-dependent autophagy states on tumor progression.
Monocarboxylate transporter 4 (MCT4) is a high affinity transporter capable of exporting lactate in high-lactate microenvironments.
Indoleamine 2,3-dioxygenase 2 depletion suppresses tumor growth in a mouse model of Lewis lung carcinoma.
High fructose diet: A risk factor for immune system dysregulation.
High-density lipoproteins: A promising tool against cancer.
Kynurenic Acid: The Janus-Faced Role of an Immunomodulatory Tryptophan Metabolite and Its Link to Pathological Conditions.
GRP78 and next generation cancer hallmarks: An underexplored molecular target in cancer chemoprevention research.
Burkitt lymphoma and lactic acidosis: A case report and review of the literature.
Cancer: a de-repression of a default survival program common to all cells?: a life-history perspective on the nature of cancer.
Regulatory roles of copper metabolism and cuproptosis in human cancers.
Glucose-induced toxicity in insulin-producing pituitary cells that coexpress GLUT2 and glucokinase. Implications for metabolic engineering.
Cell cycle on the crossroad of tumorigenesis and cancer therapy.
Tryptophan in health and disease.
Concurrent imaging of vascularization and metabolism in a mouse model of paraganglioma under anti-angiogenic treatment.
Finding your niche: immune evasion in quiescent tumor reservoirs.
Targeting tumor-intrinsic hexosamine biosynthesis sensitizes pancreatic cancer to anti-PD1 therapy.
Targeting tumor endothelial hyperglycolysis enhances immunotherapy through remodeling tumor microenvironment.
Cancer Cachexia and Dysregulated Phosphate Metabolism: Insights from Mutant p53 and Mutant Klotho Mouse Models.
The Transcriptional and Epigenetic Landscape of Cancer Cell Lineage Plasticity.
Cholesterol: An important actor on the cancer immune scene.

Uncovering the interplay between pH receptors and immune cells: Potential drug targets (Review).
Revisiting the hallmarks of cancer: The role of hyaluronan.
O-GlcNAc in cancer: An Oncometabolism-fueled vicious cycle.
Fatty acid oxidation fuels glioblastoma radioresistance with CD47-mediated immune evasion.
The IL411 Enzyme: A New Player in the Immunosuppressive Tumor Microenvironment.
Biological Functions and Analytical Strategies of Sialic Acids in Tumor.
Chinese herbal medicine anticancer cocktail soup activates immune cells to kill colon cancer cells by regulating the gut microbiota-Th17 axis.
Lactate levels with glioblastoma multiforme.
Glucose regulated protein 78: a critical link between tumor microenvironment and cancer hallmarks.
The two faces of the Integrated Stress Response in cancer progression and therapeutic strategies.
Dietary spinach reshapes the gut microbiome in an Apc-mutant genetic background: mechanistic insights from integrated multi-omics.
Cyclic Derivatives of the Chemerin C-Terminus as Metabolically Stable Agonists at the Chemokine-like Receptor 1 for Cancer Treatment.
Methionine and Kynurenine Activate Oncogenic Kinases in Glioblastoma, and Methionine Deprivation Compromises Proliferation.
Bone marrow niches in haematological malignancies.
Efficient Reprogramming of Human Fibroblasts and Blood-Derived Endothelial Progenitor Cells Using Nonmodified RNA for Reprogramming and Immune Evasion.
Excessive intake of sugar: An accomplice of inflammation.
NAD <sup>+</sup> -metabolizing ecto-enzymes shape tumor-host interactions: the chronic lymphocytic leukemia model.
Roles and Modalities of Ectonucleotidases in Remodeling the Multiple Myeloma Niche.
MYCN Drives a Tumor Immunosuppressive Environment Which Impacts Survival in Neuroblastoma.
Cluster 36 focuses on metabolic impacts on the immune system, especially on the contribution of metabolic dysfunction to the immunosuppressive tumor microenvironment (256)
Metabolic Hallmarks of Tumor and Immune Cells in the Tumor Microenvironment.
Immunometabolism: A new target for improving cancer immunotherapy.
Emerging Landscapes of Tumor Immunity and Metabolism.
Emerging role of metabolic reprogramming in tumor immune evasion and immunotherapy.
Metabolic reprogramming of immune cells in pancreatic cancer progression.
Lipid Metabolism and Tumor Antigen Presentation.
Immune metabolism in PD-1 blockade-based cancer immunotherapy.
Metabolic Plasticity in Cancers-Distinct Role of Glycolytic Enzymes GPI, LDHs or Membrane Transporters MCTs.
Genetics of enzymatic dysfunctions in metabolic disorders and cancer.
Lipid Metabolism Regulation Based on Nanotechnology for Enhancement of Tumor Immunity.
Harnessing Metabolic Reprogramming to Improve Cancer Immunotherapy.
Metabolism of Dendritic Cells in Tumor Microenvironment: For Immunotherapy.
Targeting immune-onco-metabolism for precision cancer therapy.
Metabolic influence on the differentiation of suppressive myeloid cells in cancer.
[Effect of abnormal lipid metabolism on immune microenvironment in tumors].
CD147-mediated reprogrammed glycolytic metabolism potentially induces immune escape in the tumor microenvironment (Review).
Immunometabolism in cancer at a glance.

Lactate Contribution to the Tumor Microenvironment: Mechanisms, Effects on Immune Cells and Therapeutic Relevance.
Nanodrug regulates lactic acid metabolism to reprogram the immunosuppressive tumor microenvironment for enhanced cancer immunotherapy.
Tumor Microenvironment: A Metabolic Player that Shapes the Immune Response.
[Metabolic Competition in Tumor Microenvironment].
Tumor-associated macrophages: new insights on their metabolic regulation and their influence in cancer immunotherapy.
Targeting Metabolism to Control Immune Responses in Cancer and Improve Checkpoint Blockade Immunotherapy.
Unveiling the veil of lactate in tumor-associated macrophages: a successful strategy for immunometabolic therapy.
Novel strategy for oncogenic alteration-induced lipid metabolism reprogramming in pancreatic cancer.
Aerobic glycolysis and high level of lactate in cancer metabolism and microenvironment.
Cell Intrinsic and Systemic Metabolism in Tumor Immunity and Immunotherapy.
Tumor Microenvironment-Derived Metabolites: A Guide to Find New Metabolic Therapeutic Targets and Biomarkers.
Linking Immune evasion and Metabolic Reprogramming in B-Cell-Derived Lymphomas.
Mevalonate metabolism in cancer.
Immunometabolism in the tumor microenvironment and its related research progress.
Lipid Metabolic Regulatory Crosstalk Between Cancer Cells and Tumor-Associated Macrophages.
Targeting Tumor Metabolism: A New Challenge to Improve Immunotherapy.
Adapt and conquer: Metabolic flexibility in cancer growth, invasion and evasion.
Targeted Glucose or Glutamine Metabolic Therapy Combined With PD-1/PD-L1 Checkpoint Blockade Immunotherapy for the Treatment of Tumors - Mechanisms and Strategies.
Lactate in the Regulation of Tumor Microenvironment and Therapeutic Approaches.
Lactate in the tumor microenvironment: A rising star for targeted tumor therapy.
Metabolic Reprogramming and Immune Evasion in Nasopharyngeal Carcinoma.
Fueling the Revolution: Targeting Metabolism to Enhance Immunotherapy.
Cancer Metabolism: Fueling More than Just Growth.
Control of the Antitumor Immune Response by Cancer Metabolism.
Contradictory roles of lipid metabolism in immune response within the tumor microenvironment.
Hypoxia, cancer metabolism and the therapeutic benefit of targeting lactate/H(+) symporters.
Lactate-Lactylation Hands between Metabolic Reprogramming and Immunosuppression.
A novel strategy to fuel cancer immunotherapy: targeting glucose metabolism to remodel the tumor microenvironment.
Influence of the Metabolism on Myeloid Cell Functions in Cancers: Clinical Perspectives.
Lipid metabolic features of T cells in the Tumor Microenvironment.
Sculpting tumor microenvironment with immune system: from immunometabolism to immunoediting.
Aberrant Lipid Metabolism in Cancer: Current Status and Emerging Therapeutic Perspectives.
Even Cancer Cells Watch Their Cholesterol!
Tumor cell metabolism: cancer's Achilles' heel.

Table A2-39, Cluster 38

Cluster 38 focuses on Zika virus primarily and flaviviruses and Dengue virus secondarily, emphasizing immune system evasion (201)
Nonstructural Proteins Are Preferential Positive Selection Targets in Zika Virus and Related Flaviviruses.
Immune Response to Dengue and Zika.
A Mutation Identified in Neonatal Microcephaly Destabilizes Zika Virus NS1 Assembly in Vitro.
Screening Pipeline for Flavivirus Based Inhibitors for Zika Virus NS1.
Contribution of intertwined loop to membrane association revealed by Zika virus full-length NS1 structure.
Zika virus NS1 structure reveals diversity of electrostatic surfaces among flaviviruses.
Levels of Circulating NS1 Impact West Nile Virus Spread to the Brain.
Flavivirus NS1: a multifaceted enigmatic viral protein.
Transposon Mutagenesis of the Zika Virus Genome Highlights Regions Essential for RNA Replication and Restricted for Immune Evasion.
Antibodies targeting epitopes on the cell-surface form of NS1 protect against Zika virus infection during pregnancy.
Zika Virus: Immune Evasion Mechanisms, Currently Available Therapeutic Regimens, and Vaccines.
A short N-terminal peptide motif on flavivirus nonstructural protein NS1 modulates cellular targeting and immune recognition.
Flavivirus NS1 Triggers Tissue-Specific Vascular Endothelial Dysfunction Reflecting Disease Tropism.
The Good, the Bad, and the Shocking: The Multiple Roles of Dengue Virus Nonstructural Protein 1 in Protection and Pathogenesis.
Secretory pathways and multiple functions of nonstructural protein 1 in flavivirus infection.
Advances in Zika Virus-Host Cell Interaction: Current Knowledge and Future Perspectives.
Replication in the presence of dengue convalescent serum impacts Zika virus neutralization sensitivity and fitness.
Dengue Virus Non-structural Protein 1 Modulates Infectious Particle Production via Interaction with the Structural Proteins.
Cross-reactive antibodies targeting surface-exposed non-structural protein 1 (NS1) of dengue virus-infected cells recognize epitopes on the spaghetti loop of the $\beta$ -ladder domain.
Flavivirus NS1 structures reveal surfaces for associations with membranes and the immune system.
Pathogenesis and Manifestations of Zika Virus-Associated Ocular Diseases.
Immune Recognition versus Immune Evasion Systems in Zika Virus Infection.
Pathogenesis of Zika Virus Infection.
Immune evasion strategies of flaviviruses.
Zika virus pathogenesis and current therapeutic advances.
Zika Virus Subgenomic Flavivirus RNA Generation Requires Cooperativity between Duplicated RNA Structures That Are Essential for Productive Infection in Human Cells.
Evasion of Innate and Intrinsic Antiviral Pathways by the Zika Virus.
Arbovirus-vector protein interactomics identifies Loquacious as a co-factor for dengue virus replication in Aedes mosquitoes.
Generation of Zika virus-specific T cells from seropositive and virus-naïve donors for potential use as an autologous or "off-the-shelf" immunotherapeutic.
Structure-guided insights on the role of NS1 in flavivirus infection.
Current Advances in Zika Vaccine Development.
Innate Immune Subversion Strategies of Human Flaviviruses.

Modulation of cellular machineries by Zika virus-encoded proteins.
Neuroimmune Evasion of Zika Virus to Facilitate Viral Pathogenesis.
N-glycosylation of Viral E Protein Is the Determinant for Vector Midgut Invasion by Flaviviruses.
Human Fetal Astrocytes Infected with Zika Virus Exhibit Delayed Apoptosis and Resistance to Interferon: Implications for Persistence.
Mapping the interactions of dengue virus NS1 protein with human liver proteins using a yeast two-hybrid system: identification of C1q as an interacting partner.
The Complement System in Flavivirus Infections.
An evolutionary NS1 mutation enhances Zika virus evasion of host interferon induction.
Zika Virus Pathogenesis: A Battle for Immune Evasion.
ApoA1 Neutralizes Proinflammatory Effects of Dengue Virus NS1 Protein and Modulates Viral Immune Evasion.
Abrogation of TLR3 inhibition by discrete amino acid changes in the C-terminal half of the West Nile virus NS1 protein.
Comparative Analysis of African and Asian Lineage-Derived Zika Virus Strains Reveals Differences in Activation of and Sensitivity to Antiviral Innate Immunity.
Long non-coding subgenomic flavivirus RNAs have extended 3D structures and are flexible in solution.
Secreted NS1 Protects Dengue Virus from Mannose-Binding Lectin-Mediated Neutralization.
[Flaviviruses].
Immune Evasion Strategies Used by Zika Virus to Infect the Fetal Eye and Brain.
Adaptation to host cell environment during experimental evolution of Zika virus.
The flavivirus NS1 protein: molecular and structural biology, immunology, role in pathogenesis and application as a diagnostic biomarker.
Innate and adaptive immune evasion by dengue virus.
Antagonism of the complement component C4 by flavivirus nonstructural protein NS1.
In silico analysis revealed Zika virus miRNAs associated with viral pathogenesis through alteration of host genes involved in immune response and neurological functions.
Structural basis of Flavivirus NS1 assembly and antibody recognition.
The innate immune response in Zika virus infection.
Dengue Virus Glycosylation: What Do We Know?
Innate Antiviral Immunity against Dengue Virus.
Novel approaches to flavivirus drug discovery.
Self-association features of NS1 proteins from different flaviviruses.
Evasion of the human innate immune system by dengue virus.
The Zika Virus Capsid Disrupts Corticogenesis by Suppressing Dicer Activity and miRNA Biogenesis.
Zika virus infection differentially affects genome-wide transcription in neuronal cells and myeloid dendritic cells.
Dengue virus genomic variation associated with mosquito adaptation defines the pattern of viral non-coding RNAs and fitness in human cells.
An Evolutionary Insight into Zika Virus Strains Isolated in the Latin American Region.
Viral immune evasion in dengue: toward evidence-based revisions of clinical practice guidelines.
Molecularly barcoded Zika virus libraries to probe in vivo evolutionary dynamics.
Molecular Insight into Dengue Virus Pathogenesis and Its Implications for Disease Control.
Regulatory Role of Host MicroRNAs in Flaviviruses Infection.
Evasion of early innate immune response by 2'-O-methylation of dengue genomic RNA.

Organotypic hippocampal culture model reveals differential responses to highly similar Zika virus isolates.
Non structural protein of avian influenza A (H11N1) virus is a weaker suppressor of immune responses but capable of inducing apoptosis in host cells.
Influenza A Virus NS1 Protein Binds as a Dimer to RNA-Free PABP1 but Not to the PABP1·Poly(A) RNA Complex.
Critical issues in dengue vaccine development.
Flavivirus Infection Impairs Peroxisome Biogenesis and Early Antiviral Signaling.
The Dengue Virus Replication Complex: From RNA Replication to Protein-Protein Interactions to Evasion of Innate Immunity.
Dengue proteins with their role in pathogenesis, and strategies for developing an effective anti-dengue treatment: A review.
Zika virus and the nonmicrocephalic fetus: why we should still worry.
Structure of the dengue virus glycoprotein non-structural protein 1 by electron microscopy and single-particle analysis.
Dengue virus and the host innate immune response.
An epidemic Zika virus isolate suppresses antiviral immunity by disrupting antigen presentation pathways.
Differential proteomics of Zika virus (ZIKV) infection reveals molecular changes potentially involved in immune system evasion by a Brazilian strain of ZIKV.
Molecular pathogenesis of dengue virus infection in Aedes mosquitoes.
Viral receptors for flaviviruses: Not only gatekeepers.
Dendritic cells in dengue virus infection: targets of virus replication and mediators of immunity.
A Brief Review of West Nile Virus Biology.
Zika Virus Inhibits IFN- $\alpha$ Response by Human Plasmacytoid Dendritic Cells and Induces NS1-Dependent Triggering of CD303 (BDCA-2) Signaling.
Respiratory Syncytial Virus NS1 Protein Targets the Transactivator Binding Domain of MED25.
The Role of Host Cholesterol During Flavivirus Infection.
Comparison of immune responses to attenuated rabies virus and street virus in mouse brain.
A glance at subgenomic flavivirus RNAs and microRNAs in flavivirus infections.
Recent advances in dengue pathogenesis and clinical management.
Innate Immune Evasion Mediated by Flaviviridae Non-Structural Proteins.
The Unique Cofactor Region of Zika Virus NS2B-NS3 Protease Facilitates Cleavage of Key Host Proteins.
Characterization of Dendritic Cell-Derived Extracellular Vesicles During Dengue Virus Infection.
Immune evasion, a critical strategy for rabies virus.
Dengue virus infection and Nrf2 regulation of oxidative stress.
DDX21 translocates from nucleus to cytoplasm and stimulates the innate immune response due to dengue virus infection.
A detailed comparative analysis of codon usage bias in Zika virus.
Mammalian Adaptation of an Avian Influenza A Virus Involves Stepwise Changes in NS1.
Mechanisms of immune evasion induced by a complex of dengue virus and preexisting enhancing antibodies.
Partial maturation: an immune-evasion strategy of dengue virus?
Immunological aspects of rabies: a literature review.

Critical Role of K1685 and K1829 in the Large Protein of Rabies Virus in Viral Pathogenicity and Immune Evasion.
The importance of immune evasion in the pathogenesis of rabies virus.
Dengue Virus Perturbs Mitochondrial Morphodynamics to Dampen Innate Immune Responses.
Evasion of innate and adaptive immunity by flaviviruses.
Skin innate immune response to flaviviral infection.
Interaction of rabies virus P-protein with STAT proteins is critical to lethal rabies disease.
Dengue Virus Selectively Annexes Endoplasmic Reticulum-Associated Translation Machinery as a Strategy for Co-opting Host Cell Protein Synthesis.
Research Advances on the Interactions between Rabies Virus Structural Proteins and Host Target Cells: Accrued Knowledge from the Application of Reverse Genetics Systems.
Autophagy-associated dengue vesicles promote viral transmission avoiding antibody neutralization.
Let's Get Physical: Flavivirus-Host Protein-Protein Interactions in Replication and Pathogenesis.
Subversion of the Immune Response by Rabies Virus.
The Human STAT2 Coiled-Coil Domain Contains a Degron for Zika Virus Interferon Evasion.
The role of toll-like receptors in the induction of immune responses during rabies virus infection.
The journey of Zika to the developing brain.
Profile of Cytokines and Chemokines Triggered by Wild-Type Strains of Rabies Virus in Mice.
Cross-reacting antibodies enhance dengue virus infection in humans.
Role of microRNAs in antiviral responses to dengue infection.
The role of viral persistence in flavivirus biology.
Dual R108K and G189D Mutations in the NS1 Protein of A/H1N1 Influenza Virus Counteract Host Innate Immune Responses.
Flavivirus RNA methylation.
Protective immunity to DENV2 after immunization with a recombinant NS1 protein using a genetically detoxified heat-labile toxin as an adjuvant.
An epigenetic 'extreme makeover': the methylation of flaviviral RNA (and beyond).
Up-regulation of Fas ligand (FasL) in the central nervous system: a mechanism of immune evasion by rabies virus.
High flavivirus structural plasticity demonstrated by a non-spherical morphological variant.
Mechanism of Immune Evasion in Mosquito-Borne Diseases.
Flavivirus Persistence in Wildlife Populations.
Dengue virus-1 NS5 genetic variant associated with a severe clinical infection: Possible reduction of the innate immune response by inhibition of interferon type 1 and the Janus kinase-signal transducer and activator of transcription signaling pathway.
Susceptibilities of CNS Cells towards Rabies Virus Infection Is Linked to Cellular Innate Immune Responses.
Membrane topology and function of dengue virus NS2A protein.
Zika virus NS1 suppresses the innate immune responses via miR-146a in human microglial cells.
Modulation of mRNA Translation and Cell Viability by Influenza A Virus Derived Nonstructural Protein 1.
Insect-specific flavivirus infection is restricted by innate immunity in the vertebrate host.
Rabies virus clearance from the central nervous system.
Experimental evidence and molecular modeling of the interaction between hRSV-NS1 and quercetin.
Flavivirus RNA transactions from viral entry to genome replication.
The Autophagosomes Containing Dengue Virus Proteins and Full-Length Genomic RNA Are Infectious.



Identification of a novel antiviral micro-RNA targeting the NS1 protein of the H1N1 pandemic human influenza virus and a corresponding viral escape mutation.
The West Nile virus assembly process evades the conserved antiviral mechanism of the interferon-induced MxA protein.
Evasive strategies in rabies virus infection.
Innate Immune Response of Primary Human Keratinocytes to West Nile Virus Infection and Its Modulation by Mosquito Saliva.
Lyssaviruses and the Fatal Encephalitic Disease Rabies.
Rabies - epidemiology, pathogenesis, public health concerns and advances in diagnosis and control: a comprehensive review.
West Nile virus: the complex biology of an emerging pathogen.
In Silico Analysis of Dengue Virus Serotype 2 Mutations Detected at the Intrahost Level in Patients with Different Clinical Outcomes.
Circulating levels of soluble MICB in infants with symptomatic primary dengue virus infections.
Innate immune evasion by hepatitis C virus and West Nile virus.
Immune responses to an attenuated West Nile virus NS4B-P38G mutant strain.
Co-delivery of NS1 and BMP2 mRNAs to murine pluripotent stem cells leads to enhanced BMP-2 expression and osteogenic differentiation.
Viral Determinants of Virulence in Tick-Borne Flaviviruses.
Detrimental contribution of the immuno-inhibitor B7-H1 to rabies virus encephalitis.
H3N2 canine influenza virus NS1 protein inhibits canine NLRP3 inflammasome activation.
Cullin4 Is Pro-Viral during West Nile Virus Infection of Culex Mosquitoes.
Messenger RNA translation enhancement by immune evasion proteins: a comparative study between EKB (vaccinia virus) and NS1 (influenza A virus).
Intrahost Selection Pressures Drive Rapid Dengue Virus Microevolution in Acute Human Infections.
Passive carriage of rabies virus by dendritic cells.
Deficient Incorporation of Rabies Virus Glycoprotein into Virions Enhances Virus-Induced Immune Evasion and Viral Pathogenicity.
Dengue Non-coding RNA: TRIMmed for Transmission.
A Human Skin Model for Assessing Arboviral Infections.
Development and characterization of reverse genetics system for the Indian West Nile virus lineage 1 strain 68856.
Non-structural protein 5 (NS5) as a target for antiviral development against established and emergent flaviviruses.
Influenza A Virus NS1 Protein Inhibits the NLRP3 Inflammasome.
The 5' and 3' Untranslated Regions of the Flaviviral Genome.
West Nile virus differentially modulates the unfolded protein response to facilitate replication and immune evasion.
Role of virus-induced neuropeptides in the brain in the pathogenesis of rabies.
Comparison of lncRNA and mRNA expression in mouse brains infected by a wild-type and a lab-attenuated Rabies lyssavirus.
Intrinsically Disordered Side of the Zika Virus Proteome.
Non-linear enhancement of mRNA delivery efficiencies by influenza A derived NS1 protein engendering host gene inhibition property.
Definition of the immune evasion-replication interface of rabies virus P protein.

A single amino acid substitution in the central portion of the West Nile virus NS4B protein confers a highly attenuated phenotype in mice.
Structural comparison of the C-terminal domain of functionally divergent lyssavirus P proteins.
Immune modulation by flaviviruses.
The non-structural NS1 protein unique to respiratory syncytial virus: a two-state folding monomer in quasi-equilibrium with a stable spherical oligomer.
Biomimetic inorganic camouflage circumvents antibody-dependent enhancement of infection.
Research progress in RNA interference against the infection of mosquito-borne viruses.
Lyssavirus P Protein Isoforms Diverge Significantly in Subcellular Interactions Underlying Mechanisms of Interferon Antagonism.
Molecular Basis of Functional Effects of Phosphorylation of the C-Terminal Domain of the Rabies Virus P Protein.
The Amino Acid at Position 95 in the Matrix Protein of Rabies Virus Is Involved in Antiviral Stress Granule Formation in Infected Cells.
Full genome comparison and characterization of avian H10 viruses with different pathogenicity in Mink ( <i>Mustela vison</i> ) reveals genetic and functional differences in the non-structural gene.
Quantitative Analysis of the Microtubule Interaction of Rabies Virus P3 Protein: Roles in Immune Evasion and Pathogenesis.
Immune evasion by rabies viruses through the maintenance of blood-brain barrier integrity.
Dengue virus 3 genotype I (GI) lineage 1 (L1) isolates elicit differential cytopathic effect with syncytium formation in human glioblastoma cells (U251).
Light pollution increases West Nile virus competence of a ubiquitous passerine reservoir species.
The rabies virus interferon antagonist P protein interacts with activated STAT3 and inhibits Gp130 receptor signaling.
The generation of a reverse genetics system for Kyasanur Forest Disease Virus and the ability to antagonize the induction of the antiviral state in vitro.
[Rabies virus glycoprotein: structure, immunogenicity and pathogenic role].
Phenotypic Divergence of P Proteins of Australian Bat Lyssavirus Lineages Circulating in Microbats and Flying Foxes.
Concepts in the pathogenesis of rabies.
Minocycline differentially modulates macrophage mediated peripheral immune response following Japanese encephalitis virus infection.
Structural insights into the multifunctionality of rabies virus P3 protein.
The ectodomain of rabies virus glycoprotein determines dendritic cell activation.
Mosquito immunity against arboviruses.
Neutralization escape variant of West Nile virus associated with altered peripheral pathogenicity and differential cytokine profile.
Yellow Fever Virus Down-Regulates mRNA Expression of SOCS1 in the Initial Phase of Infection in Human Cell Lines.
Differential effects of dengue virus on infected and bystander dendritic cells.
Conservation of a unique mechanism of immune evasion across the Lyssavirus genus.
Transcriptomic Analysis Suggests the M1 Polarization and Launch of Diverse Programmed Cell Death Pathways in Japanese Encephalitis Virus-Infected Macrophages.
Systems Biology Reveals NS4B-Cyclophilin A Interaction: A New Target to Inhibit YFV Replication.
Nuclear Trafficking of the Rabies Virus Interferon Antagonist P-Protein Is Regulated by an Importin-Binding Nuclear Localization Sequence in the C-Terminal Domain.
An updated review of avian-origin Tembusu virus: a newly emerging avian Flavivirus.

RIPK3 Promotes JEV Replication in Neurons via Downregulation of IFI44L.
---

Table A2-40, Cluster 39

Cluster 39 focuses on nanoparticles (especially cell membrane-coated biomimetic) for tumor drug delivery, emphasizing increased immune escape/evasion and enhanced circulation time (224)
Research Progress of Cell Membrane Biomimetic Nanoparticles for Tumor Therapy.
Cell membrane-coated nanoparticles: a novel multifunctional biomimetic drug delivery system.
Cell membrane camouflaged nanoparticles: a new biomimetic platform for cancer photothermal therapy.
Cell membrane biomimetic nanoparticles for inflammation and cancer targeting in drug delivery.
Advancements in cell membrane camouflaged nanoparticles: A bioinspired platform for cancer therapy.
Recent Advances in Cell Membrane-Camouflaged Nanoparticles for Cancer Phototherapy.
Hybrid Membrane-Coated Biomimetic Nanoparticles (HM@BNPs): A Multifunctional Nanomaterial for Biomedical Applications.
Cell Membrane Coated Nanoparticles: An Emerging Biomimetic Nanoplatform for Targeted Bioimaging and Therapy.
Research update on cell membrane camouflaged nanoparticles for cancer therapy.
Cell membrane coated nanocarriers - an efficient biomimetic platform for targeted therapy.
Safe and Immunocompatible Nanocarriers Cloaked in RBC Membranes for Drug Delivery to Treat Solid Tumors.
Targeting drugs to tumours using cell membrane-coated nanoparticles.
Membrane engineering of cell membrane biomimetic nanoparticles for nanoscale therapeutics.
Cancer-Erythrocyte Hybrid Membrane-Camouflaged Magnetic Nanoparticles with Enhanced Photothermal-Immunotherapy for Ovarian Cancer.
Hybrid cell membrane-coated nanoparticles: A multifunctional biomimetic platform for cancer diagnosis and therapy.
Platelet-Tumor Cell Hybrid Membrane-Camouflaged Nanoparticles for Enhancing Therapy Efficacy in Glioma.
Cell membrane coating integrity affects the internalization mechanism of biomimetic nanoparticles.
Biomimetic platelet membrane-coated nanoparticles for targeted therapy.
Cancer cell membrane-coated biomimetic platform for targeted therapy of breast cancer in an orthotopic mouse model.
Application of cell membrane-functionalized biomimetic nanoparticles in the treatment of glioma.
Nanoparticle elasticity regulates the formation of cell membrane-coated nanoparticles and their nano-bio interactions.
Current progress of mesenchymal stem cell membrane-camouflaged nanoparticles for targeted therapy.
Cancer cell membrane coated PLGA nanoparticles as biomimetic drug delivery system for improved cancer therapy.
Cancer cell membrane biomimetic nanosystem for homologous targeted dual-mode imaging and combined therapy.
Cell membrane-coated nanosized active targeted drug delivery systems homing to tumor cells: A review.

Preparation and evaluation of long circulating erythrocyte membrane-cloaked anti-cancer drug delivery system.
Cell membrane-coated nanomaterials for cancer therapy.
Biomimetic hybrid membrane-based nanoplatfroms: synthesis, properties and biomedical applications.
Cell Membrane-Camouflaged Nanoparticles: A Promising Biomimetic Strategy for Cancer Theragnostics.
Leutosome: A Biomimetic Nanoplatfrom Integrating Plasma Membrane Components of Leukocytes and Tumor Cells for Remarkably Enhanced Solid Tumor Homing.
Advanced targeted drug delivery by bioengineered white blood cell-membrane camouflaged nanoparticulate delivery nanostructures.
NIR-responsive cancer cytomembrane-cloaked carrier-free nanosystems for highly efficient and self-targeted tumor drug delivery.
Light-Triggered Biomimetic Nanoerythrocyte for Tumor-Targeted Lung Metastatic Combination Therapy of Malignant Melanoma.
A bionic "Trojan horse"-like gene delivery system hybridized with tumor and macrophage cell membrane for cancer therapy.
Bone-targeted erythrocyte-cancer hybrid membrane-camouflaged nanoparticles for enhancing photothermal and hypoxia-activated chemotherapy of bone invasion by OSCC.
Melanoma Cell Membrane Biomimetic Versatile CuS Nanoprobes for Homologous Targeting Photoacoustic Imaging and Photothermal Chemotherapy.
Platelet-Like Gold Nanostars for Cancer Therapy: The Ability to Treat Cancer and Evade Immune Reactions.
Nanoparticles disguised as red blood cells to evade the immune system.
Light/pH-Triggered Biomimetic Red Blood Cell Membranes Camouflaged Small Molecular Drug Assemblies for Imaging-Guided Combinational Chemo-Photothermal Therapy.
Cell membrane biomimetic nanoparticles in drug delivery.
Cell membrane cloaked nanomedicines for bio-imaging and immunotherapy of cancer: Improved pharmacokinetics, cell internalization and anticancer efficacy.
Bioactive cytomembrane@poly(citrate-peptide)-miRNA365 nanoplatfrom with immune escape and homologous targeting for colon cancer therapy.
Construction of homologous cancer cell membrane camouflage in a nano-drug delivery system for the treatment of lymphoma.
Recent advances in cell membrane coated metal-organic frameworks (MOFs) for tumor therapy.
Macrophage membrane-coated iron oxide nanoparticles for enhanced photothermal tumor therapy.
Cancer Cell Membrane-Camouflaged Nanorods with Endoplasmic Reticulum Targeting for Improved Antitumor Therapy.
Cell-Based Biohybrid Drug Delivery Systems: The Best of the Synthetic and Natural Worlds.
An erythrocyte membrane-camouflaged biomimetic nanoplatfrom for enhanced chemo-photothermal therapy of breast cancer.
Tailoring the Inherent Properties of Biobased Nanoparticles for Nanomedicine.
Biomimetic Upconversion Nanoparticles and Gold Nanoparticles for Novel Simultaneous Dual-Modal Imaging-Guided Photothermal Therapy of Cancer.
Platelet-membrane-camouflaged bismuth sulfide nanorods for synergistic radio-photothermal therapy against cancer.
Harnessing the nano-bio interface: Application of membrane coating to long acting silica particles.

Long circulation and tumor-targeting biomimetic nanoparticles for efficient chemo/photothermal synergistic therapy.
Nano Drug Delivery System for Tumor Immunotherapy: Next-Generation Therapeutics.
Cancer cell membrane-coated biomimetic platform for tumor targeted photodynamic therapy and hypoxia-amplified bio-reductive therapy.
Bioinspired tumor-homing nanoplatfor for co-delivery of paclitaxel and siRNA-E7 to HPV-related cervical malignancies for synergistic therapy.
A erythrocyte-platelet hybrid membrane coated biomimetic nanosystem based on ginsenosides and PFH combined with ultrasound for targeted delivery in thrombus therapy.
A biomimetic cascade nanoreactor for tumor targeted starvation therapy-amplified chemotherapy.
Cancer Cell Membrane-Coated Upconversion Nanoprobes for Highly Specific Tumor Imaging.
A cancer cell membrane coated nanoparticles-based gene delivery system for enhancing cancer therapy.
Cell membrane nanomaterials composed of phospholipids and glycoproteins for drug delivery in inflammatory bowel disease: A review.
Drug Targeting via Platelet Membrane-Coated Nanoparticles.
Development of a nanodrug-delivery system camouflaged by erythrocyte membranes for the chemo/phototherapy of cancer.
Cancer Cell Membrane Camouflaged Mesoporous Silica Nanoparticles Combined with Immune Checkpoint Blockade for Regulating Tumor Microenvironment and Enhancing Antitumor Therapy.
Biomimetic nanoparticles for tumor immunotherapy.
PEGylated WS(2) nanodrug system with erythrocyte membrane coating for chemo/photothermal therapy of cervical cancer.
Cancer Cell Membrane-Coated Nanosuspensions for Enhanced Chemotherapeutic Treatment of Glioma.
Self-Assembly of Extracellular Vesicle-like Metal-Organic Framework Nanoparticles for Protection and Intracellular Delivery of Biofunctional Proteins.
Targeted co-delivery of PD-L1 monoclonal antibody and sorafenib to circulating tumor cells via platelet-functionalized nanocarriers.
Enhanced photothermal therapy of biomimetic polypyrrole nanoparticles through improving blood flow perfusion.
Red Blood Cell Membrane Camouflaged Mesoporous Silica Nanorods as Nanocarriers for Synergistic Chemo-Photothermal Therapy.
Polymeric capsule-cushioned leukocyte cell membrane vesicles as a biomimetic delivery platform.
Immune cell membrane-based biomimetic nanomedicine for treating cancer metastasis.
Cancer-Cell-Membrane-Coated Nanoparticles with a Yolk-Shell Structure Augment Cancer Chemotherapy.
Red-blood-cell-mimetic gene delivery systems for long circulation and high transfection efficiency in ECs.
Carrier-free highly drug-loaded biomimetic nanosuspensions encapsulated by cancer cell membrane based on homology and active targeting for the treatment of glioma.
Biomimetic recombinant of red blood cell membranes for improved photothermal therapy.
Tumor Microenvironment-Adaptive Nanoplatfor Synergistically Enhances Cascaded Chemodynamic Therapy.
Platelets for advanced drug delivery in cancer.
Tumor microenvironment-responsive versatile "Trojan horse" theranostic nanoplatfor for magnetic resonance imaging-guided multimodal synergistic antitumor treatment.

Microenvironment-driven sequential ferroptosis, photodynamic therapy, and chemotherapy for targeted breast cancer therapy by a cancer-cell-membrane-coated nanoscale metal-organic framework.
Cancer-cell-biomimetic Upconversion nanoparticles combining chemo-photodynamic therapy and CD73 blockade for metastatic triple-negative breast cancer.
Pro-efferocytic macrophage membrane biomimetic nanoparticles for the synergistic treatment of atherosclerosis via competition effect.
MSCs-engineered biomimetic PMAA nanomedicines for multiple bioimaging-guided and photothermal-enhanced radiotherapy of NSCLC.
Microfluidic Sonication To Assemble Exosome Membrane-Coated Nanoparticles for Immune Evasion-Mediated Targeting.
Self-Assembly of Podophyllotoxin-Loaded Lipid Bilayer Nanoparticles for Highly Effective Chemotherapy and Immunotherapy via Downregulation of Programmed Cell Death Ligand 1 Production.
Experimental Quantification of Interactions Between Drug Delivery Systems and Cells In Vitro: A Guide for Preclinical Nanomedicine Evaluation.
Nano-Chitosan Particles in Anticancer Drug Delivery: An Up-to-Date Review.
A biomimetic nanoreactor for synergistic chemiexcited photodynamic therapy and starvation therapy against tumor metastasis.
Extracellular matrix modulating enzyme functionalized biomimetic Au nanoplatfrom-mediated enhanced tumor penetration and synergistic antitumor therapy for pancreatic cancer.
RBC membrane camouflaged prussian blue nanoparticles for gamabutolin loading and combined chemo/photothermal therapy of breast cancer.
Orchestration of biomimetic membrane coating and nanotherapeutics in personalized anticancer therapy.
Engineering of Neutrophil Membrane Camouflaging Nanoparticles Realizes Targeted Drug Delivery for Amplified Antitumor Therapy.
Radiolabeled theranostics: magnetic and gold nanoparticles.
Cell Membrane-Camouflaged Nanocarriers for Cancer Diagnostic and Therapeutic.
Cancer Cell Membrane Camouflaged Cascade Bioreactor for Cancer Targeted Starvation and Photodynamic Therapy.
Tumor targeted cancer membrane-camouflaged ultra-small Fe nanoparticles for enhanced collaborative apoptosis and ferroptosis in glioma.
Targeting macrophage endocytosis via platelet membrane coating for advanced osteoimmunomodulation.
Erythrocyte-Camouflaged Mesoporous Titanium Dioxide Nanoplatfrom for an Ultrasound-Mediated Sequential Therapies of Breast Cancer.
B16F10 Cell Membrane-Based Nanovesicles for Melanoma Therapy Are Superior to Hyaluronic Acid-Modified Nanocarriers.
Synthetic nanoparticles functionalized with biomimetic leukocyte membranes possess cell-like functions.
Cell membrane camouflaged cerium oxide nanocubes for targeting enhanced tumor-selective therapy.
Engineered Red Blood Cell Biomimetic Nanovesicle with Oxygen Self-Supply for Near-Infrared-II Fluorescence-Guided Synergetic Chemo-Photodynamic Therapy against Hypoxic Tumors.
Surface functionalization of gold nanoparticles with red blood cell membranes.
Sialylation: An alternative to designing long-acting and targeted drug delivery system.

Autologous cell membrane coatings on tissue engineering xenografts for suppression and alleviation of acute host immune responses.
Platelet-biomimetic nanoparticles for in vivo targeted photodynamic therapy of breast cancer.
Biomimetic Nanotechnology: A Natural Path Forward for Tumor-Selective and Tumor-Specific NIR Activable Photonanomedicines.
A Multifunctional Biomimetic Nanoplatfrom for Relieving Hypoxia to Enhance Chemotherapy and Inhibit the PD-1/PD-L1 Axis.
Polydopamine Nanoparticles Camouflaged by Stem Cell Membranes for Synergistic Chemo-Photothermal Therapy of Malignant Bone Tumors.
Gold nanoparticles-mediated photothermal therapy and immunotherapy.
The strategy of precise targeting and in situ oxygenating for enhanced triple-negative breast cancer chemophototherapy.
Tumor-targeted biomimetic nanoplatfrom precisely integrates photodynamic therapy and autophagy inhibition for collaborative treatment of oral cancer.
Cancer-cell-membrane-camouflaged supramolecular self-assembly of antisense oligonucleotide and chemodrug for targeted combination therapy.
A biomimetic theranostic O(2)-meter for cancer targeted photodynamic therapy and phosphorescence imaging.
Immune-regulating camouflaged nanoplatforms: A promising strategy to improve cancer nano-immunotherapy.
Platelet-mimicking nanoparticles co-loaded with W(18)O(49) and metformin alleviate tumor hypoxia for enhanced photodynamic therapy and photothermal therapy.
Polymeric coating lubricates nanocontainers to escape macrophage uptake for bioreceptor recognition.
Biomimetic 2D layered double hydroxide nanocomposites for hyperthermia-facilitated homologous targeting cancer photo-chemotherapy.
Cancer Cyto-membrane-Cloaked Prussian Blue Nanoparticles Enhance the Efficacy of Mild-Temperature Photothermal Therapy by Disrupting Mitochondrial Functions of Cancer Cells.
'Stealth' nanoparticles evade neural immune cells but also evade major brain cell populations: Implications for PEG-based neurotherapeutics.
Cell Membrane-Coated Mimics: A Methodological Approach for Fabrication, Characterization for Therapeutic Applications, and Challenges for Clinical Translation.
Hybrid membrane camouflaged copper sulfide nanoparticles for photothermal-chemotherapy of hepatocellular carcinoma.
Cell Membrane-Camouflaged NIR II Fluorescent Ag(2) Te Quantum Dots-Based Nanobioprobes for Enhanced In Vivo Homotypic Tumor Imaging.
"Cluster Bomb" Based on Redox-Responsive Carbon Dot Nanoclusters Coated with Cell Membranes for Enhanced Tumor Theranostics.
Cancer Cell Membrane Biomimetic Mesoporous Nanozyme System with Efficient ROS Generation for Antitumor Chemoresistance.
Biomimetic nanoreactor for targeted cancer starvation therapy and cascade amplified chemotherapy.
Exosome-based nanomedicine for cancer treatment by targeting inflammatory pathways: Current status and future perspectives.
Lipid-hybrid cell-derived biomimetic functional materials: A state-of-the-art multifunctional weapon against tumors.

A size-tunable nanoplatform: enhanced MMP2-activated chemo-photodynamic immunotherapy based on biodegradable mesoporous silica nanoparticles.
Brain Tumor Cell Membrane-Coated Lanthanide-Doped Nanoparticles for NIR-IIb Luminescence Imaging and Surgical Navigation of Glioma.
Emerging self-assembling peptide nanomaterial for anti-cancer therapy.
Ultrasound and laser-promoted dual-gas nano-generator for combined photothermal and immune tumor therapy.
Gene-engineered exosomes-thermosensitive liposomes hybrid nanovesicles by the blockade of CD47 signal for combined photothermal therapy and cancer immunotherapy.
Platelet Membranes Coated Gold Nanocages for Tumor Targeted Drug Delivery and Amplificated Low-Dose Radiotherapy.
How Nanotherapeutic Platforms Play a Key Role in Glioma? A Comprehensive Review of Literature.
Recent Advances in Perfluorocarbon-Based Delivery Systems for Cancer Theranostics.
Mapping the cell-membrane proteome of the SKBR3/HER2+ cell line to the cancer hallmarks.
Light-triggered multifunctional nanoplatform for efficient cancer photo-immunotherapy.
PEI-modified macrophage cell membrane-coated PLGA nanoparticles encapsulating Dendrobium polysaccharides as a vaccine delivery system for ovalbumin to improve immune responses.
Co-delivery of doxorubicin and epacadostat via heparin coated pH-sensitive liposomes to suppress the lung metastasis of melanoma.
Delivery of an ectonucleotidase inhibitor with ROS-responsive nanoparticles overcomes adenosine-mediated cancer immunosuppression.
Novel, Self-Distinguished, Dual Stimulus-Responsive Therapeutic Nanoplatform for Intracellular On-Demand Drug Release.
"Trojan horse" nanoparticle-delivered cancer cell membrane vaccines to enhance cancer immunotherapy by overcoming immune-escape.
Development of Adamantane-Conjugated TLR7/8 Agonists for Supramolecular Delivery and Cancer Immunotherapy.
Cationic lipid-assisted nanoparticles for simultaneous delivery of CD47 siRNA and R848 to promote antitumor immune responses.
A pH-Driven indomethacin-loaded nanomedicine for effective rheumatoid arthritis therapy by combining with photothermal therapy.
A tumor microenvironment responsive biodegradable CaCO <sub>3</sub> /MnO <sub>2</sub> - based nanoplatform for the enhanced photodynamic therapy and improved PD-L1 immunotherapy.
An erythrocyte membrane-modified biomimetic synergistic nanosystem for cancer anti-vascular therapy and initial efficacy monitoring.
M2 Macrophage Hybrid Membrane-Camouflaged Targeted Biomimetic Nanosomes to Reprogram Inflammatory Microenvironment for Enhanced Enzyme-Thermo-Immunotherapy.
Reproducible and Characterized Method for Ponatinib Encapsulation into Biomimetic Lipid Nanoparticles as a Platform for Multi-Tyrosine Kinase-Targeted Therapy.
Integration of Peptides for Enhanced Uptake of PEGylated Gold Nanoparticles.
Controlling the Biological Fate of Micellar Nanoparticles: Balancing Stealth and Targeting.
Cannabidiol-loaded biomimetic macrophage membrane vesicles against post-traumatic stress disorder assisted by ultrasound.
Red Phosphorus/P25 Nanophotosensitizers Coated with Platelet Membrane for Enhancing Cancer Cells Photodynamic Therapy.
Self-Assembly Catalase Nanocomplex Conveyed by Bacterial Vesicles for Oxygenated Photodynamic Therapy and Tumor Immunotherapy.



Engineering of small molecular organic nanoparticles for mitochondria-targeted mild photothermal therapy of malignant breast cancers.
Protein-Polymer Delivery: Chemistry from the Cold Chain to the Clinic.
Multifunctional nanogels for siRNA delivery.
Recent advances in drug delivery systems for targeting brain tumors.
Novel biomimetic dual-mode nanodroplets as ultrasound contrast agents with potential ability of precise detection and photothermal ablation of tumors.
A Dual Drug Delivery Platform for Cancer-Bacteria Cotargeting.
Near-Infrared II Phototherapy Induces Deep Tissue Immunogenic Cell Death and Potentiates Cancer Immunotherapy.
Bispecific prodrug nanoparticles circumventing multiple immune resistance mechanisms for promoting cancer immunotherapy.
Active targeting of orthotopic glioma using biomimetic liposomes co-loaded elemene and cabazitaxel modified by transferritin.
Platelet-camouflaged nanococktail: Simultaneous inhibition of drug-resistant tumor growth and metastasis via a cancer cells and tumor vasculature dual-targeting strategy.
pH-responsive complexes using prefunctionalized polymers for synchronous delivery of doxorubicin and siRNA to cancer cells.
Self-Delivering Nanodrugs Developed via Small-Molecule-Directed Assembly and Macrophage Cloaking for Sonodynamic-Augmented Immunotherapy.
Antitumor immunity triggered by photothermal therapy and photodynamic therapy of a 2D MoS(2) nanosheet-incorporated injectable polypeptide-engineered hydrogel combined with chemotherapy for 4T1 breast tumor therapy.
Biomimetic Nanomaterials Triggered Ferroptosis for Cancer Theranostics.
Immunogenic-cell-killing and immunosuppression-inhibiting nanomedicine.
Nitrogen-Doped Carbon Nanotube Cups for Cancer Therapy.
A tumor extracellular pH-sensitive PD-L1 binding peptide nanoparticle for chemo-immunotherapy of cancer.
A MXene-Based Bionic Cascaded-Enzyme Nanoreactor for Tumor Phototherapy/Enzyme Dynamic Therapy and Hypoxia-Activated Chemotherapy.
Site-specific MOF-based immunotherapeutic nanoplatforms via synergistic tumor cells-targeted treatment and dendritic cells-targeted immunomodulation.
A Biomimetic Nanoplatfrom with Improved Inflammatory Targeting Behavior for ROS Scavenging-Based Treatment of Ulcerative Colitis.
Maximizing the Supported Bilayer Phenomenon: Liposomes Comprised Exclusively of PEGylated Phospholipids for Enhanced Systemic and Lymphatic Delivery.
Precise engineering of targeted nanoparticles by using self-assembled biointegrated block copolymers.
Exosome-Modified Liposomes Targeted Delivery of Thalidomide to Regulate Treg Cells for Antitumor Immunotherapy.
Time-Programmed Delivery of Sorafenib and Anti-CD47 Antibody via a Double-Layer-Gel Matrix for Postsurgical Treatment of Breast Cancer.
Macrophage-Laden Gold Nanoflowers Embedded with Ultrasmall Iron Oxide Nanoparticles for Enhanced Dual-Mode CT/MR Imaging of Tumors.
Induction of anti-tumor cytotoxic T cell responses through PLGA-nanoparticle mediated antigen delivery.
Engineering antiphagocytic biomimetic drug carriers.

Neutrophil membrane fusogenic nanoliposomal leonurine for targeted ischemic stroke therapy via remodeling cerebral niche and restoring blood-brain barrier integrity.
Phototheranostics Using Erythrocyte-Based Particles.
A near-infrared light-responsive extracellular vesicle as a "Trojan horse" for tumor deep penetration and imaging-guided therapy.
Cell-derived membrane biomimetic nanocarriers for targeted therapy of pulmonary disease.
Sequentially pH-Responsive Drug-Delivery Nanosystem for Tumor Immunogenic Cell Death and Cooperating with Immune Checkpoint Blockade for Efficient Cancer Chemoimmunotherapy.
Transforming Cold Tumors into Hot Ones with a Metal-Organic Framework-Based Biomimetic Nanosystem for Enhanced Immunotherapy.
'Smartening' anticancer therapeutic nanosystems using biomolecules.
Regulating Glucose Metabolism with Prodrug Nanoparticles for Promoting Photoimmunotherapy of Pancreatic Cancer.
Cancer Immunotherapy Based on Cell Membrane-Coated Nanocomposites Augmenting cGAS/STING Activation by Efferocytosis Blockade.
Understanding the Effects of Nanocapsular Mechanical Property on Passive and Active Tumor Targeting.
Microfluidic Manufacturing of Multitargeted PLGA/PEG Nanoparticles for Delivery of Taxane Chemotherapeutics.
Small Interfering RNA for Gliomas Treatment: Overcoming Hurdles in Delivery.
Promise and peril in nanomedicine: the challenges and needs for integrated systems biology approaches to define health risk.
Tumor Microenvironment Cascade-Responsive Nanodrug with Self-Targeting Activation and ROS Regeneration for Synergistic Oxidation-Chemotherapy.
Artemisinin and Procyanidins loaded multifunctional nanocomplexes alleviate atherosclerosis via simultaneously modulating lipid influx and cholesterol efflux.
A "bulldozer" driven by anoxic bacteria for pancreatic cancer chemo-immunotherapy.
Cascade-Responsive Hierarchical Nanosystems for Multisite Specific Drug Exposure and Boosted Chemoimmunotherapy.
A Bimetallic Metal-Organic-Framework-Based Biomimetic Nanoplatfrom Enhances Anti-Leukemia Immunity via Synchronizing DNA Demethylation and RNA Hypermethylation.
Blocking CD47 with restructured peptide nanoparticles for motivating phagocytosis to inhibit tumor progression.
Challenges and progress toward tumor-targeted therapy by systemic delivery of polymer-complexed oncolytic adenoviruses.
Macrophages-Cancer Membrane-Encapsulated Metal-Organic Frameworks with Copper-Depleting Moiety for Mitochondria-Targeted Therapeutics.
T(1)- and T(2)-weighted Magnetic Resonance Dual Contrast by Single Core Truncated Cubic Iron Oxide Nanoparticles with Abrupt Cellular Internalization and Immune Evasion.
Multifunctional nanoparticles of sinomenine hydrochloride for treat-to-target therapy of rheumatoid arthritis via modulation of proinflammatory cytokines.
Suppressing or Enhancing Macrophage Engulfment through the Use of CD47 and Related Peptides.
Tumor-derived biomimetic nanozyme with immune evasion ability for synergistically enhanced low dose radiotherapy.
Platelet bio-nanobubbles as microvascular recanalization nanoformulation for acute ischemic stroke lesion theranostics.

Efficient lung orthotopic tumor-growth suppression of oncolytic adenovirus complexed with RGD-targeted bioreducible polymer.
From conventional to stealth liposomes: a new frontier in cancer chemotherapy.
Immunogenic cancer cell death selectively induced by near infrared photoimmunotherapy initiates host tumor immunity.
Ultradense Erythrocyte Bionic Layer Used to Capture Circulating Tumor Cells and Plasma-Assisted High-Purity Release.
Treatment of multi-position condyloma acuminatum using topical CO(2) laser combined with photodynamic therapy - Report of 6 cases.
From conventional to stealth liposomes: a new Frontier in cancer chemotherapy.
Low dose photodynamic-therapy induce immune escape of tumor cells in a HIF-1 $\alpha$ dependent manner through PI3K/Akt pathway.
Biological Function and Immunotherapy Utilizing Phosphatidylserine-based Nanoparticles.
Nanocarrier-mediated brain delivery of bioactives for treatment/prevention of neurodegenerative diseases.
Sustained release of PKR inhibitor C16 from mesoporous silica nanoparticles significantly enhances mRNA translation and anti-tumor vaccination.
Polynorbornene-based bioconjugates by aqueous grafting-from ring-opening metathesis polymerization reduce protein immunogenicity.
Preclinical testing of an anal bulking agent coated with a zwitterionic polymer in a fecal incontinence rat model.
Hydrogel loading functionalized PAMAM/shRNA complex for postsurgical glioblastoma treatment.
Bioreducible polymer-conjugated oncolytic adenovirus for hepatoma-specific therapy via systemic administration.
Quercetin encapsulated in folic acid-modified liposomes is therapeutic against osteosarcoma by non-covalent binding to the JH2 domain of JAK2 Via the JAK2-STAT3-PDL1.

Table A2-41, Cluster 40

Cluster 40 focuses on tumors, especially their escape and evasion from immune system control (402)
Dual Effect of Immune Cells within Tumour Microenvironment: Pro- and Anti-Tumour Effects and Their Triggers.
Tumour escape mechanisms and their therapeutic implications in combination tumour therapy.
Targeting hypoxia-related metabolism molecules: How to improve tumour immune and clinical treatment?
Immunosenescence, suppression and tumour progression.
The role of the tumour microenvironment in immunotherapy.
Tumour-derived IL-10 within tumour microenvironment represses the antitumour immunity of Socs1-silenced and sustained antigen expressing DCs.
Immunosuppressive networks in the tumour environment and their therapeutic relevance.
Mechanisms of immune activation and regulation: lessons from melanoma.
Role of immune escape in different digestive tumours.
Tumour exosomes inhibit binding of tumour-reactive antibodies to tumour cells and reduce ADCC.
Activation of cytotoxic T cells by solid tumours?
TGFbeta is responsible for skin tumour infiltration by macrophages enabling the tumours to escape immune destruction.

The tumour glyco-code as a novel immune checkpoint for immunotherapy.
Evasion of tumours from the control of the immune system: consequences of brief encounters.
Combination cancer immunotherapies tailored to the tumour microenvironment.
Allogeneic IgG combined with dendritic cell stimuli induce antitumour T-cell immunity.
Mesenchymal stromal cells (MSCs) and colorectal cancer: a troublesome twosome for the anti-tumour immune response?
Crosstalk between cancer and immune cells: role of STAT3 in the tumour microenvironment.
Angiostasis as a way to improve immunotherapy.
The immune system--is it relevant to cancer development, progression and treatment?
Tumour microenvironment of pancreatic cancer: immune landscape is dictated by molecular and histopathological features.
Lymphatic and interstitial flow in the tumour microenvironment: linking mechanobiology with immunity.
Immune escape of colorectal tumours via local LRH-1/Cyp11b1-mediated synthesis of immunosuppressive glucocorticoids.
Tumour cell conditioned medium reveals greater M2 skewing of macrophages in the absence of properdin.
Harnessing the immune response to treat cancer.
Metabolic control of tumour progression and antitumour immunity.
Therapy sculpts the complex interplay between cancer and the immune system during tumour evolution.
Escape strategies and reasons for failure in the interaction between tumour cells and the immune system: how can we tilt the balance towards immune-mediated cancer control?
Circulating and disseminated tumour cells - mechanisms of immune surveillance and escape.
Tumour microenvironment and metabolic plasticity in cancer and cancer stem cells: Perspectives on metabolic and immune regulatory signatures in chemoresistant ovarian cancer stem cells.
The emerging promise of tumour mechanobiology in cancer treatment.
The determinants of tumour immunogenicity.
Cleavage of CD95 by matrix metalloproteinase-7 induces apoptosis resistance in tumour cells.
Inflammation and immunity in ovarian cancer.
Molecular insights into tumour metastasis: tracing the dominant events.
Tumour immunotherapy: the adjuvant treatment of the 21st century?
Tumour-mediated TRAIL-Receptor expression indicates effective apoptotic depletion of infiltrating CD8+ immune cells in clinical colorectal cancer.
The acidic tumour microenvironment: Manipulating the immune response to elicit escape.
Hostile takeover: how tumours hijack pre-existing vascular environments to thrive.
Glycosylation in cancer: Selected roles in tumour progression, immune modulation and metastasis.
Immune evasion by tumours: involvement of the CD95 (APO-1/Fas) system and its clinical implications.
Immunosuppressive networks in the tumour environment and their effect in dendritic cells.
Epigenetic promoter alterations in GI tumour immune-editing and resistance to immune checkpoint inhibition.
Tumour hypoxia promotes tolerance and angiogenesis via CCL28 and T(reg) cells.
Immunological responses can have both pro- and antitumour effects: implications for immunotherapy.
Antitumour immune responses.

Tumour Cell Secretome in Chemoresistance and Tumour Recurrence.
Does the tumour microenvironment alter tumorigenesis and clinical response in transmissible venereal tumour in dogs?
Nano-immunotherapy: Overcoming tumour immune evasion.
The hallmarks of ovarian cancer: Focus on angiogenesis and micro-environment and new models for their characterisation.
Targeting tumour-reprogrammed myeloid cells: the new battleground in cancer immunotherapy.
Advances in the understanding of cancer immunotherapy.
Tumour Derived Extracellular Vesicles: Challenging Target to Blunt Tumour Immune Evasion.
Stromal protein $\beta$ ig-h3 reprogrammes tumour microenvironment in pancreatic cancer.
Nature of tumour rejection antigens in ovarian cancer.
Long non-coding RNAs within the tumour microenvironment and their role in tumour-stroma cross-talk.
Radiotherapy and immunotherapy: a synergistic effect in cancer care.
Inflammation and cancer: till death tears them apart.
Saga of monokines in shaping tumour-immune microenvironment: Origin to execution.
Coagulation and inflammation in cancer: Limitations and prospects for treatment.
Myeloid cells are required for PD-1/PD-L1 checkpoint activation and the establishment of an immunosuppressive environment in pancreatic cancer.
Immune selection in neoplasia: towards a microevolutionary model of cancer development.
The role of tumour heterogeneity and clonal cooperativity in metastasis, immune evasion and clinical outcome.
Immunobiology of cholangiocarcinoma.
Trajectory of immune evasion and cancer progression in hepatocellular carcinoma.
Myeloid-derived suppressor cells: Bridging the gap between inflammation and pancreatic adenocarcinoma.
Redirecting oncolytic viruses: Engineering opportunists to take control of the tumour microenvironment.
Tuning cancer fate: the unremitting role of host immunity.
Cancer metastasis as a non-healing wound.
Caspase activation in tumour-infiltrating lymphocytes is associated with lymph node metastasis in oral squamous cell carcinoma.
Progress in research on the role of amino acid metabolic reprogramming in tumour therapy: A review.
Antigenic targets for renal cell carcinoma immunotherapy.
PI3K $\beta$ controls immune evasion in PTEN-deficient breast tumours.
Cancer stem cell-immune cell crosstalk in tumour progression.
Advances in Anti-Cancer Immunotherapy: Car-T Cell, Checkpoint Inhibitors, Dendritic Cell Vaccines, and Oncolytic Viruses, and Emerging Cellular and Molecular Targets.
Targeted depletion of tumour-associated macrophages by an alendronate-glucomannan conjugate for cancer immunotherapy.
Lipid metabolism and tumor immunotherapy.
Phenotype, function and clinical implications of myeloid-derived suppressor cells in cancer patients.
Role of extracellular vesicles in glioma progression.
New roads open up for implementing immunotherapy in mesothelioma.
Host response to colorectal cancer.

Immunology in the clinic review series; focus on cancer: tumour-associated macrophages: undisputed stars of the inflammatory tumour microenvironment.
Local tumour ablative therapies: opportunities for maximising immune engagement and activation.
Advances in the Immunomodulatory Properties of Glycoantigens in Cancer.
Cell biology-metabolic crosstalk in glioma.
Galectins in prostate and bladder cancer: tumorigenic roles and clinical opportunities.
Loss of MAPK-activated protein kinase 2 enables potent dendritic cell-driven anti-tumour T cell response.
Analysis of tumour-immune evasion with chemo-immuno therapeutic treatment with quadratic optimal control.
From Immunosuppression to Immunomodulation - Turning Cold Tumours into Hot.
In silico simulation of a clinical trial with anti-CTLA-4 and anti-PD-L1 immunotherapies in metastatic breast cancer using a systems pharmacology model.
CRIP1 fosters MDSC trafficking and resets tumour microenvironment via facilitating NF- $\kappa$ B/p65 nuclear translocation in pancreatic ductal adenocarcinoma.
Regulating the regulators in cancer-immunosuppression in multiple myeloma (MM).
The Role of Different Immunocompetent Cell Populations in the Pathogenesis of Head and Neck Cancer-Regulatory Mechanisms of Pro- and Anti-Cancer Activity and Their Impact on Immunotherapy.
Is human cytomegalovirus a target in cancer therapy?
Understanding the epigenetic regulation of tumours and their microenvironments: opportunities and problems for epigenetic therapy.
The Tumour Microenvironment and Circulating Tumour Cells: A Partnership Driving Metastasis and Glycan-Based Opportunities for Cancer Control.
miR-25/93 mediates hypoxia-induced immunosuppression by repressing cGAS.
Role of extracellular vesicles in tumour microenvironment.
m(6)A methylation: a process reshaping the tumour immune microenvironment and regulating immune evasion.
CAF-immune cell crosstalk and its impact in immunotherapy.
Tumour-educated macrophages display a mixed polarisation and enhance pancreatic cancer cell invasion.
Targeting neoantigens to augment antitumour immunity.
Mechanisms of HIF-driven immunosuppression in tumour microenvironment.
Insights into the role of sialylation in cancer progression and metastasis.
Metabolic networks in mutant KRAS-driven tumours: tissue specificities and the microenvironment.
Controlling escape from angiogenesis inhibitors.
Epigenetics and immunotherapy: The current state of play.
The Multifaceted Role of Macrophages in Oncolytic Virotherapy.
The hypoxic tumour microenvironment: A safe haven for immunosuppressive cells and a therapeutic barrier to overcome.
T-cell-directed cancer vaccines: mechanisms of immune escape and immune tolerance.
Cancer immune evasion through KRAS and PD-L1 and potential therapeutic interventions.
Immunotherapy of colorectal cancer.
Mutant p53 gain of function mediates cancer immune escape that is counteracted by APR-246.
Viro-immune therapy: A new strategy for treatment of pancreatic cancer.
SLC38A2 and glutamine signalling in cDC1s dictate anti-tumour immunity.
Tumour tissue microenvironment can inhibit dendritic cell maturation in colorectal cancer.

Transforming growth factor-beta1 immobilises dendritic cells within skin tumours and facilitates tumour escape from the immune system.
Challenges in glioblastoma immunotherapy: mechanisms of resistance and therapeutic approaches to overcome them.
Immune biomarkers for predicting response to adoptive cell transfer as cancer treatment.
Tumour growth and immune evasion as targets for a new strategy in advanced cancer.
OAS3 is a Co-Immune Biomarker Associated With Tumour Microenvironment, Disease Staging, Prognosis, and Treatment Response in Multiple Cancer Types.
TGF- $\beta$ signalling in tumour associated macrophages.
Apoptosis drives cancer cells proliferate and metastasize.
Ovarian cancer mutational processes drive site-specific immune evasion.
Crosstalk between fatty acid metabolism and tumour-associated macrophages in cancer progression.
Microglia promote glioblastoma via mTOR-mediated immunosuppression of the tumour microenvironment.
Serum lactate as a potential biomarker of malignancy in primary adult brain tumours.
Combination of epigenetic regulation with gene therapy-mediated immune checkpoint blockade induces anti-tumour effects and immune response in vivo.
Loss of viral genome with altered immune microenvironment during tumour progression of Epstein-Barr virus-associated gastric carcinoma.
Aggressive serous epithelial ovarian cancer is potentially propagated by EpCAM(+)/CD45(+) phenotype.
Targeting PPAR-gamma counteracts tumour adaptation to immune-checkpoint blockade in hepatocellular carcinoma.
The biology and analysis of single disseminated tumour cells.
An organoid-based screen for epigenetic inhibitors that stimulate antigen presentation and potentiate T-cell-mediated cytotoxicity.
T-cell modulation by cyclophosphamide for tumour therapy.
Interleukin-10 attenuates tumour growth by inhibiting interleukin-6/signal transducer and activator of transcription 3 signalling in myeloid-derived suppressor cells.
KDM5B promotes immune evasion by recruiting SETDB1 to silence retroelements.
Mechanisms of Drug Resistance in Melanoma.
Combining radiation and cancer gene therapy: a potential marriage of physical and biological targeting?
Enhancing cancer immunotherapy using antiangiogenics: opportunities and challenges.
Genomic-transcriptomic evolution in lung cancer and metastasis.
Expression pattern of immune suppressive cytokines and growth factors in oesophageal adenocarcinoma reveal a tumour immune escape-promoting microenvironment.
Role of N-acetylgalactosaminyltransferase 6 in early tumorigenesis and formation of metastasis.
The immune landscape of neuroblastoma: Challenges and opportunities for novel therapeutic strategies in pediatric oncology.
Challenges and future perspectives of T cell immunotherapy in cancer.
Hypoxia and acidosis: immune suppressors and therapeutic targets.
Cancer immunoediting and resistance to T cell-based immunotherapy.
Mast cells as regulators of adaptive immunity to tumours.
Vaccines for established cancer: overcoming the challenges posed by immune evasion.
Non-steroidal anti-inflammatory drugs, tumour immunity and immunotherapy.

The effect of acute acid exposure on immunomodulatory protein secretion, cell survival, and cell cycle progression in tumour cell lines.
The oncogenic and clinical implications of lactate induced immunosuppression in the tumour microenvironment.
Genetically modified tumour vaccines: an obstacle race to break host tolerance to cancer.
Leptomeningeal Metastases in Melanoma Patients: An Update on and Future Perspectives for Diagnosis and Treatment.
Vaccines as early therapeutic interventions for cancer therapy: neutralising the immunosuppressive tumour environment and increasing T cell avidity may lead to improved responses.
[Tumour-derived exosomes and their roles in cancer].
[Dissecting the pathways of tumour escape: "question of life and death?"].
Targeting stromal cell Syndecan-2 reduces breast tumour growth, metastasis and limits immune evasion.
PI3K $\delta$ inhibitor plus radiation enhances the antitumour immune effect of PD-1 blockade in syngenic murine breast cancer and humanised patient-derived xenograft model.
New insights into cancer immunoediting and its three component phases--elimination, equilibrium and escape.
Cancer Associated Fibroblasts - An Impediment to Effective Anti-Cancer T Cell Immunity.
Genomic and transcriptomic heterogeneity of colorectal tumours arising in Lynch syndrome.
TGF $\beta$ drives immune evasion in genetically reconstituted colon cancer metastasis.
Lung cancer and Toll-like receptors.
What have we learned from cancer immunotherapy in the last 3 years?
Epithelial Ovarian Cancer and the Immune System: Biology, Interactions, Challenges and Potential Advances for Immunotherapy.
Hallmarks of the Tumour Microenvironment of Gliomas and Its Interaction with Emerging Immunotherapy Modalities.
Tumour-infiltrating Gr-1+ myeloid cells antagonize senescence in cancer.
Vaccination therapy in malignant disease.
Actual status of therapeutic vaccination in non-small cell lung cancer.
Role of tissue microenvironment resident adipocytes in colon cancer.
Indoleamine 2,3-Dioxygenase: A Novel Immunotherapeutic Target for Osteosarcoma.
Hypermutated tumours in the era of immunotherapy: The paradigm of personalised medicine.
Immunology and immunotherapy of human cancer: present concepts and clinical developments.
Immune cell promotion of metastasis.
Research progress on tumour-associated macrophages in gastric cancer (Review).
Harnessing immunity for therapy in human papillomavirus driven cancers.
Tumour neoantigen heterogeneity thresholds provide a time window for combination immunotherapy.
Does the devil facial tumour produce immunosuppressive cytokines as an immune evasion strategy?
Rise of the natural red pigment 'prodigiosin' as an immunomodulator in cancer.
Local Breast Microbiota: A "New" Player on the Block.
Understanding patterns of invasion: a novel approach to assessment of podoplanin expression in the prediction of lymph node metastasis in oral squamous cell carcinoma.
Galectin inhibitory disaccharides promote tumour immunity in a breast cancer model.
Recent insights into the biology of pancreatic cancer.



Intercellular communication in the tumour microecosystem: Mediators and therapeutic approaches for hepatocellular carcinoma.
New approaches in the immunotherapy of haematological malignancies.
Inhibitory B7-family molecules in the tumour microenvironment.
Organoids as an Enabler of Precision Immuno-Oncology.
[Gene therapy of childhood cancers: current status and perspectives].
Targeted inhibition of STAT3 induces immunogenic cell death of hepatocellular carcinoma cells via glycolysis.
Y chromosome loss in cancer drives growth by evasion of adaptive immunity.
Leukaemia cell of origin identified by chromatin landscape of bulk tumour cells.
The role of microglia and P2X7 receptors in gliomas.
Immunotherapy of colorectal cancer: Challenges for therapeutic efficacy.
Enhancing anti-tumour innate immunity by targeting the DNA damage response and pattern recognition receptors in combination with radiotherapy.
MEK inhibition abrogates sunitinib resistance in a renal cell carcinoma patient-derived xenograft model.
Molecules in cancer immunotherapy: benefits and side effects.
Targeting of tumour-infiltrating macrophages via CCL2/CCR2 signalling as a therapeutic strategy against hepatocellular carcinoma.
$\gamma$ -Secretase inhibitor reduces immunosuppressive cells and enhances tumour immunity in head and neck squamous cell carcinoma.
Barrett's to oesophageal cancer sequence: a model of inflammatory-driven upper gastrointestinal cancer.
Immune cell infiltration of intrinsic and metastatic intracranial tumours.
Evolving immunotherapeutic strategies in bladder and renal cancer.
How tumours escape mass destruction.
Comparative analysis of peripheral and localised cytokine secretion in glioblastoma patients.
Hypoxia skews dendritic cells to a T helper type 2-stimulating phenotype and promotes tumour cell migration by dendritic cell-derived osteopontin.
Renal Carcinoma and Angiogenesis: Therapeutic Target and Biomarkers of Response in Current Therapies.
Hepatitis C virus drives the pathogenesis of hepatocellular carcinoma: from immune evasion to carcinogenesis.
Antitumour and immune-adjuvant activities of protein-tyrosine kinase inhibitors.
Novel cancer therapies: treatments driven by tumour biology.
5th European conference on Progress in Vaccination Against Cancer. 20-21 September 2005, Athens, Greece.
Evolution of Cancer Vaccines-Challenges, Achievements, and Future Directions.
Impaired CD95 expression predisposes for recurrence in curatively resected colon carcinoma: clinical evidence for immunoselection and CD95L mediated control of minimal residual disease.
Galectin-1 as a potential cancer target.
Transforming growth factor $\beta$ as regulator of cancer stemness and metastasis.
An overview on the methods of determining the activity of Indoleamine 2, 3-Dioxygenase 1.
Platelet CLEC-2 and podoplanin in cancer metastasis.
Harnessing the immune system in glioblastoma.
Cell-programmed nutrient partitioning in the tumour microenvironment.

Immunological battlefield in gastric cancer and role of immunotherapies.
New approaches to targeting the bone marrow microenvironment in multiple myeloma.
Clinical impact of herpesvirus entry mediator expression in human hepatocellular carcinoma.
Supernatants from lymphocytes stimulated with Bacillus Calmette-Guerin can modify the antigenicity of tumours and stimulate allogeneic T-cell responses.
The classical Hodgkin's lymphoma microenvironment and its role in promoting tumour growth and immune escape.
Combinatorial Immunotherapies for Metastatic Colorectal Cancer.
CD24 signalling through macrophage Siglec-10 is a target for cancer immunotherapy.
Specific cannabinoids revive adaptive immunity by reversing immune evasion mechanisms in metastatic tumours.
The sweet and sour of cancer: glycans as novel therapeutic targets.
Expression and DNA methylation of TNF, IFNG and FOXP3 in colorectal cancer and their prognostic significance.
Plasmacytoid dendritic cells, a role in neoplastic prevention and progression.
The Cancer Stem Cell Niche in Ovarian Cancer and Its Impact on Immune Surveillance.
Immune checkpoint molecule herpes virus entry mediator is overexpressed and associated with poor prognosis in human glioblastoma.
The great escape: tumour cell plasticity in resistance to targeted therapy.
Targeting Aberrant Sialylation to Treat Cancer.
Hypoxia and its impact on the tumour microenvironment of gastroesophageal cancers.
[Circulating Myeloid Suppressor Cells and Their Role in Tumour Immunology].
Review. Colon cancer vaccines: an update.
The cyclooxygenase-2/prostaglandin E(2) pathway and its role in the pathogenesis of human and dog hematological malignancies.
Immunology and immunotherapy approaches for prostate cancer.
MAPK signalling-induced phosphorylation and subcellular translocation of PDHE1 $\alpha$ promotes tumour immune evasion.
Peripheral blood leucocytes show differential expression of tumour progression-related genes in colorectal cancer patients who have a postoperative intra-abdominal infection: a prospective matched cohort study.
TrkB-Target Galectin-1 Impairs Immune Activation and Radiation Responses in Neuroblastoma: Implications for Tumour Therapy.
Therapeutically exploiting STAT3 activity in cancer - using tissue repair as a road map.
Extracellular Vesicles in the Tumour Microenvironment: Eclectic Supervisors.
Adenovirus subversion of immune surveillance, apoptotic and growth regulatory pathways: a model for tumorigenesis.
Competitive glucose metabolism as a target to boost bladder cancer immunotherapy.
Stereotactic ablative body radiotherapy combined with immunotherapy: present status and future perspectives.
The tumour microenvironment of the upper and lower gastrointestinal tract differentially influences dendritic cell maturation.
Local irradiation of murine melanoma affects the development of tumour-specific immunity.
Immune crosstalk in cancer progression and metastatic spread: a complex conversation.
Revisiting cancer immunoediting by understanding cancer immune complexity.
In vivo CRISPR screening identifies Ptpn2 as a cancer immunotherapy target.

The contribution of tumour-derived exosomes to the hallmarks of cancer.
Discovery and saturation analysis of cancer genes across 21 tumour types.
Ferroptosis at the crossroads of cancer-acquired drug resistance and immune evasion.
Mechanisms Of Hypoxia-Induced Immune Escape In Cancer And Their Regulation By Nitric Oxide.
Immunosuppressive effect of bone marrow-derived mesenchymal stem cells in inflammatory microenvironment favours the growth of B16 melanoma cells.
Reversal of gastrointestinal carcinoma-induced immunosuppression and induction of antitumoural immunity by a combination of cyclophosphamide and gene transfer of IL-12.
m(6)A Regulator-Mediated Methylation Modification Patterns and Characterisation of Tumour Microenvironment Infiltration in Non-Small Cell Lung Cancer.
Unraveling tumour microenvironment heterogeneity in nasopharyngeal carcinoma identifies biologically distinct immune subtypes predicting prognosis and immunotherapy responses.
Preclinical intravital microscopy of the tumour-stroma interface: invasion, metastasis, and therapy response.
Genetic dysregulation of gene coding tumor necrosis factor alpha receptors (TNFalpha Rs) in colorectal cancer cells.
Neoantigen quality predicts immunoediting in survivors of pancreatic cancer.
Involvement of Kynurenine Pathway in Hepatocellular Carcinoma.
Tumour-associated CD204(+) microglia/macrophages accumulate in perivascular and perinecrotic niches and correlate with an interleukin-6-enriched inflammatory profile in glioblastoma.
Immune Checkpoint Inhibitors and the Kidney: A Focus on Diagnosis and Management for Personalised Medicine.
Model-based rational design of an oncolytic virus with improved therapeutic potential.
p53, cancer and the immune response.
CD73 in small extracellular vesicles derived from HNSCC defines tumour-associated immunosuppression mediated by macrophages in the microenvironment.
Elucidating tumour-associated microglia/macrophage diversity along glioblastoma progression and under ACOD1 deficiency.
TRAIL receptor-induced features of epithelial-to-mesenchymal transition increase tumour phenotypic heterogeneity: potential cell survival mechanisms.
Epigenetic DNA-methylation regulation of genes coding for lipid raft-associated components: a role for raft proteins in cell transformation and cancer progression (review).
Tumour DDR1 promotes collagen fibre alignment to instigate immune exclusion.
STAT3 and NF-κB are Simultaneously Suppressed in Dendritic Cells in Lung Cancer.
Expression of RCAS1 protein in microglia/macrophages accompanying brain tumours. An immunofluorescence study.
Tumour-infiltrating lymphocytes in melanoma prognosis and cancer immunotherapy.
Colorectal Cancer Stem Cells Fuse with Monocytes to Form Tumour Hybrid Cells with the Ability to Migrate and Evade the Immune System.
Glycan targeting nanoparticle for photodynamic immunotherapy of melanoma.
Gene Expression Analysis of Immune Regulatory Genes in Circulating Tumour Cells and Peripheral Blood Mononuclear Cells in Patients with Colorectal Carcinoma.
Therapy resistance: opportunities created by adaptive responses to targeted therapies in cancer.
Bone marrow stroma protects myeloma cells from cytotoxic damage via induction of the oncoprotein MUC1.
Galectin-1 as a potential therapeutic target in autoimmune disorders and cancer.
Cryoablation and immunotherapy: an overview of evidence on its synergy.

Overcoming TGFβ-mediated immune evasion in cancer.
Therapeutic vaccination in patients with gastrointestinal malignancies. A review of immunological and clinical results.
The RANK-RANKL axis: an opportunity for drug repurposing in cancer?
Anti-tumour immunotherapy with Vγ9Vδ2 T lymphocytes: from the bench to the bedside.
High endothelial venules are associated with microsatellite instability, hereditary background and immune evasion in colorectal cancer.
Aneuploidy as a promoter and suppressor of malignant growth.
Emerging role of PTEN loss in evasion of the immune response to tumours.
Bacteria in cancer initiation, promotion and progression.
Endocytosis in cancer and cancer therapy.
Inactivation of DNA repair triggers neoantigen generation and impairs tumour growth.
Overexpression of B7-H1 correlates with malignant cell proliferation in pancreatic cancer.
Immune landscape, evolution, hypoxia-mediated viral mimicry pathways and therapeutic potential in molecular subtypes of pancreatic neuroendocrine tumours.
ST6GAL1-mediated aberrant sialylation promotes prostate cancer progression.
cGAS-STING signalling in cancer: striking a balance with chromosomal instability.
E3 ubiquitin ligase HECTD2 mediates melanoma progression and immune evasion.
DNA methylation loss promotes immune evasion of tumours with high mutation and copy number load.
DKK1 drives immune suppressive phenotypes in intrahepatic cholangiocarcinoma and can be targeted with anti-DKK1 therapeutic DKN-01.
Immuno-genomic characterisation of high-grade serous ovarian cancer reveals immune evasion mechanisms and identifies an immunological subtype with a favourable prognosis and improved therapeutic efficacy.
Regulative Roles of Metabolic Plasticity Caused by Mitochondrial Oxidative Phosphorylation and Glycolysis on the Initiation and Progression of Tumorigenesis.
Upregulation of GALNT7 in prostate cancer modifies O-glycosylation and promotes tumour growth.
Cancer vaccines.
Immune evasion mechanisms in colorectal cancer liver metastasis patients vaccinated with TroVax (MVA-5T4).
Epigenetic silencing by SETDB1 suppresses tumour intrinsic immunogenicity.
The metabolic milieu in melanoma: Role of immune suppression by CD73/adenosine.
A novel cell-based screen identifies chemical entities that reverse the immune-escape phenotype of metastatic tumours.
STAT proteins as novel targets for cancer drug discovery.
Melanoma-derived gangliosides impair migratory and antigen-presenting function of human epidermal Langerhans cells and induce their apoptosis.
Reviving the Autopsy for Modern Cancer Evolution Research.
Hypoxia-induced EMAP-II transcription in colorectal cancer.
The hallmarks of cancer: relevance to the pathogenesis of polycystic kidney disease.
TGF-beta signalling and immunity in prostate tumourigenesis.
EMAP-II-dependent lymphocyte killing is associated with hypoxia in colorectal cancer.
EMT: A mechanism for escape from EGFR-targeted therapy in lung cancer.
Galectins as pivotal components in oncogenesis and immune exclusion in human malignancies.

Discordant prognosis of mismatch repair deficiency in colorectal and endometrial cancer reflects variation in antitumour immune response and immune escape.
Interferon $\alpha$ in cancer immunoediting: From elimination to escape.
Demonstration of immune responses against devil facial tumour disease in wild Tasmanian devils.
The immune contexture in cancer prognosis and treatment.
Mesenchymal plasticity of devil facial tumour cells during in vivo vaccine and immunotherapy trials.
Mitochondria autophagy: a potential target for cancer therapy.
Targeting TBK1 to overcome resistance to cancer immunotherapy.
Gangliosides inhibit the development from monocytes to dendritic cells.
Metastatic melanoma cells escape from immunosurveillance through the novel mechanism of releasing nitric oxide to induce dysfunction of immunocytes.
Endothelin B receptor promotes the proliferation and immune escape of malignant gliomas.
Dendritic cells in autoimmune disorders and cancer of the thyroid.
Dysregulation at multiple points of the kynurenine pathway is a ubiquitous feature of renal cancer: implications for tumour immune evasion.
Add fuel to the fire: Inflammation and immune response in lung cancer combined with COVID-19.
Aberrant Sialylation in Cancer: Therapeutic Opportunities.
The roles and implications of RNA m(6)A modification in cancer.
A concerted HIF-1 $\alpha$ /MT1-MMP signalling axis regulates the expression of the 3BP2 adaptor protein in hypoxic mesenchymal stromal cells.
The c-MYC oncoprotein as a treatment target in cancer and other disorders of cell growth.
Tryptophan metabolism and disposition in cancer biology and immunotherapy.
G-protein-coupled receptors and cancer.
Hypoxia enhances the malignant nature of bladder cancer cells and concomitantly antagonizes protein O-glycosylation extension.
Advances in the Study of Antitumour Immunotherapy for Newcastle Disease Virus.
Mannosylated liposomes improve therapeutic effects of paclitaxel in colon cancer models.
Upregulation of tumour associated antigen RCAS1 is implicated in high stages of colorectal cancer.
The future of biology in driving the field of hyperthermia.
Significance of KDM6A mutation in bladder cancer immune escape.
Expression profiling of genes regulated by TGF- $\beta$ : differential regulation in normal and tumour cells.
Alternative RNA splicing in tumour heterogeneity, plasticity and therapy.
Identification of thrombin-like activity in ovarian cancer associated ascites and modulation of multiple cytokine networks.
Cytokine patterns in patients with cancer: a systematic review.
Artificial intelligence and digital pathology: Opportunities and implications for immuno-oncology.
An immune edited tumour versus a tumour edited immune system: Prospects for immune therapy of acute myeloid leukaemia.
Predicting clinical benefit of immunotherapy by antigenic or functional mutations affecting tumour immunogenicity.
The intracellular uptake of CD95 modified paclitaxel-loaded poly(lactic-co-glycolic acid) microparticles.
RNA splicing dysregulation and the hallmarks of cancer.
Dendritic cell and macrophage infiltration in microsatellite-unstable and microsatellite-stable colorectal cancer.

Immune modulatory effects of oncogenic KRAS in cancer.
Allele-specific recognition by LILRB3 and LILRA6 of a cytokeratin 8-associated ligand on necrotic glandular epithelial cells.
Extreme intratumour heterogeneity and driver evolution in mismatch repair deficient gastro-oesophageal cancer.
A tale of two tumours: comparison of the immune escape strategies of contagious cancers.
Complex Immune Contextures Characterise Malignant Peritoneal Mesothelioma: Loss of Adaptive Immunological Signature in the More Aggressive Histological Types.
The MYC oncogene - the grand orchestrator of cancer growth and immune evasion.
Role of histone methyltransferase SETDB1 in regulation of tumourigenesis and immune response.
Cancer-cell-derived GABA promotes $\beta$ -catenin-mediated tumour growth and immunosuppression.
The autophagic network and cancer.
Erratum: A novel cell-based screen identifies chemical entities that reverse the immune-escape phenotype of metastatic tumours.
Emerging growth factor receptor antagonists for ovarian cancer treatment.
Serum levels of intercellular adhesion molecule 1 (ICAM-1) in patients with colorectal cancer: inhibitory effect on cytotoxicity.
Generation of Non-Nucleotide CD73 Inhibitors Using a Molecular Docking and 3D-QSAR Approach.
FUT8 Alpha-(1,6)-Fucosyltransferase in Cancer.
P53 and cancer-associated sialylated glycans are surrogate markers of cancerization of the bladder associated with Schistosoma haematobium infection.
Novel microenvironment-based classification of intrahepatic cholangiocarcinoma with therapeutic implications.
Curcuphenol possesses an unusual histone deacetylase enhancing activity that counters immune escape in metastatic tumours.
HNRNPL induced circFAM13B increased bladder cancer immunotherapy sensitivity via inhibiting glycolysis through IGF2BP1/PKM2 pathway.
Hallmarks of alternative splicing in cancer.
Understanding the impact of immune-mediated selection on lung cancer evolution.
Targeting Mesothelin in Solid Tumours: Anti-mesothelin Antibody and Drug Conjugates.
Between the Devil and the Deep Blue Sea: Non-Coding RNAs Associated with Transmissible Cancers in Tasmanian Devil, Domestic Dog and Bivalves.
First-in-human phase 1 dose-escalation study of CAN04, a first-in-class interleukin-1 receptor accessory protein (IL1RAP) antibody in patients with solid tumours.
High CCL27 immunoreactivity in 'supratumoral' epidermis correlates with better prognosis in patients with cutaneous malignant melanoma.
Clonal architecture in mesothelioma is prognostic and shapes the tumour microenvironment.
[Immunologic and tumor biology aspects of placentation in humans].
The role of physics in multiomics and cancer evolution.
Challenges of Neoantigen Targeting in Lynch Syndrome and Constitutional Mismatch Repair Deficiency Syndrome.
The ExPeCT (Examining Exercise, Prostate Cancer and Circulating Tumour Cells) trial: study protocol for a randomised controlled trial.
Transplantation tolerance: the big picture. Where do we stand, where should we go?
The imbalance between Survivin and Bim mediates tumour growth and correlates with poor survival in patients with multiple myeloma.
Cloak and dagger in the avoidance of immune surveillance.

Porphyromonas gingivalis and digestive system cancers.
The NFκB Signaling Pathway in Papillomavirus-induced Lesions: Friend or Foe?
Post-Translational Modification of ZEB Family Members in Cancer Progression.
Filling the gap between chemical carcinogenesis and the hallmarks of cancer: A temporal perspective.
Cancer/testis antigens, gametogenesis and cancer.
Annexin-1 is an oncogene in glioblastoma and causes tumour immune escape through the indirect upregulation of interleukin-8.
The cancer which survived: insights from the genome of an 11000 year-old cancer.
Complementary roles of EP2 and EP4 receptors in malignant glioma.
MAPK inhibitors dynamically affect melanoma release of immune NKG2D-ligands, as soluble protein and extracellular vesicle-associated.
Impact of cancer evolution on immune surveillance and checkpoint inhibitor response.
Human Galectin-1 and Its Inhibitors: Privileged Target for Cancer and HIV.
Fulminant myocarditis caused by immune checkpoint inhibitor: a case report and possible treatment inspiration.
STAT3 and its targeting inhibitors in osteosarcoma.
Two of a kind: transmissible Schwann cell cancers in the endangered Tasmanian devil (Sarcophilus harrisii).
CD47 promotes the proliferation and migration of adamantinomatous craniopharyngioma cells by activating the MAPK/ERK pathway, and CD47 blockade facilitates microglia-mediated phagocytosis. [PIBF - Progesterone-Induced Blocking Factor].
The effects of RNA editing in cancer tissue at different stages in carcinogenesis.
KRAS mutation: from undruggable to druggable in cancer.
S100 proteins in cancer.
Overview of serpin B9 and its roles in cancer (Review).

Table A2-42, Cluster 41

Cluster 41 focuses on immune checkpoint inhibitors for cancer immunotherapy, including mechanisms of resistance to immune checkpoint blockade (273)
Mechanism and potential predictive biomarkers of immune checkpoint inhibitors in NSCLC.
Immune Checkpoint Inhibitors in 10 Years: Contribution of Basic Research and Clinical Application in Cancer Immunotherapy.
Current Clinical Applications and Future Perspectives of Immune Checkpoint Inhibitors in Non-Hodgkin Lymphoma.
Immune-Checkpoint Blockade Therapy in Lymphoma.
Programmed death 1 immune checkpoint inhibitors.
Clinical impact of checkpoint inhibitors as novel cancer therapies.
Immunotherapy against endocrine malignancies: immune checkpoint inhibitors lead the way.
Immune checkpoint blockade therapy for cancer: An overview of FDA-approved immune checkpoint inhibitors.
Immune Checkpoint Inhibitors in Melanoma and HIV Infection.
Immune checkpoint inhibition in ovarian cancer.
The role of PD-1/PD-L1 and application of immune-checkpoint inhibitors in human cancers.
Progress of immune checkpoint therapy in the clinic (Review).
Modern Aspects of Immunotherapy with Checkpoint Inhibitors in Melanoma.

Current Progress and Future Perspectives of Immune Checkpoint in Cancer and Infectious Diseases.
Immune Checkpoint Inhibitors: Recent Clinical Advances and Future Prospects.
Mechanisms of Resistance to Immune Checkpoint Blockade.
Immune Checkpoint Inhibitors: A Promising Choice for Endometrial Cancer Patients?
Predictors of Response, Progression-Free Survival, and Overall Survival in Patients With Lung Cancer Treated With Immune Checkpoint Inhibitors.
Genomics- and Transcriptomics-Based Patient Selection for Cancer Treatment With Immune Checkpoint Inhibitors: A Review.
Recent advances in the clinical development of immune checkpoint blockade therapy.
The emerging role of immune checkpoint inhibition in malignant lymphoma.
Beyond CTLA-4 and PD-1 Inhibition: Novel Immune Checkpoint Molecules for Melanoma Treatment.
Mechanisms underlying response and resistance to immune checkpoint blockade in cancer immunotherapy.
Immune checkpoint-based therapy in myeloid malignancies: a promise yet to be fulfilled.
Immune checkpoint inhibitors for the treatment of melanoma.
Combining Immune Checkpoint Inhibitors: Established and Emerging Targets and Strategies to Improve Outcomes in Melanoma.
Current Advances in Checkpoint Inhibitors: Lessons from Non-Central Nervous System Cancers and Potential for Glioblastoma.
Immune Checkpoint Inhibitors in HBV-Caused Hepatocellular Carcinoma Therapy.
Immune checkpoint blockade opens an avenue of cancer immunotherapy with a potent clinical efficacy.
Immune Checkpoint Blockade for the Treatment of Hodgkin Lymphoma.
Targets of Immune Escape Mechanisms in Cancer: Basis for Development and Evolution of Cancer Immune Checkpoint Inhibitors.
Checkpoint inhibitors in hematological malignancies.
Programmed death-1 immune checkpoint blockade in the treatment of hematological malignancies.
Immune checkpoint inhibitors in clinical trials.
Mechanisms of immune-related adverse events during the treatment of cancer with immune checkpoint inhibitors.
Targeting the PD-1 pathway: a promising future for the treatment of melanoma.
Pivotal role of PD-1/PD-L1 immune checkpoints in immune escape and cancer progression: Their interplay with platelets and FOXP3+Tregs related molecules, clinical implications and combinational potential with phytochemicals.
Intellectual property issues of immune checkpoint inhibitors.
Immune checkpoint inhibitors and cellular treatment for lymphoma immunotherapy.
Next steps in immuno-oncology: enhancing antitumor effects through appropriate patient selection and rationally designed combination strategies.
Identifying and treating candidates for checkpoint inhibitor therapies in multiple myeloma and lymphoma.
Novel Immune Checkpoints in Esophageal Cancer: From Biomarkers to Therapeutic Targets.
Cardiovascular Complications Associated With Novel Cancer Immunotherapies.
Cancer Immunotherapy by Blocking Immune Checkpoints on Innate Lymphocytes.
Monoclonal Antibodies for the Treatment of Cancer.
[Clinical Diagnosis and Treatment Recommendations for Cardiac Adverse Reactions Related to Immune Checkpoint Inhibitor].



Strategies to Improve the Antitumor Effect of Immunotherapy for Hepatocellular Carcinoma.
Identification of responders to immune checkpoint therapy: which biomarkers have the highest value?
Next generation of immune checkpoint therapy in cancer: new developments and challenges.
Histone Deacetylase Inhibitors to Overcome Resistance to Targeted and Immuno Therapy in Metastatic Melanoma.
Research Progresses in Immunological Checkpoint Inhibitors for Breast Cancer Immunotherapy.
Next generation of immune checkpoint inhibitors and beyond.
Immunomodulatory Properties of Immune Checkpoint Inhibitors-More than Boosting T-Cell Responses?
Checkpoint Inhibition in Hodgkin Lymphoma - a Review.
LAG-3 Inhibitors: Novel Immune Checkpoint Inhibitors Changing the Landscape of Immunotherapy.
Checkpoint Inhibition: Programmed Cell Death 1 and Programmed Cell Death 1 Ligand Inhibitors in Hodgkin Lymphoma.
PD-1/PD-L1 checkpoint inhibitors in advanced hepatocellular carcinoma immunotherapy.
Immune checkpoint inhibition and its relationship with hypermutation phenotype as a potential treatment for Glioblastoma.
[IMMUNOTHERAPY WITH CHECKPOINT INHIBITORS (ICPI) AND IMMUNE RELATED ADVERSE EVENTS (IRAE'S)].
[Clinical Diagnosis and Treatment of Immune-related Adverse Events in Digestive System Related to Immune Checkpoint Inhibitors].
Mechanisms of resistance to immune checkpoint inhibitors in melanoma: What we have to overcome?
Immune Checkpoint Inhibitor Therapy in Colorectal Cancer-The Role of Cellular Pathology.
Immune Checkpoint Inhibitors in Thoracic Malignancies: Review of the Existing Evidence by an IASLC Expert Panel and Recommendations.
Immune checkpoint inhibitors in melanoma provide the cornerstones for curative therapies.
Immune-Related Adverse Events Associated With Immune Checkpoint Inhibitor Therapy.
Checkpoint Inhibitors and Their Application in Breast Cancer.
The blockade of immune checkpoints in cancer immunotherapy.
Immune Checkpoint Inhibitors to Treat Malignant Lymphomas.
Indicators of responsiveness to immune checkpoint inhibitors.
Tumor matrix remodeling and novel immunotherapies: the promise of matrix-derived immune biomarkers.
Checkpoint blockade in lymphoma.
Resistance of melanoma to immune checkpoint inhibitors is overcome by targeting the sphingosine kinase-1.
Emerging role of immune checkpoint inhibitors and predictive biomarkers in head and neck cancers.
The emerging role of immunotherapy in advanced urothelial cancers.
Immune checkpoint inhibitors in advanced and recurrent/metastatic cervical cancer.
Programmed death-1 checkpoint blockade in acute myeloid leukemia.
PD-1/PD-L1 immune checkpoint blockade-based combinational treatment: Immunotherapeutic amplification strategies against colorectal cancer.
Understanding and overcoming the resistance of cancer to PD-1/PD-L1 blockade.
Immune checkpoint inhibitors break whose heart? Perspectives from cardio-immuno-oncology.
Emerging drugs targeting PD-1 and PD-L1: reality or hope?

Application of PD-1 Blockade in Cancer Immunotherapy.
Novel immunotherapy approaches for metastatic urothelial and renal cell carcinoma.
Immune checkpoint blockade therapy: the 2014 Tang Prize in Biopharmaceutical Science.
Cutaneous manifestations associated with immune checkpoint inhibitors.
Immune evasion mechanisms in acute myeloid leukemia: A focus on immune checkpoint pathways.
The Evolving Role of Immune Checkpoint Inhibitors in Hepatocellular Carcinoma Treatment.
The Role of the Lymphocyte Functional Crosstalk and Regulation in the Context of Checkpoint Inhibitor Treatment-Review.
[The "immune checkpoints", how does it work].
Immune Dysfunction in Non-Hodgkin Lymphoma: Avenues for New Immunotherapy-Based Strategies.
On the Horizon: Targeting Next-Generation Immune Checkpoints for Cancer Treatment.
Modulating the wayward T cell: New horizons with immune checkpoint inhibitor treatments in autoimmunity, transplant, and cancer.
Recent pharmacological approaches for the treatment of renal cell carcinoma.
Progress of immune checkpoint LAG-3 in immunotherapy.
Immune checkpoint inhibition in early-stage triple-negative breast cancer.
Advances in T-cell checkpoint immunotherapy for head and neck squamous cell carcinoma.
Molecular Farming of Pembrolizumab and Nivolumab.
Novel immunologic approaches in lymphoma: unleashing the brakes on the immune system.
Immune Checkpoint Inhibitors and RAS-ERK Pathway-Targeted Drugs as Combined Therapy for the Treatment of Melanoma.
Checkpoint Proteins in Pediatric Brain and Extracranial Solid Tumors: Opportunities for Immunotherapy.
Nivolumab: A Review in Advanced Melanoma.
Current landscape and future directions of biomarkers for predicting responses to immune checkpoint inhibitors.
Successful treatment of severe immune checkpoint inhibitor associated autoimmune hepatitis with basiliximab: a case report.
PD-1/PD-L1 Checkpoint Inhibitors in Tumor Immunotherapy.
Current status and future potential of predictive biomarkers for immune checkpoint inhibitors in gastric cancer.
Optimizing checkpoint inhibitors therapy for relapsed or progressive classic Hodgkin lymphoma by multiplex immunohistochemistry of the tumor microenvironment.
Current Understanding of the Mechanisms Underlying Immune Evasion From PD-1/PD-L1 Immune Checkpoint Blockade in Head and Neck Cancer.
Response to Immune Checkpoint Inhibitors Is Affected by Deregulations in the Antigen Presentation Machinery: A Systematic Review and Meta-Analysis.
Immune Checkpoint Inhibition in Classical Hodgkin Lymphoma: From Early Achievements towards New Perspectives.
Current Approaches of Immune Checkpoint Therapy in Chronic Lymphocytic Leukemia.
Harnessing the immune system through programmed death-1 blockade in the management of Hodgkin lymphoma.
Inhibition of immune checkpoints PD-1, CTLA-4, and IDO1 coordinately induces immune-mediated liver injury in mice.
Checkpoint Inhibition in Hodgkin Lymphoma: Saving the Best for Last?

Immune Checkpoint Inhibitors and Beyond: An Overview of Immune-Based Therapies in Merkel Cell Carcinoma.
[New Treatment Options for Renal Cell Carcinoma - Efficacy and Safety of Immune Checkpoint Inhibitors].
Novel Immunotherapy Combinations.
Therapeutic Advances and Treatment Options in Metastatic Melanoma.
Fibrinogen-like protein 1 (FGL1): the next immune checkpoint target.
Adverse Events of Concurrent Immune Checkpoint Inhibitors and Antiangiogenic Agents: A Systematic Review.
Simultaneous inhibition of PD-1 and LAG-3: the future of immunotherapy?
Role of Clinical Pharmacology in the Development and Approval of Immunotherapies Targeting Immune Checkpoints.
Immune checkpoint inhibitors reverse tolerogenic mechanisms induced by melanoma targeted radionuclide therapy.
Checkpoint-inhibition in ovarian cancer: rising star or just a dream?
Immunomodulatory Drugs: Immune Checkpoint Agents in Acute Leukemia.
Immune checkpoint blockade: a common denominator approach to cancer therapy.
Autoimmune rhabdomyolysis and a multiorgan display of PD-1 inhibitor induced immune related adverse events during treatment of metastatic melanoma.
Endocrine adverse events related with immune checkpoint inhibitors: an update for clinicians.
Immunotherapy and chimeric antigen receptor T-cell therapy in hepatocellular carcinoma.
Pathophysiology of Immune Checkpoint Inhibitor-Induced Myocarditis.
Immuno-Oncology: Emerging Targets and Combination Therapies.
Immune Checkpoint Modulators: An Emerging Antiglioma Armamentarium.
Pembrolizumab: PD-1 inhibition as a therapeutic strategy in cancer.
The performance and perspectives of dendritic cell vaccines modified by immune checkpoint inhibitors or stimulants.
Immune checkpoint inhibitors and endocrine side effects, a narrative review.
Pembrolizumab and its role in relapsed/refractory classical Hodgkin's lymphoma: evidence to date and clinical utility.
Anti-programmed death-1 and anti-programmed death-ligand 1 antibodies in cancer therapy.
Systematic Review of Immune Checkpoint Inhibition in Urological Cancers.
The Role of Immune Checkpoint Inhibitors in Classical Hodgkin Lymphoma.
Pembrolizumab in classical Hodgkin's lymphoma.
Immune Checkpoints in Leprosy: Immunotherapy As a Feasible Approach to Control Disease Progression.
[Advances in the study of programmed cell death protein 1 and its ligand inhibitors in the treatment of late stage HCC].
[PD-1 blockade therapy in lymphoid malignancies].
Checkpoint Inhibition in Myeloma: Opportunities and Challenges.
Checkpoint Inhibition and Infectious Diseases: A Good Thing?
LAG3-PD-1 Combo Overcome the Disadvantage of Drug Resistance.
Immune Checkpoint Inhibition in Acute Myeloid Leukemia and Myelodysplastic Syndromes.
Immunotherapy of Hodgkin Lymphoma: Mobilizing the Patient's Immune Response.
Atezolizumab-Induced Bell's Palsy in a Patient With Small Cell Lung Cancer.
Programmed death-1 & its ligands: promising targets for cancer immunotherapy.

Identification of MicroRNA-mRNA Networks in Melanoma and Their Association with PD-1 Checkpoint Blockade Outcomes.
Multinational Association of Supportive Care in Cancer (MASCC) 2020 clinical practice recommendations for the management of immune-related adverse events: pulmonary toxicity.
Integrin signaling gene alterations and outcomes of cancer patients receiving immune checkpoint inhibitors.
Emerging principles of brain immunology and immune checkpoint blockade in brain metastases.
A Cancer Cell Program Promotes T Cell Exclusion and Resistance to Checkpoint Blockade.
Functions of Immune Checkpoint Molecules Beyond Immune Evasion.
Role of CD155/TIGIT in Digestive Cancers: Promising Cancer Target for Immunotherapy.
An updated review of gastrointestinal toxicity induced by PD-1 inhibitors: from mechanisms to management.
Tankyrase inhibition sensitizes melanoma to PD-1 immune checkpoint blockade in syngeneic mouse models.
Immune checkpoints and reproductive immunology: Pioneers in the future therapy of infertility related Disorders?
High PD-1/PD-L1 Checkpoint Interaction Infers Tumor Selection and Therapeutic Sensitivity to Anti-PD-1/PD-L1 Treatment.
Immune checkpoint inhibitors in malignant lymphoma: Advances and perspectives.
Immune checkpoint blockade in human cancer therapy: lung cancer and hematologic malignancies.
Genetic and Epigenetic Biomarkers of Immune Checkpoint Blockade Response.
Soluble B7-CD28 Family Inhibitory Immune Checkpoint Proteins and Anti-Cancer Immunotherapy.
Deep immune profiling reveals targetable mechanisms of immune evasion in immune checkpoint inhibitor-refractory glioblastoma.
Resistance to anti-PD-1-based immunotherapy in basal cell carcinoma: a case report and review of the literature.
Therapeutic targeting of PD-1/PD-L1 blockade by novel small-molecule inhibitors recruits cytotoxic T cells into solid tumor microenvironment.
Immune Checkpoint Inhibitors in AML-A New Frontier.
Clinical utility of nivolumab in the treatment of advanced melanoma.
Checkpoint inhibition: new treatment options in urologic cancer.
Bypassing anti-PD-(L)1 therapy: Mechanisms and management strategies.
Immune Checkpoint Molecules in Reproductive Immunology.
Harnessing the Power of Onco-Immunotherapy with Checkpoint Inhibitors.
Targeting Histone 3 Variants Epigenetic Landscape and Inhibitory Immune Checkpoints: An Option for Paediatric Brain Tumours Therapy.
Inflammatory Myeloradiculitis Secondary to Pembrolizumab: A Case Report and Literature Review.
Pharmacological combination of nivolumab with dendritic cell vaccines in cancer immunotherapy: An overview.
Epigenetic abnormalities of classical Hodgkin lymphoma and its effect on immune escape.
Myeloid-derived suppressor cells and their role in CTLA-4 blockade therapy.
Management of Adverse Events in Cancer Patients Treated With PD-1/PD-L1 Blockade: Focus on Asian Populations.
Rheumatic complications of checkpoint inhibitors: Lessons from autoimmunity.
Advanced Melanoma: Current Treatment Options, Biomarkers, and Future Perspectives.

Successful Use of Immunotherapy in a Patient with Metastatic Squamous Cell Lung Cancer and Underlying Autoimmune Disease.
Pembrolizumab: a novel antiprogrammed death 1 (PD-1) monoclonal antibody for treatment of metastatic melanoma.
The Role of Immune Checkpoint Molecules for Relapse After Allogeneic Hematopoietic Cell Transplantation.
Inhibitory Effect of PD-1/PD-L1 and Blockade Immunotherapy in Leukemia.
TIGIT and CD96: new checkpoint receptor targets for cancer immunotherapy.
Nivolumab in renal cell carcinoma.
Harnessing the power of the immune system via blockade of PD-1 and PD-L1: a promising new anticancer strategy.
Anticoagulation with Factor Xa Inhibitors Is Associated with Improved Overall Response and Progression-Free Survival in Patients with Metastatic Malignant Melanoma Receiving Immune Checkpoint Inhibitors-A Retrospective, Real-World Cohort Study.
Acute heart failure due to autoimmune myocarditis under pembrolizumab treatment for metastatic melanoma.
Atezolizumab for the treatment of colorectal cancer: the latest evidence and clinical potential.
A multicenter phase 2 single arm study of cabozantinib in patients with advanced or unresectable renal cell carcinoma pre-treated with one immune-checkpoint inhibitor: The BREAKPOINT trial (Meet-Uro trial 03).
Tumor-Specific Delivery of Immune Checkpoint Inhibitors by Engineered AAV Vectors.
Making urothelial carcinomas less immune to immunotherapy.
Hyperprogressive Disease In a Metastatic Renal Cell Carcinoma Patient After Receiving Immune Checkpoint Inhibitors: A Case Report.
PD-L1 and immune escape: insights from melanoma and other lineage-unrelated malignancies.
Epi-drugs in combination with immunotherapy: a new avenue to improve anticancer efficacy.
Clinical and molecular characteristics associated with response to therapeutic PD-1/PD-L1 inhibition in advanced Merkel cell carcinoma.
Pembrolizumab-Induced Psoriasis in Metastatic Melanoma: Activity and Safety of Apremilast, a Case Report.
Characterization of the impact of immune checkpoint inhibitors on platelet activation and aggregation.
Immunotherapy in ovarian cancer: fake news or the real deal?
Nivolumab in the Treatment of Refractory Pediatric Hodgkin Lymphoma.
Immune Checkpoint Blockade and Hematopoietic Stem Cell Transplant.
The role of CTLA-4 and PD-1 in anti-tumor immune response and their potential efficacy against osteosarcoma.
The Where, the When, and the How of Immune Monitoring for Cancer Immunotherapies in the Era of Checkpoint Inhibition.
From biomarkers to therapeutic targets: the promise of PD-L1 in thyroid autoimmunity and cancer.
The effects of PD-1/PD-L1 checkpoint inhibitors on recurrent/metastatic head and neck squamous cell carcinoma: a critical review of the literature and meta-analysis.
Discovery of acquired molecular signature on immune checkpoint inhibitors in paired tumor tissues.
Prognostic role of soluble PD-1 and BTN2A1 in overweight melanoma patients treated with nivolumab or pembrolizumab: finding the missing links in the symbiotic immune-metabolic interplay.
Tumor immune checkpoints and their associated inhibitors.

Results of clinical trials with anti-programmed death 1/programmed death ligand 1 inhibitors in lung cancer.
Microsatellite Instability Predicts Response to Anti-PD1 Immunotherapy in Metastatic Melanoma.
Immune checkpoint molecules in acute myeloid leukaemia: managing the double-edged sword.
Microsatellite instability is a biomarker for immune checkpoint inhibitors in endometrial cancer.
Post-hematopoietic stem cell transplantation relapse: Role of checkpoint inhibitors.
Checkpoint inhibitors in the treatment of cutaneous malignant melanoma.
TGFβ2 Induces the Soluble Isoform of CTLA-4 - Implications for CTLA-4 Based Checkpoint Inhibitor Antibodies in Malignant Melanoma.
Emerging targets in cancer immunotherapy: beyond CTLA-4 and PD-1.
Advances in Hodgkin's lymphoma pharmacotherapy: a focus on histone deacetylase inhibitors.
Dietary Considerations for Inflammatory Bowel Disease Are Useful for Treatment of Checkpoint Inhibitor-Induced Colitis.
Nivolumab in the treatment of advanced renal cell carcinoma: clinical trial evidence and experience.
PD-1-siRNA delivered by attenuated Salmonella enhances the antimelanoma effect of pimozide.
Co-inhibitory immune checkpoints in head and neck squamous cell carcinoma.
[Consideration on the Research and Development of Anti-tumor Bispecific Antibody Drugs].
Nivolumab for previously treated unresectable metastatic anal cancer (NCI9673): a multicentre, single-arm, phase 2 study.
Programmed Cell Death-One Inhibition Therapy in Classical Hodgkin Lymphoma.
Serum cell division cycle 42 in advanced hepatocellular carcinoma patients: Linkage with clinical characteristics and immune checkpoint inhibitor-related treatment outcomes.
Novel immunotherapeutic drugs for the treatment of lung cancer.
Immunotherapy and Targeted Therapy in the Management of Oral Cancers.
Efficacy and safety of nivolumab in Japanese patients with previously untreated advanced melanoma: A phase II study.
Avelumab: A Novel Anti-PD-L1 Agent in the Treatment of Merkel Cell Carcinoma and Urothelial Cell Carcinoma.
A Novel CD73 Inhibitor SHR170008 Suppresses Adenosine in Tumor and Enhances Anti-Tumor Activity with PD-1 Blockade in a Mouse Model of Breast Cancer.
Novel Immunologic Approaches to Melanoma Treatment.
Pembrolizumab for the treatment of Hodgkin Lymphoma.
Immunotherapy in MSI/dMMR tumors in the perioperative setting: The IMHOTEP trial.
Investigational PD-1 inhibitors in HL and NHL and biomarkers for predictors of response and outcome.
Au-CGKRK Nanoconjugates for Combating Cancer through T-Cell-Driven Therapeutic RNA Interference.
NLRP3 promotes immune escape by regulating immune checkpoints: A pan-cancer analysis.
Retrospective study of nivolumab for patients with recurrent high grade gliomas.
Safety and Efficacy of Pembrolizumab Prior to Allogeneic Stem Cell Transplantation for Acute Myelogenous Leukemia.
T lymphocyte membrane-decorated epigenetic nanoinducer of interferons for cancer immunotherapy.
Oncolytic vaccinia virus reinvigorates peritoneal immunity and cooperates with immune checkpoint inhibitor to suppress peritoneal carcinomatosis in colon cancer.
Tackling the dysregulated immune-checkpoints in classical Hodgkin lymphoma: bidirectional regulations between the microenvironment and Hodgkin/Reed-Sternberg cells.

Efficacy and Safety of PD-1/PD-L1 Inhibitor as Single-Agent Immunotherapy in Endometrial Cancer: A Systematic Review and Meta-Analysis.
Earlier-Phased Cancer Immunity Cycle Strongly Influences Cancer Immunity in Operable Never-Smoker Lung Adenocarcinoma.
Serum cell division cycle 42 reflects the treatment response and survival in patients with advanced cervical cancer who receive immune checkpoint inhibitor treatment.
CSF-1R inhibitor, pexidartinib, sensitizes esophageal adenocarcinoma to PD-1 immune checkpoint blockade in a rat model.
Down-regulation of A20 promotes immune escape of lung adenocarcinomas.
Single-cell atlases link macrophages and CD8(+) T-cell subpopulations to disease progression and immunotherapy response in urothelial carcinoma.
Comparative Analysis of Immune Checkpoint Molecules and Their Potential Role in the Transmissible Tasmanian Devil Facial Tumor Disease.
Galectin-1-driven T cell exclusion in the tumor endothelium promotes immunotherapy resistance.
Aptamer-Based Immunotheranostic Strategies.
PD-1/PD-L1 immune checkpoints: Tumor vs atherosclerotic progression.
A20 promotes colorectal cancer immune evasion by upregulating STC1 expression to block "eat-me" signal.
Metastatic Merkel cell carcinoma response to nivolumab.
Immunotherapy for recurrent/metastatic head and neck cancer.
First-line pembrolizumab versus dabrafenib/trametinib treatment for BRAF V600-mutant advanced melanoma.
Roles for the FCRL6 Immunoreceptor in Tumor Immunology.
Molecular Pathways: Targeting B7-H3 (CD276) for Human Cancer Immunotherapy.
A phase 2 study of tremelimumab in patients with advanced uveal melanoma.
Introduction of novel agents in the treatment of primary CNS lymphoma.
MYC Overexpression Drives Immune Evasion in Hepatocellular Carcinoma That Is Reversible through Restoration of Proinflammatory Macrophages.
The role of IFN- $\gamma$ -signalling in response to immune checkpoint blockade therapy.
Predictive value of co-expression patterns of immune checkpoint molecules for clinical outcomes of hematological malignancies.
Clinical Relevance of the serum CTLA-4 in Cats with Mammary Carcinoma.
Anti-CTLA-4 based therapy elicits humoral immunity to galectin-3 in patients with metastatic melanoma.
Biology and management of primary effusion lymphoma.
Early low blood MALT1 expression levels forecast better efficacy of PD-1 inhibitor-based treatment in patients with metastatic colorectal cancer.
Pretreatment serum VEGF is associated with clinical response and overall survival in advanced melanoma patients treated with ipilimumab.
Vertical level of blood cell division cycle 42 predicts response and survival benefits to PD-1 inhibitor-based regimen in metastatic colorectal cancer patients.
Multimodal pooled Perturb-CITE-seq screens in patient models define mechanisms of cancer immune evasion.
Urothelial Carcinoma In Situ of the Bladder: Correlation of CK20 Expression With Adaptive Immune Resistance, Response to BCG Therapy, and Clinical Outcome.
Suppression of RAC1-driven malignant melanoma by group A PAK inhibitors.

Molecular profiling reveals immunogenic cues in anaplastic large cell lymphomas with DUSP22 rearrangements.

Table A2-43, Cluster 42

Cluster 42 focuses on Porphyromonas gingivalis primarily and Pseudomonas aeruginosa secondarily, emphasizing their immune escape and evasion mechanisms (233)
Porphyromonas gingivalis survival skills: Immune evasion.
Roles of Porphyromonas gingivalis and its virulence factors in periodontitis.
Molecular Strategies Underlying Porphyromonas gingivalis Virulence.
Intracellular localization of Porphyromonas gingivalis thiol proteinase in periodontal tissues of chronic periodontitis patients.
Genetic diversity in the oral pathogen Porphyromonas gingivalis: molecular mechanisms and biological consequences.
Macrophages: A communication network linking Porphyromonas gingivalis infection and associated systemic diseases.
Porphyromonas gingivalis regulates atherosclerosis through an immune pathway.
Possible mechanisms of Pseudomonas aeruginosa-associated lung disease.
Persistent Exposure to Porphyromonas gingivalis Promotes Proliferative and Invasion Capabilities, and Tumorigenic Properties of Human Immortalized Oral Epithelial Cells.
Porphyromonas gingivalis Stimulates TLR2-PI3K Signaling to Escape Immune Clearance and Induce Bone Resorption Independently of MyD88.
Pseudomonas aeruginosa: acute lung injury or ventilator-associated pneumonia?
Porphyromonas gingivalis suppresses adaptive immunity in periodontitis, atherosclerosis, and Alzheimer's disease.
Virulence Factors of Pseudomonas Aeruginosa and Antivirulence Strategies to Combat Its Drug Resistance.
Porphyromonas gingivalis adopts intricate and unique molecular mechanisms to survive and persist within the host: a critical update.
Synthetic host defense peptide IDR-1002 reduces inflammation in Pseudomonas aeruginosa lung infection.
Dampening Host Sensing and Avoiding Recognition in Pseudomonas aeruginosa Pneumonia.
Local and Systemic Effects of Porphyromonas gingivalis Infection.
Serratiopeptidase Affects the Physiology of Pseudomonas aeruginosa Isolates from Cystic Fibrosis Patients.
A Macrophage Subversion Factor Is Shared by Intracellular and Extracellular Pathogens.
The salicylidene acylhydrazide INP0341 attenuates Pseudomonas aeruginosa virulence in vitro and in vivo.
Differential capacity for complement receptor-mediated immune evasion by Porphyromonas gingivalis depending on the type of innate leukocyte.
Natural competence is a major mechanism for horizontal DNA transfer in the oral pathogen Porphyromonas gingivalis.
Porphyromonas gingivalis Outer Membrane Vesicles Induce Selective Tumor Necrosis Factor Tolerance in a Toll-Like Receptor 4- and mTOR-Dependent Manner.
Subversion of innate immunity by periodontopathic bacteria via exploitation of complement receptor-3.



Pseudomonas aeruginosa in Chronic Lung Infections: How to Adapt Within the Host?
Pseudomonas aeruginosa hijacks the murine nitric oxide metabolic pathway to evade killing by neutrophils in the lung.
Immune evasion strategies of Porphyromonas gingivalis.
Porphyromonas gingivalis downregulates the immune response of fibroblasts.
Sialidase facilitates Porphyromonas gingivalis immune evasion by reducing M1 polarization, antigen presentation, and phagocytosis of infected macrophages.
Epstein-Barr Virus-Oral Bacterial Link in the Development of Oral Squamous Cell Carcinoma.
Hyperlipidemia impaired innate immune response to periodontal pathogen porphyromonas gingivalis in apolipoprotein E knockout mice.
Genetic characteristics and pathogenic mechanisms of periodontal pathogens.
Immune escape strategies of Pseudomonas aeruginosa to establish chronic infection.
Role of Porphyromonas gingivalis HmuY in Immunopathogenesis of Chronic Periodontitis.
Porphyromonas gingivalis and the autophagic pathway: an innate immune interaction?
Porphyromonas gingivalis induction of TLR2 association with Vinculin enables PI3K activation and immune evasion.
Flagellar motility is a key determinant of the magnitude of the inflammasome response to Pseudomonas aeruginosa.
OmpA-like proteins of Porphyromonas gingivalis contribute to serum resistance and prevent Toll-like receptor 4-mediated host cell activation.
Pseudomonas aeruginosa exploits lipid A and muropeptides modification as a strategy to lower innate immunity during cystic fibrosis lung infection.
Proteolytic activity of Porphyromonas gingivalis attenuates MCP-1 mRNA expression in LPS-stimulated THP-1 cells.
Role of nucleotide-binding oligomerization domain-like receptor family pyrin domain containing 6 in activation of inflammation in human umbilical vein endothelial cells stimulated by Porphyromonas gingivalis-an in vitro study.
Porphyromonas gingivalis interactions with complement receptor 3 (CR3): innate immunity or immune evasion?
Microbiota, Immune Subversion, and Chronic Inflammation.
Identification of PGN_1123 as the Gene Encoding Lipid A Deacylase, an Enzyme Required for Toll-Like Receptor 4 Evasion, in Porphyromonas gingivalis.
Virulence Factors of the Periodontal Pathogens: Tools to Evade the Host Immune Response and Promote Carcinogenesis.
Pouring salt on a wound: Pseudomonas aeruginosa virulence factors alter Na <sup>+</sup> and Cl <sup>-</sup> flux in the lung.
The C5a receptor impairs IL-12-dependent clearance of Porphyromonas gingivalis and is required for induction of periodontal bone loss.
Mechanism and implications of CXCR4-mediated integrin activation by Porphyromonas gingivalis.
Fimbrial proteins of porphyromonas gingivalis mediate in vivo virulence and exploit TLR2 and complement receptor 3 to persist in macrophages.
Pseudomonas aeruginosa evasion of phagocytosis is mediated by loss of swimming motility and is independent of flagellum expression.
Loss of lipopolysaccharide receptor CD14 from the surface of human macrophage-like cells mediated by Porphyromonas gingivalis outer membrane vesicles.
Extracellular Vesicles from Pseudomonas aeruginosa Suppress MHC-Related Molecules in Human Lung Macrophages.
Pseudomonas aeruginosa invades corneal epithelial cells during experimental infection.

Pseudomonas aeruginosa: the potential to immunise against infection.
Recognition of the carbohydrate modifications to the RgpA protease of Porphyromonas gingivalis by periodontal patient serum IgG.
P. aeruginosa type III and type VI secretion systems modulate early response gene expression in type II pneumocytes in vitro.
Metapopulation Structure of CRISPR-Cas Immunity in Pseudomonas aeruginosa and Its Viruses.
The capsule of Porphyromonas gingivalis reduces the immune response of human gingival fibroblasts.
The capsule of Porphyromonas gingivalis leads to a reduction in the host inflammatory response, evasion of phagocytosis, and increase in virulence.
Pseudomonas aeruginosa Airway Infection Recruits and Modulates Neutrophilic Myeloid-Derived Suppressor Cells.
Purinergic signaling during Porphyromonas gingivalis infection.
Pseudomonas aeruginosa biofilms: mechanisms of immune evasion.
Exposure to Bile Leads to the Emergence of Adaptive Signaling Variants in the Opportunistic Pathogen Pseudomonas aeruginosa.
Human Toll-like receptor 4 responses to P. gingivalis are regulated by lipid A 1- and 4'-phosphatase activities.
Internalization of Pseudomonas aeruginosa Strain PAO1 into Epithelial Cells Is Promoted by Interaction of a T6SS Effector with the Microtubule Network.
Comparative Analysis of Peptidoglycans From Pseudomonas aeruginosa Isolates Recovered From Chronic and Acute Infections.
Porphyromonas gingivalis evasion of autophagy and intracellular killing by human myeloid dendritic cells involves DC-SIGN-TLR2 crosstalk.
The subversion of the immune system by Francisella tularensis.
The Pseudomonas aeruginosa Type III secretion system plays a dual role in the regulation of caspase-1 mediated IL-1 $\beta$ maturation.
The Microbial Endocrinology of Pseudomonas aeruginosa: Inflammatory and Immune Perspectives.
Pseudomonas aeruginosa ExoT inhibits in vitro lung epithelial wound repair.
Pseudomonas aeruginosa Infections in Cancer Patients.
The Pseudomonas aeruginosa Lectin LecB Causes Integrin Internalization and Inhibits Epithelial Wound Healing.
Inactivation of tumor necrosis factor- $\alpha$ by proteinases (gingipains) from the periodontal pathogen, Porphyromonas gingivalis. Implications of immune evasion.
Granulocytic Myeloid-Derived Suppressor Cells in Cystic Fibrosis.
A Library of Promoter-gfp Fusion Reporters for Studying Systematic Expression Pattern of Cyclic-di-GMP Metabolism-Related Genes in Pseudomonas aeruginosa.
Pseudomonas aeruginosa proteolytically alters the interleukin 22-dependent lung mucosal defense.
Uncovering the components of the Francisella tularensis virulence stealth strategy.
Genome-Wide Survey of Pseudomonas aeruginosa PA14 Reveals a Role for the Glyoxylate Pathway and Extracellular Proteases in the Utilization of Mucin.
NOD1 and NOD2 mediate sensing of periodontal pathogens.
Modulation of inflammasome activity by Porphyromonas gingivalis in periodontitis and associated systemic diseases.
Biology and pathogenesis of cytomegalovirus in periodontal disease.
Type III Secretion Protein, PcrV, Impairs Pseudomonas aeruginosa Biofilm Formation by Increasing M1 Macrophage-Mediated Anti-bacterial Activities.

Porphyromonas gingivalis attenuates ATP-mediated inflammasome activation and HMGB1 release through expression of a nucleoside-diphosphate kinase.
Evaluating Bacterial Pathogenesis Using a Model of Human Airway Organoids Infected with Pseudomonas aeruginosa Biofilms.
Host-pathogen interactions and immune evasion strategies in Francisella tularensis pathogenicity.
Type IV pilus glycosylation mediates resistance of Pseudomonas aeruginosa to opsonic activities of the pulmonary surfactant protein A.
Identification of Small Molecules Blocking the Pseudomonas aeruginosa type III Secretion System Protein PcrV.
The human pathogen Pseudomonas aeruginosa utilizes conserved virulence pathways to infect the social amoeba Dictyostelium discoideum.
Kdo hydrolase is required for Francisella tularensis virulence and evasion of TLR2-mediated innate immunity.
Structure-Based Design of $\alpha$ -Substituted Mercaptoacetamides as Inhibitors of the Virulence Factor LasB from Pseudomonas aeruginosa.
From the Outside-In: The Francisella tularensis Envelope and Virulence.
Cutting edge: mutation of Francisella tularensis mviN leads to increased macrophage absent in melanoma 2 inflammasome activation and a loss of virulence.
The lipid A phosphate position determines differential host Toll-like receptor 4 responses to phylogenetically related symbiotic and pathogenic bacteria.
Tripartite interactions between filamentous Pf4 bacteriophage, Pseudomonas aeruginosa, and bacterivorous nematodes.
Cyclic-di-GMP regulates lipopolysaccharide modification and contributes to Pseudomonas aeruginosa immune evasion.
Pathogen induction of CXCR4/TLR2 cross-talk impairs host defense function.
Genome-wide patterns of recombination in the opportunistic human pathogen Pseudomonas aeruginosa.
Spheres of influence: Porphyromonas gingivalis outer membrane vesicles.
O-Specific Antigen-Dependent Surface Hydrophobicity Mediates Aggregate Assembly Type in Pseudomonas aeruginosa.
Antioxidant Defenses of Francisella tularensis Modulate Macrophage Function and Production of Proinflammatory Cytokines.
Opportunistic infections in lung disease: Pseudomonas infections in cystic fibrosis.
Multifaceted effects of Francisella tularensis on human neutrophil function and lifespan.
Gene Loss and Acquisition in Lineages of Pseudomonas aeruginosa Evolving in Cystic Fibrosis Patient Airways.
[Immune evasion of Pseudomonas aeruginosa].
The LasB Elastase of Pseudomonas aeruginosa Acts in Concert with Alkaline Protease AprA To Prevent Flagellin-Mediated Immune Recognition.
Pseudomonas aeruginosa invasion of and multiplication within corneal epithelial cells in vitro.
Targeting of a Fixed Bacterial Immunogen to Fc Receptors Reverses the Anti-Inflammatory Properties of the Gram-Negative Bacterium, Francisella tularensis, during the Early Stages of Infection.
A Secreted Bacterial Peptidylarginine Deiminase Can Neutralize Human Innate Immune Defenses.
Identification of a novel Francisella tularensis factor required for intramacrophage survival and subversion of innate immune response.
Bacterial Nucleotidyl Cyclase Inhibits the Host Innate Immune Response by Suppressing TAK1 Activation.

Francisella tularensis inhibits the intrinsic and extrinsic pathways to delay constitutive apoptosis and prolong human neutrophil lifespan.
Pseudomonas aeruginosa quorum sensing modulates immune responses: An updated review article.
How has neutrophil research improved our understanding of periodontal pathogenesis?
Scavenging of reactive oxygen species by tryptophan metabolites helps Pseudomonas aeruginosa escape neutrophil killing.
Cleavage of IgG1 and IgG3 by gingipain K from Porphyromonas gingivalis may compromise host defense in progressive periodontitis.
Pseudomonas aeruginosa MucD protease mediates keratitis by inhibiting neutrophil recruitment and promoting bacterial survival.
Complement Receptor 3-Mediated Inhibition of Inflammasome Priming by Ras GTPase-Activating Protein During Francisella tularensis Phagocytosis by Human Mononuclear Phagocytes.
Persistent infection with Pseudomonas aeruginosa in ventilator-associated pneumonia.
Identification of a potent inhibitor of type II secretion system from Pseudomonas aeruginosa.
The Mechanistic Pathways of Periodontal Pathogens Entering the Brain: The Potential Role of Treponema denticola in Tracing Alzheimer's Disease Pathology.
Diguanylate Cyclases and Phosphodiesterases Required for Basal-Level c-di-GMP in Pseudomonas aeruginosa as Revealed by Systematic Phylogenetic and Transcriptomic Analyses.
The orange spotted cockroach (Blattella germanica) is a permissive experimental host for Francisella tularensis.
Innate immune response to oral bacteria and the immune evasive characteristics of periodontal pathogens.
Francisella tularensis SchuS4 and SchuS4 lipids inhibit IL-12p40 in primary human dendritic cells by inhibition of IRF1 and IRF8.
The link between periodontal disease and rheumatoid arthritis: an updated review.
Proteome-wide mapping and reverse vaccinology-based B and T cell multi-epitope subunit vaccine designing for immune response reinforcement against Porphyromonas gingivalis.
Pseudomonas aeruginosa quorum-sensing metabolite induces host immune cell death through cell surface lipid domain dissolution.
Lipids derived from virulent Francisella tularensis broadly inhibit pulmonary inflammation via toll-like receptor 2 and peroxisome proliferator-activated receptor $\alpha$ .
A Francisella tularensis locus required for spermine responsiveness is necessary for virulence.
Specific interactions between the alkaline protease of P. aeruginosa and its natural peptide inhibitor: ab initio molecular simulations.
Bacterial degradation of immunoglobulin A1 in relation to periodontal diseases.
The presence of CD14 overcomes evasion of innate immune responses by virulent Francisella tularensis in human dendritic cells in vitro and pulmonary cells in vivo.
Roles of extracellular vesicles in periodontal homeostasis and their therapeutic potential.
Deletion of ripA alleviates suppression of the inflammasome and MAPK by Francisella tularensis.
Qualitative and Quantitative Determination of Quorum Sensing Inhibition In Vitro.
Soluble lytic transglycosylase SLT of Francisella novicida is involved in intracellular growth and immune suppression.
Janus kinase 3 activity is necessary for phosphorylation of cytosolic phospholipase A2 and prostaglandin E2 synthesis by macrophages infected with Francisella tularensis live vaccine strain.
Fine tuning inflammation at the front door: macrophage complement receptor 3-mediate phagocytosis and immune suppression for Francisella tularensis.

Temporal Manipulation of Mitochondrial Function by Virulent <i>Francisella tularensis</i> To Limit Inflammation and Control Cell Death.
Characterization of <i>Francisella tularensis</i> outer membrane proteins.
<i>Pseudomonas aeruginosa</i> and <i>Klebsiella pneumoniae</i> Adaptation to Innate Immune Clearance Mechanisms in the Lung.
N-alkylimidazole derivatives as potential inhibitors of quorum sensing in <i>Pseudomonas aeruginosa</i> .
Neutrophils in the periodontium: Interactions with pathogens and roles in tissue homeostasis and inflammation.
HigB of <i>Pseudomonas aeruginosa</i> Enhances Killing of Phagocytes by Up-Regulating the Type III Secretion System in Ciprofloxacin Induced Persister Cells.
Human neutrophils and oral microbiota: a constant tug-of-war between a harmonious and a discordant coexistence.
Clonal Clusters, Molecular Resistance Mechanisms and Virulence Factors of Gram-Negative Bacteria Isolated from Chronic Wounds in Ghana.
Virulence mechanism of bacteria in mixed infection: attenuation of cytokine levels and evasion of polymorphonuclear leukocyte phagocytosis.
Indoleamine 2,3-dioxygenase 1 is a lung-specific innate immune defense mechanism that inhibits growth of <i>Francisella tularensis</i> tryptophan auxotrophs.
Virulent <i>Francisella tularensis</i> destabilize host mRNA to rapidly suppress inflammation.
<i>Francisella tularensis</i> -infected macrophages release prostaglandin E2 that blocks T cell proliferation and promotes a Th2-like response.
Neutrophils: potential therapeutic targets in tularemia?
Virulence factors of the oral spirochete <i>Treponema denticola</i> .
The Cyclic AMP-Vfr Signaling Pathway in <i>Pseudomonas aeruginosa</i> Is Inhibited by Cyclic Di-GMP.
MyD88-dependent signaling prolongs survival and reduces bacterial burden during pulmonary infection with virulent <i>Francisella tularensis</i> .
<i>Francisella tularensis</i> antioxidants harness reactive oxygen species to restrict macrophage signaling and cytokine production.
Effective, broad spectrum control of virulent bacterial infections using cationic DNA liposome complexes combined with bacterial antigens.
Development and bioanalytical method validation of an LC-MS/MS assay for simultaneous quantitation of 2-alkyl-4(1H)-quinolones for application in bacterial cell culture and lung tissue.
Correlation of phenotypic and genotypic virulence markers, antimicrobial susceptibility pattern, and outcome of <i>Pseudomonas aeruginosa</i> sepsis infection.
Rapid and efficient inactivation of IL-6 gingipains, lysine- and arginine-specific proteinases from <i>Porphyromonas gingivalis</i> .
O-Polysaccharide Plays a Major Role on the Virulence and Immunostimulatory Potential of <i>Aggregatibacter actinomycetemcomitans</i> During Periodontal Infection.
Proteolysis of CD14 on human gingival fibroblasts by arginine-specific cysteine proteinases from <i>Porphyromonas gingivalis</i> leading to down-regulation of lipopolysaccharide-induced interleukin-8 production.
Exposure to extracellular vesicles from <i>Pseudomonas aeruginosa</i> result in loss of DNA methylation at enhancer and DNase hypersensitive site regions in lung macrophages.
Evasion of IFN- $\gamma$ signaling by <i>Francisella novicida</i> is dependent upon <i>Francisella</i> outer membrane protein C.
[Research progress in <i>Filifactor alocis</i> interacting with periodontitis].
Extracellular Bacterial Proteases in Chronic Wounds: A Potential Therapeutic Target?

Host cell-surface proteins as substrates of gingipains, the main proteases of <i>Porphyromonas gingivalis</i> .
<i>Pseudomonas</i> expression of an oxygen sensing prolyl hydroxylase homologue regulates neutrophil host responses in vitro and in vivo.
Method for the isolation of <i>Francisella tularensis</i> outer membranes.
<i>Aggregatibacter actinomycetemcomitans</i> and <i>Filifactor alocis</i> : Two exotoxin-producing oral pathogens.
Garlic as an inhibitor of <i>Pseudomonas aeruginosa</i> quorum sensing in cystic fibrosis--a pilot randomized controlled trial.
Oral infection of mice with <i>Fusobacterium nucleatum</i> results in macrophage recruitment to the dental pulp and bone resorption.
Calcium Regulates the Activity and Structural Stability of Tpr, a Bacterial Calpain-like Peptidase.
Characterization of the <i>Treponema denticola</i> Virulence Factor Dentilisin.
In-silico structural characterization and phylogenetic analysis of Nucleoside diphosphate kinase: A novel antiapoptotic protein of <i>Porphyromonas gingivalis</i> .
Insights into the Evolution of <i>P. aeruginosa</i> Antimicrobial Resistance in a Patient Undergoing Intensive Therapy.
The intracellular pathogen <i>Francisella tularensis</i> escapes from adaptive immunity by metabolic adaptation.
Infection with <i>Francisella tularensis</i> LVS clpB leads to an altered yet protective immune response.
New insights into the emerging role of oral spirochaetes in periodontal disease.
Extracellular deoxyribonuclease production by periodontal bacteria.
A novel role for plasmin-mediated degradation of opsonizing antibody in the evasion of host immunity by virulent, but not attenuated, <i>Francisella tularensis</i> .
Proteolysis of ICAM-1 on human oral epithelial cells by gingipains.
Immune Response Modulation by <i>Pseudomonas aeruginosa</i> Persister Cells.
The role of the <i>Pseudomonas aeruginosa</i> hypermutator phenotype on the shift from acute to chronic virulence during respiratory infection.
Human Single-Chain Antibodies That Neutralize Elastolytic Activity of <i>Pseudomonas aeruginosa</i> LasB.
Intracellular localisation and innate immune responses following <i>Francisella noatunensis</i> infection of Atlantic cod ( <i>Gadus morhua</i> ) macrophages.
Comparison of the virulence of exopolysaccharide-producing <i>Prevotella intermedia</i> to exopolysaccharide non-producing periodontopathic organisms.
Remodeling of Lipid A in <i>Pseudomonas syringae</i> pv. <i>phaseolicola</i> In Vitro.
<i>Pseudomonas aeruginosa</i> promotes autophagy to suppress macrophage-mediated bacterial eradication.
<i>Aggregatibacter actinomycetemcomitans</i> leukotoxin: from threat to therapy.
Exploring structural features and potential lipid interactions of <i>Pseudomonas aeruginosa</i> type three secretion effector PemB by spectroscopic and calorimetric experiments.
A <i>Pseudomonas aeruginosa</i> TIR effector mediates immune evasion by targeting UBAP1 and TLR adaptors.
Subversion of innate immune responses by <i>Francisella</i> involves the disruption of TRAF3 and TRAF6 signalling complexes.
Therapeutic possibilities of ceftazidime nanoparticles in devastating <i>pseudomonas</i> ophthalmic infections; keratitis and endophthalmitis.
Self-assembled hybrid supraparticles that proteolytically degrade tumor necrosis factor- $\alpha$ .
Unique <i>Francisella</i> Phosphatidylethanolamine Acts as a Potent Anti-Inflammatory Lipid.

Metabolic labeling to characterize the overall composition of Francisella lipid A and LPS grown in broth and in human phagocytes.
Intracellular survival and innate immune evasion of Burkholderia cepacia: Improved understanding of quorum sensing-controlled virulence factors, biofilm, and inhibitors.
Adaptation by Ancient Horizontal Acquisition of Butyrate Metabolism Genes in Aggregatibacter actinomycetemcomitans.
Proteomics of Aggregatibacter actinomycetemcomitans Outer Membrane Vesicles.
The bacterial pigment pyocyanin inhibits the NLRP3 inflammasome through intracellular reactive oxygen and nitrogen species.
Infection of human gingival fibroblasts with Aggregatibacter actinomycetemcomitans: An in vitro study.
Insecticidal activity of the metalloprotease AprA occurs through suppression of host cellular and humoral immunity.
Transcriptome Profiling of Wild-Type and pga-Knockout Mutant Strains Reveal the Role of Exopolysaccharide in Aggregatibacter actinomycetemcomitans.
A galU mutant of Francisella tularensis is attenuated for virulence in a murine pulmonary model of tularemia.
The role of NLRP3 in regulating gingival epithelial cell responses evoked by Aggregatibacter actinomycetemcomitans.
Repression of aerobic leukotoxin transcription by integration host factor in Aggregatibacter actinomycetemcomitans.
The Msp Protein of Treponema denticola Interrupts Activity of Phosphoinositide Processing in Neutrophils.
Pf Filamentous Phage Requires UvrD for Replication in Pseudomonas aeruginosa.
Expression, purification and crystallization of the phosphate-binding PstS protein from Pseudomonas aeruginosa.
Proteolytic inactivation of LL-37 by karilysin, a novel virulence mechanism of Tannerella forsythia.
Sialic Acid-Siglec-E Interactions During Pseudomonas aeruginosa Infection of Macrophages Interferes With Phagosome Maturation by Altering Intracellular Calcium Concentrations.
Aryl hydrocarbon receptor is required for immune response in Epinephelus coioides and Danio rerio infected by Pseudomonas plecoglossicida.
Proteolytic cleavage of the long pentraxin PTX3 in the airways of cystic fibrosis patients.
Anti-inflammatory potential of pyocyanin in LPS-stimulated murine macrophages.
Francisella tularensis elicits IL-10 via a PGE <sub>2</sub> -inducible factor, to drive macrophage MARCH1 expression and class II down-regulation.
Identification of a Fusobacterium nucleatum 65 kDa serine protease.
Anaerobic regulation of Actinobacillus actinomycetemcomitans leukotoxin transcription is ArcA/FnrA-independent and requires a novel promoter element.
Systematic investigation on the turning point of over-inflammation to immunosuppression in CLP mice model and their characteristics.
The regulation of leukotoxin production in Actinobacillus actinomycetemcomitans strain JP2.
The Zinc Nutritional Immunity of Epinephelus coioides Contributes to the Importance of znuC During Pseudomonas plecoglossicida Infection.
Filifactor alocis modulates human neutrophil antimicrobial functional responses.
Aggregatibacter actinomycetemcomitans Outer Membrane Proteins 29 and 29 Parologue Induce Evasion of Immune Response.

Mlc is a transcriptional activator with a key role in integrating cyclic AMP receptor protein and integration host factor regulation of leukotoxin RNA synthesis in <i>Aggregatibacter actinomycetemcomitans</i> .
An unbroken network of interactions connecting flagellin domains is required for motility in viscous environments.
Sialoglycoproteins adsorbed by <i>Pseudomonas aeruginosa</i> facilitate their survival by impeding neutrophil extracellular trap through siglec-9.
<i>Pseudomonas syringae</i> evades host immunity by degrading flagellin monomers with alkaline protease AprA.
Potential of the <i>Tannerella forsythia</i> S-layer to delay the immune response.
Exploiting type 3 complement receptor for TNF- $\alpha$ suppression, immune evasion, and progressive pulmonary fungal infection.
Gangliosides block <i>Aggregatibacter Actinomycetemcomitans</i> leukotoxin (LtxA)-mediated hemolysis.
ATP Induces IL-1 $\beta$ Secretion in <i>Neisseria gonorrhoeae</i> -Infected Human Macrophages by a Mechanism Not Related to the NLRP3/ASC/Caspase-1 Axis.
Albumin grafted coaxial electrosprayed polycaprolactone-zinc oxide nanoparticle for sustained release and activity enhanced antibacterial drug delivery.
Gonococcal nitric oxide reductase is encoded by a single gene, <i>norB</i> , which is required for anaerobic growth and is induced by nitric oxide.

Table A2-44, Cluster 43

Cluster 43 focuses on HIV escape mutations, emphasizing the role of the HIV-1 accessory protein Nef in viral pathogenesis, viral replication, and immune escape of HIV-infected cells (553)
Consequences of HLA-B*13-Associated Escape Mutations on HIV-1 Replication and Nef Function.
Adaptation of HIV-1 to human leukocyte antigen class I.
HIV-specific CD8+ T cells: serial killers condemned to die?
Translation of HLA-HIV associations to the cellular level: HIV adapts to inflate CD8 T cell responses against Nef and HLA-adapted variant epitopes.
Modelling the spread of HIV immune escape mutants in a vaccinated population.
Role of HIV-specific CD8(+) T cells in pediatric HIV cure strategies after widespread early viral escape.
Virological and immunological correlates of HIV post-treatment control after temporal antiretroviral therapy during acute HIV infection.
Relative Resistance of HLA-B to Downregulation by Naturally Occurring HIV-1 Nef Sequences.
Temporal loss of Nef-epitope CTL recognition following macaque lipopeptide immunization and SIV challenge.
Microdevices for examining immunological responses of single cells to HIV.
Anti-HIV potency of T-cell responses elicited by dendritic cell therapeutic vaccination.
Nef: a pleiotropic modulator of primate lentivirus infectivity and pathogenesis.
HIV: current opinion in escapology.
Evolution of human immunodeficiency virus type 1 cytotoxic T-lymphocyte epitopes: fitness-balanced escape.
Clustering patterns of cytotoxic T-lymphocyte epitopes in human immunodeficiency virus type 1 (HIV-1) proteins reveal imprints of immune evasion on HIV-1 global variation.
Identification of SIV Nef CD8(+) T cell epitopes restricted by a MHC class I haplotype associated with lower viral loads in a macaque AIDS model.



CTL Escape and Viral Fitness in HIV/SIV Infection.
Differential escape patterns within the dominant HLA-B*57:03-restricted HIV Gag epitope reflect distinct clade-specific functional constraints.
Effects of naturally-arising HIV Nef mutations on cytotoxic T lymphocyte recognition and Nef's functionality in primary macrophages.
CTL response to HIV type 1 subtype C is poorly predicted by known epitope motifs.
Trivalent live attenuated influenza-simian immunodeficiency virus vaccines: efficacy and evolution of cytotoxic T lymphocyte escape in macaques.
CD8+ T cell epitope-flanking mutations disrupt proteasomal processing of HIV-1 Nef.
Fitness costs and diversity of the cytotoxic T lymphocyte (CTL) response determine the rate of CTL escape during acute and chronic phases of HIV infection.
HIV-specific T cell responses reflect substantive in vivo interactions with antigen despite long-term therapy.
HLA-associated clinical progression correlates with epitope reversion rates in early human immunodeficiency virus infection.
Precise identification of a human immunodeficiency virus type 1 antigen processing mutant.
Escape is a more common mechanism than avidity reduction for evasion of CD8+ T cell responses in primary human immunodeficiency virus type 1 infection.
HIV escape mutations occur preferentially at HLA-binding sites of CD8 T-cell epitopes.
Cytotoxic T lymphocytes and viral adaptation in HIV infection.
Selection of escape mutation by Pol154-162-specific cytotoxic T cells among chronically HIV-1-infected HLA-B*5401-positive individuals.
HIV evolution: CTL escape mutation and reversion after transmission.
Impact of vaccination on cytotoxic T lymphocyte immunodominance and cooperation against simian immunodeficiency virus replication in rhesus macaques.
Switching and emergence of CTL epitopes in HIV-1 infection.
The HIV hide and seek game: an immunogenomic analysis of the HIV epitope repertoire.
Fitness-Balanced Escape Determines Resolution of Dynamic Founder Virus Escape Processes in HIV-1 Infection.
Molecular Basis for the Recognition of HIV Nef138-8 Epitope by a Pair of Human Public T Cell Receptors.
Evidence of differential HLA class I-mediated viral evolution in functional and accessory/regulatory genes of HIV-1.
HIV reservoir: antiviral immune responses and immune interventions for curing HIV infection.
Escape mutations in HIV infection and its impact on CD8+ T cell responses.
Nef Is Dispensable for Resistance of Simian Immunodeficiency Virus-Infected Macrophages to CD8+ T Cell Killing.
The effects of natural altered peptide ligands on the whole blood cytotoxic T lymphocyte response to human immunodeficiency virus.
Evasion of adaptive immunity by HIV through the action of host APOBEC3G/F enzymes.
Clinical and evolutionary consequences of HIV adaptation to HLA: implications for vaccine and cure.
Maintenance of AP-2-Dependent Functional Activities of Nef Restricts Pathways of Immune Escape from CD8 T Lymphocyte Responses.
Nef-mediated resistance of human immunodeficiency virus type 1 to antiviral cytotoxic T lymphocytes.
Determinant of HIV-1 mutational escape from cytotoxic T lymphocytes.
HIV molecular epidemiology: transmission and adaptation to human populations.

Bioorganic approaches towards HIV vaccine design.
Immune responses induced in HHD mice by multiepitope HIV vaccine based on cryptic epitope modification.
Genotypic and Mechanistic Characterization of Subtype-Specific HIV Adaptation to Host Cellular Immunity.
Potential immune escape mutations under inferred selection pressure in HIV-1 strains circulating in Medellín, Colombia.
Genotypic and functional impact of HIV-1 adaptation to its host population during the North American epidemic.
Variable fitness impact of HIV-1 escape mutations to cytotoxic T lymphocyte (CTL) response.
B-cell responses to HIV infection.
Accumulation of Pol Mutations Selected by HLA-B*52:01-C*12:02 Protective Haplotype-Restricted Cytotoxic T Lymphocytes Causes Low Plasma Viral Load Due to Low Viral Fitness of Mutant Viruses.
Dynamics of viral evolution and CTL responses in HIV-1 infection.
HLA Class-II Associated HIV Polymorphisms Predict Escape from CD4+ T Cell Responses.
HLA class I-driven evolution of human immunodeficiency virus type 1 subtype c proteome: immune escape and viral load.
Antigen specificities of HIV-infected cells: A role in infection and persistence?
Multiple T-cell responses are associated with better control of acute HIV-1 infection: An observational study.
The flexibility of the TCR allows recognition of a large set of naturally occurring epitope variants by HIV-specific cytotoxic T lymphocytes.
Conditional CD8+ T cell escape during acute simian immunodeficiency virus infection.
Adaptation of CD8 T cell responses to changing HIV-1 sequences in a cohort of HIV-1 infected individuals not selected for a certain HLA allele.
HIV-1 Nef blocks transport of MHC class I molecules to the cell surface via a PI 3-kinase-dependent pathway.
Paediatric non-progression following grandmother-to-child HIV transmission.
HIV immune escape at an immunodominant epitope in HLA-B*27-positive individuals predicts viral load outcome.
HIV antibodies for treatment of HIV infection.
Immunology of pediatric HIV infection.
Computational inference methods for selective sweeps arising in acute HIV infection.
HIV control through a single nucleotide on the HLA-B locus.
Longitudinal within-host evolution of HIV Nef-mediated CD4, HLA and SERINC5 downregulation activity: a case study.
Physiological relevance of ACOT8-Nef interaction in HIV infection.
HIV immune evasion disruption of antigen presentation by the HIV Nef protein.
High-avidity, high-IFN $\gamma$ -producing CD8 T-cell responses following immune selection during HIV-1 infection.
Transmission of HIV-1 Gag immune escape mutations is associated with reduced viral load in linked recipients.
Two Distinct Mechanisms Leading to Loss of Virological Control in the Rare Group of Antiretroviral Therapy-Naive, Transiently Aviremic Children Living with HIV.
Transmission of human immunodeficiency virus type 1 from a patient who developed AIDS to an elite suppressor.
Structured observations reveal slow HIV-1 CTL escape.

A longitudinal analysis of immune escapes from HLA-B*13-restricted T-cell responses at early stage of CRF01_AE subtype HIV-1 infection and implications for vaccine design.
Epitope-specific CD8+ T cell kinetics rather than viral variability determine the timing of immune escape in simian immunodeficiency virus infection.
Biphasic CD8+ T-Cell Defense in Simian Immunodeficiency Virus Control by Acute-Phase Passive Neutralizing Antibody Immunization.
Impact of HLA-B*52:01-Driven Escape Mutations on Viral Replicative Capacity.
Selection, transmission, and reversion of an antigen-processing cytotoxic T-lymphocyte escape mutation in human immunodeficiency virus type 1 infection.
How HIV evades CTL recognition.
Live and let die: Nef functions beyond HIV replication.
Modelling the evolution and spread of HIV immune escape mutants.
Transmission and accumulation of CTL escape variants drive negative associations between HIV polymorphisms and HLA.
Potential elucidation of a novel CTL epitope in HIV-1 protease by the protease inhibitor resistance mutation L90M.
Advances in HIV therapeutics and cure strategies: findings obtained through non-human primate studies.
Cytotoxic T lymphocyte lysis inhibited by viable HIV mutants.
Phylogenetic dependency networks: inferring patterns of CTL escape and codon covariation in HIV-1 Gag.
A participant-derived xenograft model of HIV enables long-term evaluation of autologous immunotherapies.
Sequential broadening of CTL responses in early HIV-1 infection is associated with viral escape.
Antiretroviral Drug Discovery Targeting the HIV-1 Nef Virulence Factor.
Immune escape precedes breakthrough human immunodeficiency virus type 1 viremia and broadening of the cytotoxic T-lymphocyte response in an HLA-B27-positive long-term-nonprogressing child.
Lentiviral Nef proteins manipulate T cells in a subset-specific manner.
Differential selection pressure exerted on HIV by CTL targeting identical epitopes but restricted by distinct HLA alleles from the same HLA supertype.
An integrative bioinformatic approach for studying escape mutations in human immunodeficiency virus type 1 gag in the Pumwani Sex Worker Cohort.
Broad CTL Response in Early HIV Infection Drives Multiple Concurrent CTL Escapes.
Enhancement of viral escape in HIV-1 Nef by STEP vaccination.
Viral replication capacity as a correlate of HLA B57/B5801-associated nonprogressive HIV-1 infection.
Compensatory mutation partially restores fitness and delays reversion of escape mutation within the immunodominant HLA-B*5703-restricted Gag epitope in chronic human immunodeficiency virus type 1 infection.
Resistance of HIV-infected cells to cytotoxic T lymphocytes.
Maintenance of Nef-mediated modulation of major histocompatibility complex class I and CD4 after sexual transmission of human immunodeficiency virus type 1.
The differential ability of HLA B*5701+ long-term nonprogressors and progressors to restrict human immunodeficiency virus replication is not caused by loss of recognition of autologous viral gag sequences.
Late seroconversion in HIV-resistant Nairobi prostitutes despite pre-existing HIV-specific CD8+ responses.

The role of recombination for the coevolutionary dynamics of HIV and the immune response.
Functionally impaired HIV-specific CD8 T cells show high affinity TCR-ligand interactions.
The evolutionary significance of certain amino acid substitutions and their consequences for HIV-1 immunogenicity toward HLA's A*0201 and B*27.
Structure, function, and inhibitor targeting of HIV-1 Nef-effector kinase complexes.
Epitope targeting and viral inoculum are determinants of Nef-mediated immune evasion of HIV-1 from cytotoxic T lymphocytes.
Current understanding of HIV-1 and T-cell adaptive immunity: progress to date.
Effects of Mutations on Replicative Fitness and Major Histocompatibility Complex Class I Binding Affinity Are Among the Determinants Underlying Cytotoxic-T-Lymphocyte Escape of HIV-1 Gag Epitopes.
HIV Diversity and Genetic Compartmentalization in Blood and Testes during Suppressive Antiretroviral Therapy.
Dynamics of immune escape during HIV/SIV infection.
HIV escape and attenuation by cytotoxic T lymphocytes.
Triple bypass: complicated paths to HIV escape.
Mucosal immunology of HIV infection.
Dendritic cells focus CTL responses toward highly conserved and topologically important HIV-1 epitopes.
Antigenic oscillations and shifting immunodominance in HIV-1 infections.
High frequency of transmitted HIV-1 Gag HLA class I-driven immune escape variants but minimal immune selection over the first year of clade C infection.
Nef promotes evasion of human immunodeficiency virus type 1-infected cells from the CTLA-4-mediated inhibition of T-cell activation.
HIV evolution in response to HLA-restricted CTL selection pressures: a population-based perspective.
Human immunodeficiency virus type 1 population genetics and adaptation in newly infected individuals.
Weaker HLA Footprints on HIV in the Unique and Highly Genetically Admixed Host Population of Mexico.
Resistance of Major Histocompatibility Complex Class B (MHC-B) to Nef-Mediated Downregulation Relative to that of MHC-A Is Conserved among Primate Lentiviruses and Influences Antiviral T Cell Responses in HIV-1-Infected Individuals.
New approaches to HIV vaccine development.
Replicative capacity of human immunodeficiency virus type 1 transmitted from mother to child is associated with pediatric disease progression rate.
Differences in expression of tumor suppressor, innate immune, inflammasome, and potassium/gap junction channel host genes significantly predict viral reservoir size during treated HIV infection.
Acute phase CD8+ T lymphocytes against alternate reading frame epitopes select for rapid viral escape during SIV infection.
In chronic infection, HIV gag-specific CD4+ T cell receptor diversity is higher than CD8+ T cell receptor diversity and is associated with less HIV quasispecies diversity.
Cytotoxic T-lymphocyte escape mutations identified by HLA association favor those which escape and revert rapidly.
Genetic subunit vaccines : a novel approach for genetic immunization.
Effective downregulation of HLA-A*2 and HLA-B*57 by primary human immunodeficiency virus type 1 isolates cultured from elite suppressors.

Global Database-Driven Assessment of HIV-1 Adaptation to the Immune Repertoires of Human Populations.
Impaired ability of Nef to counteract SERINC5 is associated with reduced plasma viremia in HIV-infected individuals.
Impact of pre-adapted HIV transmission.
Molecular and functional analysis of a conserved CTL epitope in HIV-1 p24 recognized from a long-term nonprogressor: constraints on immune escape associated with targeting a sequence essential for viral replication.
Impact of changes in antigen level on CD38/PD-1 co-expression on HIV-specific CD8 T cells in chronic, untreated HIV-1 infection.
Epitope escape mutation and decay of human immunodeficiency virus type 1-specific CTL responses.
Linking pig-tailed macaque major histocompatibility complex class I haplotypes and cytotoxic T lymphocyte escape mutations in simian immunodeficiency virus infection.
HIV-1 Mutation and Recombination Rates Are Different in Macrophages and T-cells.
An HIV-1 Nef genotype that diminishes immune control mediated by protective human leucocyte antigen alleles.
HLA-associated immune escape pathways in HIV-1 subtype B Gag, Pol and Nef proteins.
Timing of immune escape linked to success or failure of vaccination.
Tactics used by HIV-1 to evade host innate, adaptive, and intrinsic immunities.
Impact of Suboptimal APOBEC3G Neutralization on the Emergence of HIV Drug Resistance in Humanized Mice.
The rate of immune escape vanishes when multiple immune responses control an HIV infection.
New insights into the immunology and evolution of HIV.
HLA-Cw*03-restricted CD8+ T-cell responses targeting the HIV-1 gag major homology region drive virus immune escape and fitness constraints compensated for by intracodon variation.
Sexual transmission of single human immunodeficiency virus type 1 virions encoding highly polymorphic multisite cytotoxic T-lymphocyte escape variants.
Differential patterns of immune escape at Tat-specific cytotoxic T cell epitopes in pigtail macaques.
The T-cell response to HIV.
The enduring tale of T cells in HIV immunopathogenesis.
Role of CD8(+) T Cells in the Selection of HIV-1 Immune Escape Mutations.
Current progress in the development of HIV vaccines.
Mother-to-child transmission of HIV infection and CTL escape through HLA-A2-SLYNTVATL epitope sequence variation.
Immunological Control of HIV-1 Disease Progression by Rare Protective HLA Allele.
Estimating the effectiveness of simian immunodeficiency virus-specific CD8+ T cells from the dynamics of viral immune escape.
Conditional Immune Escape during Chronic Simian Immunodeficiency Virus Infection.
APOBEC3G Regulation of the Evolutionary Race Between Adaptive Immunity and Viral Immune Escape Is Deeply Imprinted in the HIV Genome.
Impaired immune evasion in HIV through intracellular delays and multiple infection of cells.
HIV-1 Nef Antagonizes SERINC5 Restriction by Downregulation of SERINC5 via the Endosome/Lysosome System.
Distribution, persistence, and efficacy of adoptively transferred central and effector memory-derived autologous simian immunodeficiency virus-specific CD8+ T cell clones in rhesus macaques during acute infection.

HLA-B57/B*5801 human immunodeficiency virus type 1 elite controllers select for rare gag variants associated with reduced viral replication capacity and strong cytotoxic T-lymphocyte [corrected] recognition.
The cytotoxic T-lymphocyte response in HIV-1 infection.
Correlates of protective cellular immunity revealed by analysis of population-level immune escape pathways in HIV-1.
Evidence of HIV-1 adaptation to host HLA alleles following chimp-to-human transmission.
Fitness cost of escape mutations in p24 Gag in association with control of human immunodeficiency virus type 1.
HIV-1 Nef sequence and functional compartmentalization in the gut is not due to differential cytotoxic T lymphocyte selective pressure.
Variable processing and cross-presentation of HIV by dendritic cells and macrophages shapes CTL immunodominance and immune escape.
Viral evolution and challenges in the development of HIV vaccines.
Reversion in vivo after inoculation of a molecular proviral DNA clone of simian immunodeficiency virus with a cytotoxic-T-lymphocyte escape mutation.
HIV-1-infected T-cells dynamics and prognosis: An evolutionary game model.
Investigating the Consequences of Interference between Multiple CD8+ T Cell Escape Mutations in Early HIV Infection.
Recent Insights into the HIV/AIDS Pandemic.
Defective pneumococcal-specific Th1 responses in HIV-infected adults precedes a loss of control of pneumococcal colonization.
HLA-driven convergence of HIV-1 viral subtypes B and F toward the adaptation to immune responses in human populations.
HIV Subtype and Nef-Mediated Immune Evasion Function Correlate with Viral Reservoir Size in Early-Treated Individuals.
Rapid viral escape at an immunodominant simian-human immunodeficiency virus cytotoxic T-lymphocyte epitope exacts a dramatic fitness cost.
Frequent detection of escape from cytotoxic T-lymphocyte recognition in perinatal human immunodeficiency virus (HIV) type 1 transmission: the ariel project for the prevention of transmission of HIV from mother to infant.
Impact of HLA in mother and child on disease progression of pediatric human immunodeficiency virus type 1 infection.
Defects in blood dendritic cell subsets in HIV-1 subtype c infected Indians.
Contribution of proteasome-catalyzed peptide cis-splicing to viral targeting by CD8(+) T cells in HIV-1 infection.
Cytomegalovirus and HIV Persistence: Pouring Gas on the Fire.
HIV-1 adaptation to antigen processing results in population-level immune evasion and affects subtype diversification.
Immune activation promotes evolutionary conservation of T-cell epitopes in HIV-1.
Cytotoxic T lymphocytes in asymptomatic long-term nonprogressing HIV-1 infection. Breadth and specificity of the response and relation to in vivo viral quasispecies in a person with prolonged infection and low viral load.
Immunological and virological mechanisms of vaccine-mediated protection against SIV and HIV.
Replication-competent simian immunodeficiency virus (SIV) Gag escape mutations archived in latent reservoirs during antiretroviral treatment of SIV-infected macaques.

Pyrosequencing reveals restricted patterns of CD8+ T cell escape-associated compensatory mutations in simian immunodeficiency virus.
iHIVARNA phase IIa, a randomized, placebo-controlled, double-blinded trial to evaluate the safety and immunogenicity of iHIVARNA-01 in chronically HIV-infected patients under stable combined antiretroviral therapy.
Impact of HLA selection pressure on HIV fitness at a population level in Mexico and Barbados.
HIV evolutionary dynamics within and among hosts.
Pitfalls of Antiretroviral Therapy: Current Status and Long-Term CNS Toxicity.
The Effect of Interference on the CD8(+) T Cell Escape Rates in HIV.
Tight-Binding Hydroxypyrazole HIV-1 Nef Inhibitors Suppress Viral Replication in Donor Mononuclear Cells and Reverse Nef-Mediated MHC-I Downregulation.
Relative expression levels of the HLA class-I proteins in normal and HIV-infected cells.
HIV Nef and Vpu protect HIV-infected CD4+ T cells from antibody-mediated cell lysis through down-modulation of CD4 and BST2.
Recognition of two overlapping CTL epitopes in HIV-1 p17 by CTL from a long-term nonprogressing HIV-1-infected individual.
Human leukocyte antigen (HLA) class I down-regulation by human immunodeficiency virus type 1 negative factor (HIV-1 Nef): what might we learn from natural sequence variants?
Aminopeptidase substrate preference affects HIV epitope presentation and predicts immune escape patterns in HIV-infected individuals.
Escape from the dominant HLA-B27-restricted cytotoxic T-lymphocyte response in Gag is associated with a dramatic reduction in human immunodeficiency virus type 1 replication.
Cellular immune responses and viral diversity in individuals treated during acute and early HIV-1 infection.
Design and use of an inducibly activated human immunodeficiency virus type 1 Nef to study immune modulation.
Impact of a single HLA-A*24:02-associated escape mutation on the detrimental effect of HLA-B*35:01 in HIV-1 control.
Novel AIDS therapies based on gene editing.
HIV viral diversity and escape from cellular immunity.
Down-modulation of CD8 $\alpha\beta$ is a fundamental activity of primate lentiviral Nef proteins.
CD4+ target cell availability determines the dynamics of immune escape and reversion in vivo.
Differential Ability of Primary HIV-1 Nef Isolates To Downregulate HIV-1 Entry Receptors.
An early HIV mutation within an HLA-B*57-restricted T cell epitope abrogates binding to the killer inhibitory receptor 3DL1.
A safe, effective and affordable HIV vaccine--an urgent global need.
Clinicopathological features of papillary thyroid carcinoma in HIV-infected patients.
Transmitted virus fitness and host T cell responses collectively define divergent infection outcomes in two HIV-1 recipients.
Molecular signatures of T-cell inhibition in HIV-1 infection.
Cat and Mouse: HIV Transcription in Latency, Immune Evasion and Cure/Remission Strategies.
CD8+ and CD4+ cytotoxic T cell escape mutations precede breakthrough SIVmac239 viremia in an elite controller.
Expression of simian immunodeficiency virus Nef protein in CD4+ T cells leads to a molecular profile of viral persistence and immune evasion.
Estimating Costs and Benefits of CTL Escape Mutations in SIV/HIV Infection.

Progression to AIDS in South Africa is associated with both reverting and compensatory viral mutations.
CD8 T cell response maturation defined by anentropic specificity and repertoire depth correlates with SIVΔnef-induced protection.
The HIV-1 Nef protein as a target for antiretroviral therapy.
Genetic determinants of HIV-1 infection and progression to AIDS: immune response genes.
Characterization of CD8 T Cell-Mediated Mutations in the Immunodominant Epitope GP33-41 of Lymphocytic Choriomeningitis Virus.
Widespread impact of HLA restriction on immune control and escape pathways of HIV-1.
The immune response to HIV: implications for vaccine development.
Mathematical modeling of ultradeep sequencing data reveals that acute CD8+ T-lymphocyte responses exert strong selective pressure in simian immunodeficiency virus-infected macaques but still fail to clear founder epitope sequences.
Cellular and molecular interactions in HIV infections: a review.
Immune selection in vitro reveals human immunodeficiency virus type 1 Nef sequence motifs important for its immune evasion function in vivo.
T-cell responses to sequentially emerging viral escape mutants shape long-term HIV-1 population dynamics.
Can immunotherapy be useful as a "functional cure" for infection with Human Immunodeficiency Virus-1?
PROTAC-mediated Degradation of HIV-1 Nef Efficiently Restores Cell-surface CD4 and MHC-I Expression and Blocks HIV-1 Replication.
Multifunctional Roles of the N-Terminal Region of HIV-1(SF2)Nef Are Mediated by Three Independent Protein Interaction Sites.
Selective downregulation of rhesus macaque and sooty mangabey major histocompatibility complex class I molecules by Nef alleles of simian immunodeficiency virus and human immunodeficiency virus type 2.
HIV-1 and its causal relationship to immunosuppression and nervous system disease in AIDS: a review.
Viral evolution in HLA-B27-restricted CTL epitopes in human immunodeficiency virus type 1-infected individuals.
Validation of a novel secretion modification region (SMR) of HIV-1 Nef using cohort sequence analysis and molecular modeling.
Complement opsonization of HIV-1 results in decreased antiviral and inflammatory responses in immature dendritic cells via CR3.
Endocytosis of HIV: anything goes.
HIV-Host Cell Interactions.
Adenovirus-vectored vaccine containing multidimensionally conserved parts of the HIV proteome is immunogenic in rhesus macaques.
Impaired protective role of HLA-B*57:01/58:01 in HIV-1 CRF01_AE infection: a cohort study in Vietnam.
CCR5-edited CD4+ T cells augment HIV-specific immunity to enable post-rebound control of HIV replication.
HIV-1 and hijacking of the host immune system: the current scenario.
Early immune adaptation in HIV-1 revealed by population-level approaches.
Selection of escape mutant by HLA-C-restricted HIV-1 Pol-specific cytotoxic T lymphocytes carrying strong ability to suppress HIV-1 replication.



Amino Acid Substitutions within HLA-B*27-Restricted T Cell Epitopes Prevent Recognition by Hepatitis Delta Virus-Specific CD8(+) T Cells.
Associations between phylogenetic clustering and HLA profile among HIV-infected individuals in San Diego, California.
HIV and interferon regulatory factor 1: a story of manipulation and control.
Correlates of spontaneous viral control among long-term survivors of perinatal HIV-1 infection expressing human leukocyte antigen-B57.
Reversion of immune escape HIV variants upon transmission: insights into effective viral immunity.
Inhibitors of HIV-1 Nef-Mediated Activation of the Myeloid Src-Family Kinase Hck Block HIV-1 Replication in Macrophages and Disrupt MHC-I Downregulation.
HIV-1 epitopes presented by MHC class I types associated with superior immune containment of viremia have highly constrained fitness landscapes.
Transcription of preintegrated HIV-1 cDNA modulates cell surface expression of major histocompatibility complex class I via Nef.
Human immunodeficiency virus type 1 Nef domains required for disruption of major histocompatibility complex class I trafficking are also necessary for coprecipitation of Nef with HLA-A2.
A molecular switch in immunodominant HIV-1-specific CD8 T-cell epitopes shapes differential HLA-restricted escape.
Accumulation of specific amino acid substitutions in HLA-B35-restricted human immunodeficiency virus type 1 cytotoxic T lymphocyte epitopes.
Efficient processing of the immunodominant, HLA-A*0201-restricted human immunodeficiency virus type 1 cytotoxic T-lymphocyte epitope despite multiple variations in the epitope flanking sequences.
Nef-Mediated CD3-TCR Downmodulation Dampens Acute Inflammation and Promotes SIV Immune Evasion.
HIV's evasion of the cellular immune response.
Immune escape from HIV-specific antibody-dependent cellular cytotoxicity (ADCC) pressure.
Early viral replication in lymph nodes provides HIV with a means by which to escape NK-cell-mediated control.
Multilayered defense in HLA-B51-associated HIV viral control.
The influence of delivery vectors on HIV vaccine efficacy.
Impact of HLA-B*81-associated mutations in HIV-1 Gag on viral replication capacity.
Genetic complexity in the replication-competent latent HIV reservoir increases with untreated infection duration in infected youth.
HIV-1 infection of CD4 T cells impairs antigen-specific B cell function.
[Coreceptors of HIV infection and the development of HIV entry inhibitors: overview and targets].
Multiple transmissions of a stable human leucocyte antigen-B27 cytotoxic T-cell-escape strain of HIV-1 in The Netherlands.
Vaccination and timing influence SIV immune escape viral dynamics in vivo.
Host-specific adaptation of HIV-1 subtype B in the Japanese population.
Genetically barcoded SIV reveals the emergence of escape mutations in multiple viral lineages during immune escape.
Crystal structures and KIR3DL1 recognition of three immunodominant viral peptides complexed to HLA-B*2705.
Two Functional Variants of AP-1 Complexes Composed of either $\gamma 2$ or $\gamma 1$ Subunits Are Independently Required for Major Histocompatibility Complex Class I Downregulation by HIV-1 Nef.
An introduction to a novel population genetic approach for HIV characterization.

Selection pressure in CD8 <sup>+</sup> T-cell epitopes in the pol gene of HIV-1 infected individuals in Colombia. A bioinformatic approach.
Dynamic range of Nef-mediated evasion of HLA class II-restricted immune responses in early HIV-1 infection.
In vivo fitness costs of different Gag CD8 T-cell escape mutant simian-human immunodeficiency viruses for macaques.
Distinct HIV-1 escape patterns selected by cytotoxic T cells with identical epitope specificity.
Viral evolution and escape during primary human immunodeficiency virus-1 infection: implications for vaccine design.
Collaboration of a Detrimental HLA-B*35:01 Allele with HLA-A*24:02 in Coevolution of HIV-1 with T Cells Leading to Poorer Clinical Outcomes.
Fitness constraints on immune escape from HIV: Implications of envelope as a target for both HIV-specific T cells and antibody.
Analysis of immune selection as a potential cause for the presence of cleavage site mutation 431V in treatment-naïve HIV type-1 isolates.
Persistent survival of prevalent clonotypes within an immunodominant HIV gag-specific CD8 <sup>+</sup> T cell response.
Mutations in a dominant Nef epitope of simian immunodeficiency virus diminish TCR:epitope peptide affinity but not epitope peptide:MHC class I binding.
Emerging complexities of APOBEC3G action on immunity and viral fitness during HIV infection and treatment.
Living in oblivion: HIV immune evasion.
[Mechanism for HIV invasion via skin or mucosa].
Novel cytotoxic T-lymphocyte escape mutation by a three-amino-acid insertion in the human immunodeficiency virus type 1 p6Pol and p6Gag late domain associated with drug resistance.
Dynamics of immune escape in HIV infection.
A simple screening approach to reduce B*5701-associated abacavir hypersensitivity on the basis of sequence variation in HIV reverse transcriptase.
Human immunodeficiency virus type 1 Nef: adapting to intracellular trafficking pathways.
Nef-mediated down-regulation of CD4 and HLA class I in HIV-1 subtype C infection: association with disease progression and influence of immune pressure.
An "escape clock" for estimating the turnover of SIV DNA in resting CD4 <sup>+</sup> T cells.
HIV/SIV glycoproteins: structure-function relationships.
Natural variants of cytotoxic epitopes are T-cell receptor antagonists for antiviral cytotoxic T cells.
HIV-1 Nef Disrupts CD4(+) T Lymphocyte Polarity, Extravasation, and Homing to Lymph Nodes via Its Nef-Associated Kinase Complex Interface.
Might dolutegravir be part of a functional cure for HIV?
Immune responses against multiple epitopes.
Human T cells expressing two additional receptors (TETARs) specific for HIV-1 recognize both epitopes.
Diverse peptide presentation of rhesus macaque major histocompatibility complex class I Mamu-A 02 revealed by two peptide complex structures and insights into immune escape of simian immunodeficiency virus.
Infection with cytotoxic T-lymphocyte escape mutants results in increased mortality and growth retardation in mice infected with a neurotropic coronavirus.
The Nef protein of HIV-1 induces loss of cell surface costimulatory molecules CD80 and CD86 in APCs.
Dynamics of haplotype frequency change in a CD8+TL epitope of simian immunodeficiency virus.

A natural polymorphism of Mycobacterium tuberculosis in the esxH gene disrupts immunodomination by the TB10.4-specific CD8 T cell response.
Influence of HLA-C expression level on HIV control.
Avoiding the kiss of death: how HIV and other chronic viruses survive.
Elite Control, Gut CD4 T Cell Sparing, and Enhanced Mucosal T Cell Responses in Macaca nemestrina Infected by a Simian Immunodeficiency Virus Lacking a gp41 Trafficking Motif.
Sensing of HIV-1 Infection in Tzm-bl Cells with Reconstituted Expression of STING.
HIV-2 Surveillance with Next-Generation Sequencing Reveals Mutations in a Cytotoxic Lymphocyte-Restricted Epitope Involved in Long-Term Nonprogression.
Integrin alpha4beta7 is downregulated on the surfaces of simian immunodeficiency virus SIVmac239-infected cells.
Apoptosis as an HIV strategy to escape immune attack.
RNA structures, genomic organization and selection of recombinant HIV.
HIV impairment of immune responses in dendritic cells.
IL-10 high producing genotype predisposes HIV infected individuals to TB infection.
The interaction of HIV with dendritic cells: outcomes and pathways.
HIV-1 viral escape in infancy followed by emergence of a variant-specific CTL response.
Depletion and dysfunction of Vγ2Vδ2 T cells in HIV disease: mechanisms, impacts and therapeutic implications.
HLA-C downregulation by HIV-1 adapts to host HLA genotype.
Diversity in the acute CD8 T cell response to vaccinia virus in humans.
High-throughput profiling of anti-glycan humoral responses to SIV vaccination and challenge.
Microbial Protein Binding to gC1qR Drives PLA2G1B-Induced CD4 T-Cell Anergy.
Highly tissue substructure-specific effects of human papilloma virus in mucosa of HIV-infected patients revealed by laser-dissection microscopy-assisted gene expression profiling.
Correlation between viral RNA levels but not immune responses in plasma and tissues of macaques with long-standing SIVmac251 infection.
Broadly targeted CD8 <sup>+</sup> T cell responses restricted by major histocompatibility complex E.
HIV-derived vectors for gene therapy targeting dendritic cells.
Mutational networks of escape from transmitted HIV-1 infection.
Rare Control of SIVmac239 Infection in a Vaccinated Rhesus Macaque.
Variability of persisting MHV RNA sequences constituting immune and replication-relevant domains.
Reversion and T cell escape mutations compensate the fitness loss of a CD8 <sup>+</sup> T cell escape mutant in their cognate transmitted/founder virus.
Spatial alterations between CD4(+) T follicular helper, B, and CD8(+) T cells during simian immunodeficiency virus infection: T/B cell homeostasis, activation, and potential mechanism for viral escape.
Evolution of mutation rate and virulence among human retroviruses.
CD8 memory, immunodominance, and antigenic escape.
Challenges and Opportunities for T-Cell-Mediated Strategies to Eliminate HIV Reservoirs.
Effect of Passive Administration of Monoclonal Antibodies Recognizing Simian Immunodeficiency Virus (SIV) V2 in CH59-Like Coil/Helical or β-Sheet Conformations on Time of SIV(mac251) Acquisition.
TRIM5α Resistance Escape Mutations in the Capsid Are Transferable between Simian Immunodeficiency Virus Strains.
Persistent production of an integrase-deleted HIV-1 variant with no resistance mutation and wild-type proviral DNA in a treated patient.

HIV Vaccine Mystery and Viral Shell Disorder.
Activation of the lectin DC-SIGN induces an immature dendritic cell phenotype triggering Rho-GTPase activity required for HIV-1 replication.
Differences of cytotoxic T-lymphocyte pressure and divergent evolution of several CRF07_BC clusters circulating in men who have sex with men in China.
Determination of HLA-A*02 antigen status in Hodgkin's disease and analysis of an HLA-A*02-restricted epitope of the Epstein-Barr virus LMP-2 protein.
Immunodominant Cytomegalovirus Epitopes Suppress Subdominant Epitopes in the Generation of High-Avidity CD8 T Cells.
Determination of structural principles underlying three different modes of lymphocytic choriomeningitis virus escape from CTL recognition.
Hide, shield and strike back: how HIV-infected cells avoid immune eradication.
Characterisation of a highly potent and near pan-neutralising anti-HIV monoclonal antibody expressed in tobacco plants.
Primate TRIM34 is a broadly-acting, TRIM5-dependent lentiviral restriction factor.
Adenovirus hexon T-cell epitope is recognized by most adults and is restricted by HLA DP4, the most common class II allele.
Retroviral proteins that target the major histocompatibility complex class I.
Emergence of cytotoxic T lymphocyte escape mutants following antiretroviral treatment suspension in rhesus macaques infected with SIVmac251.
Genetic and antigenic features of the transmitted virus.
Brief report: Circulating markers of fibrosis are associated with immune reconstitution status in HIV-infected men.
Multiantibody strategies for HIV.
Shell disorder and the HIV vaccine mystery: lessons from the legendary Oswald Avery.
CD8(+) T cells of Listeria monocytogenes-infected mice recognize both linear and spliced proteasome products.
HIV reservoirs and immune surveillance evasion cause the failure of structured treatment interruptions: a computational study.
A two-pronged mechanism for HIV-1 Nef-mediated endocytosis of immune costimulatory molecules CD80 and CD86.
Comparison of Adjuvant Effects of Montanide ISA-720 and Heat Shock Protein 27 in Increasing Immunostimulatory Properties of HIV-1 Nef-Vif Fusion Protein Construct.
HIV-1 Nef hijacks clathrin coats by stabilizing AP-1:Arf1 polygons.
Tuning antiviral CD8 T-cell response via proline-altered peptide ligand vaccination.
HLA class I haplotype diversity is consistent with selection for frequent existing haplotypes.
Recruitment of epitope-specific T cell clones with a low-avidity threshold supports efficacy against mutational escape upon re-infection.
Immune evasion in malaria: altered peptide ligands of the circumsporozoite protein.
Analysis of the helper virus in murine retrovirus-induced immunodeficiency syndrome: evidence for immunoselection of the dominant and subdominant CTL epitopes of the BM5 ecotropic virus.
Innate and adaptive immunological insights into HIV pathogenesis.
Vulnerable targets in HIV-1 Pol for attenuation-based vaccine design.
Specific mutation of a gammaherpesvirus-expressed antigen in response to CD8 T cell selection in vivo.
Rapid deletion of antigen-specific CD4+ T cells following infection represents a strategy of immune evasion and persistence for Anaplasma marginale.

Improving the MVA vaccine potential by deleting the viral gene coding for the IL-18 binding protein.
The impact of a large and frequent deletion in the human TCR $\beta$ locus on antiviral immunity.
The rational design of an AIDS vaccine.
Cytomegalovirus-Specific T Cell Epitope Recognition in Congenital Cytomegalovirus Mother-Infant Pairs.
Codon optimization and improved delivery/immunization regimen enhance the immune response against wild-type and drug-resistant HIV-1 reverse transcriptase, preserving its Th2-polarity.
[Purification, crystallographic analysis of rhesus MHC-I Mamu-A*02 complexed with simian immunodeficiency virus nonapeptide].
In vitro modeling of the HIV-macrophage reservoir.
A Targeted LC-MS Strategy for Low-Abundant HLA Class-I-Presented Peptide Detection Identifies Novel Human Papillomavirus T-Cell Epitopes.
Distinct patterns of natural selection in the reverse transcriptase gene of HIV-1 in the presence and absence of antiretroviral therapy.
Cellular Determinants of HIV Persistence on Antiretroviral Therapy.
Antigen display, T-cell activation, and immune evasion during acute and chronic ehrlichiosis.
Enhancement to the RANKPEP resource for the prediction of peptide binding to MHC molecules using profiles.
Modelling the impact of acute infection dynamics on the accumulation of HIV-1 mutations.
Rapid recruitment of virus-specific CD8 T cells restructures immunodominance during protective secondary responses.
Retroviral restriction: nature's own solution.
Relapse of Symptomatic Cerebrospinal Fluid HIV Escape.
HIV-1 gp41 and TCRalpha trans-membrane domains share a motif exploited by the HIV virus to modulate T-cell proliferation.
Dynamics of physical interaction between HIV-1 Nef and ASK1: identifying the interacting motif(s).
Viral subversion of APOBEC3s: Lessons for anti-tumor immunity and tumor immunotherapy.
On the benefits of sin: can greater understanding of the 1F7-idiotypic repertoire freeze enhance HIV vaccine development?
The rationale for pathogen-inactivation treatment of blood components.
HTLV-I viral escape and host genetic changes in the development of adult T cell leukemia.
A structural basis for LCMV immune evasion: subversion of H-2D(b) and H-2K(b) presentation of gp33 revealed by comparative crystal structure.Analyses.
The Immunogenicity of a Proline-Substituted Altered Peptide Ligand toward the Cancer-Associated TEIPP Neopeptide Trh4 Is Unrelated to Complex Stability.
A strategy for cloning infectious molecular clones of retroviruses from serum or plasma.
Role of cytotoxic T lymphocytes and interferon- $\gamma$ in coronavirus infection: Lessons from murine coronavirus infections in mice.
Proviral location affects cognate peptide-induced virus production and immune recognition of HIV-1-infected T cell clones.
HIV-1 Nef induces a Rab11-dependent routing of endocytosed immune costimulatory proteins CD80 and CD86 to the Golgi.
The role of Fc receptors in HIV prevention and therapy.
Accurate assembly of minority viral haplotypes from next-generation sequencing through efficient noise reduction.
Role of genetic susceptibility to latent adenoviral infection and decreased lung function.

Matrix-Metalloprotease Resistant Mucin-16 (MUC16) Peptide Mutants Represent a Worse Lung Adenocarcinoma Outcome.

Table A2-45, Cluster 44

Cluster 44 focuses on cancer stem cells, and methods for targeting of cancer stem cells within the tumor microenvironment (318)
The Role of Cancer Stem Cells and Their Extracellular Vesicles in the Modulation of the Antitumor Immunity.
New Opportunities and Challenges to Defeat Cancer Stem Cells.
Can breast cancer stem cells evade the immune system?
Cancer stem cells targets and combined therapies to prevent cancer recurrence.
Extracellular vesicles as regulators of tumor fate: crosstalk among cancer stem cells, tumor cells and mesenchymal stem cells.
Designing precision medicine panels for drug refractory cancers targeting cancer stemness traits.
Mechanisms of cancer stem cell therapy.
Advances in Therapeutic Targeting of Cancer Stem Cells within the Tumor Microenvironment: An Updated Review.
Pancreatic cancer stem cells: new insights and perspectives.
The Immune Privilege of Cancer Stem Cells: A Key to Understanding Tumor Immune Escape and Therapy Failure.
Macrophages and cancer stem cells: a malevolent alliance.
Judith Agudo: Beware of your inner self-immune attack.
Cancer Stem Cell Relationship with Pro-Tumoral Inflammatory Microenvironment.
Novel therapeutic strategies for targeting liver cancer stem cells.
Impact of Cancer Stem Cells and Cancer Stem Cell-Driven Drug Resiliency in Lung Tumor: Options in Sight.
Stem cell programs in cancer initiation, progression, and therapy resistance.
Cancer stem cells in esophageal cancer and response to therapy.
Non-coding RNAs Functioning in Colorectal Cancer Stem Cells.
The Role of the Microenvironment and Immune System in Regulating Stem Cell Fate in Cancer.
Evolving Strategies for Therapeutically Targeting Cancer Stem Cells.
Recent Advances in Cancer Stem Cell Targeted Therapy.
Cancer stem cell immunology: key to understanding tumorigenesis and tumor immune escape?
Hide-and-seek: the interplay between cancer stem cells and the immune system.
Cancer Stem Cells: Emerging Key Players in Immune Evasion of Cancers.
Immune evader cancer stem cells direct the perspective approaches to cancer immunotherapy.
Cancer stem cells in gliomas: evolving concepts and therapeutic implications.
Bidirectional Crosstalk Between Cancer Stem Cells and Immune Cell Subsets.
The intersection of cancer, cancer stem cells, and the immune system: therapeutic opportunities.
Lung Cancer Stem Cells and Their Clinical Implications.
Current understanding of cancer stem cells: Immune evasion and targeted immunotherapy in gastrointestinal malignancies.
Approaches for targeting cancer stem cells drug resistance.
Immune evasion by cancer stem cells.

Cancer Stem Cells Targeting; the Lessons from the Interaction of the Immune System, the Cancer Stem Cells and the Tumor Niche.
Altered BMP2/4 Signaling in Stem Cells and Their Niche: Different Cancers but Similar Mechanisms, the Example of Myeloid Leukemia and Breast Cancer.
Targeting Cancer Stem Cells-A Renewed Therapeutic Paradigm.
The significance of the pluripotency and cancer stem cell-related marker NANOG in diagnosis and treatment of ovarian carcinoma.
Specific targeting of cancer stem cells by immunotherapy: A possible stratagem to restrain cancer recurrence and metastasis.
Role of Microenvironment on the Fate of Disseminating Cancer Stem Cells.
[Immunotherapy Against Head and Neck Cancer Stem Cells].
Involvement of mesenchymal stem cells in cancer progression and metastases.
Ovarian cancer stem cells: Critical roles in anti-tumor immunity.
Inflammatory regulation of stem-like cells in melanoma.
Melanoma stem cells: not rare, but well done.
SOX2 in cancer stemness: tumor malignancy and therapeutic potentials.
Urothelial carcinoma: stem cells on the edge.
Roles of microRNAs in Regulating Cancer Stemness in Head and Neck Cancers.
Colorectal cancer stem cells: from the crypt to the clinic.
Targeting cancer stem cells: emerging role of Nanog transcription factor.
CD59 Regulation by SOX2 Is Required for Epithelial Cancer Stem Cells to Evade Complement Surveillance.
Roles Played by YY1 in Embryonic, Adult and Cancer Stem Cells.
Cancer Stem Cells: Devil or Savior-Looking behind the Scenes of Immunotherapy Failure.
Cancer non-stem cells as a potent regulator of tumor microenvironment: a lesson from chronic myeloid leukemia.
Drug and apoptosis resistance in cancer stem cells: a puzzle with many pieces.
Linking Tumor Microenvironment to Plasticity of Cancer Stem Cells: Mechanisms and Application in Cancer Therapy.
Quiescent Tissue Stem Cells Evade Immune Surveillance.
Cancer stem cells, CD200 and immunoevasion.
Glioblastoma Cancer Stem Cells Evade Innate Immune Suppression of Self-Renewal through Reduced TLR4 Expression.
Understanding and Overcoming Immunosuppression Shaped by Cancer Stem Cells.
Influence of Innate Immunity on Cancer Cell Stemness.
Cancer Stem Cells in the Immune Microenvironment.
Molecular Chaperones in Cancer Stem Cells: Determinants of Stemness and Potential Targets for Antitumor Therapy.
Cancer Stem Cells Are Possible Key Players in Regulating Anti-Tumor Immune Responses: The Role of Immunomodulating Molecules and MicroRNAs.
Role of cancer stem cell-derived extracellular vesicles in cancer progression and metastasis.
Cancer stem cell antigens as targets for new combined anti-cancer therapies.
Characterization of gastric cancer stem-like molecular features, immune and pharmacogenomic landscapes.
The role of cancer stem cells in the modulation of anti-tumor immune responses.
Overcoming resistance to immunotherapy by teaching old drugs new tricks.

Hepatoblastoma Cancer Stem Cells Express PD-L1, Reveal Plasticity and Can Emerge upon Chemotherapy.
Cancer stem cell-immune cell collusion in immunotherapy.
Effects of pericytes and colon cancer stem cells in the tumor microenvironment.
Mitotic quiescence in hepatic cancer stem cells: An incognito mode.
Tumor dormancy and cancer stem cells: two sides of the same coin?
Histone modifications in drug-resistant cancers: From a cancer stem cell and immune evasion perspective.
The therapeutic promise of the cancer stem cell concept.
Co-expression of the toleragenic glycoprotein, CD200, with markers for cancer stem cells.
Stem cell transcription factor NANOG in cancers--is eternal youth a curse?
The cross talk between gastric cancer stem cells and the immune microenvironment: a tumor-promoting factor.
PDE4 inhibitor eliminates breast cancer stem cells via noncanonical activation of mTOR.
Cancer Stem Cells and Neuroblastoma: Characteristics and Therapeutic Targeting Options.
Emerging functional markers for cancer stem cell-based therapies: Understanding signaling networks for targeting metastasis.
Epigenetic and Immune Regulation of Colorectal Cancer Stem Cells.
[PD-L1-Mediated Immune Escape Mechanism of Cancer Stem-Like Cells].
Meeting the Challenge of Targeting Cancer Stem Cells.
E3 ubiquitin ligases in cancer stem cells: key regulators of cancer hallmarks and novel therapeutic opportunities.
Mechanisms of immune evasion by head and neck cancer stem cells.
Effect of melanoma stem cells on melanoma metastasis.
KYA1797K down-regulates PD-L1 in colon cancer stem cells to block immune evasion by suppressing the $\beta$ -catenin/STT3 signaling pathway.
Tuning Cancer Fate: Tumor Microenvironment's Role in Cancer Stem Cell Quiescence and Reawakening.
Targeting Leukemia Stem Cells in the Bone Marrow Niche.
Wharton's jelly-derived mesenchymal stem cells: phenotypic characterization and optimizing their therapeutic potential for clinical applications.
Activation and Differentiation of Mesenchymal Stem Cells.
Gastrointestinal cancer stem cells as targets for innovative immunotherapy.
Metabolic Plasticity of Stem Cells and Macrophages in Cancer.
Cancer stem cells-driven tumor growth and immune escape: the Janus face of neurotrophins.
Type I IFNs promote cancer cell stemness by triggering the epigenetic regulator KDM1B.
Mesenchymal stem cells avoid allogeneic rejection.
The paracrine induction of prostate cancer progression by caveolin-1.
Inflammatory Mediators in Glioma Microenvironment Play a Dual Role in Gliomagenesis and Mesenchymal Stem Cell Homing: Implication for Cellular Therapy.
Identification and editing of stem-like cells in methylcholanthrene-induced sarcomas.
Surface PD-L1, E-cadherin, CD24, and VEGFR2 as markers of epithelial cancer stem cells associated with rapid tumorigenesis.
Crucial role of glucosylceramide synthase in the regulation of stem cell-like cancer cells in B16F10 murine melanoma.



Interactions between cancer stem cells, immune system and some environmental components: Friends or foes?
TGF-beta in neural stem cells and in tumors of the central nervous system.
Cancer stem cell metabolism: a potential target for cancer therapy.
Activation and differentiation of mesenchymal stem cells.
TLRgeting Evasion of Immune Pathways in Glioblastoma.
MDSCs sneak CSCs out of (immuno)surveillance.
Antitumor immunity and cancer stem cells.
Epigenetic Silencing of TAP1 in Aldefluor(+) Breast Cancer Stem Cells Contributes to Their Enhanced Immune Evasion.
WNT signaling and cancer stemness.
Immune evasion through competitive inhibition: the shielding effect of cancer non-stem cells.
Accept or Reject: The Role of Immune Tolerance in the Development of Stem Cell Therapies and Possible Future Approaches.
Type I interferon induces cancer stem cells-mediated chemotherapy resistance.
The metabolic addiction of cancer stem cells.
T(reg)/Th17 polarization by distinct subsets of breast cancer cells is dictated by the interaction with mesenchymal stem cells.
Aldehyde dehydrogenase-positive melanoma stem cells in tumorigenesis, drug resistance and anti-neoplastic immunotherapy.
TGF-β: A novel predictor and target for anti-PD-1/PD-L1 therapy.
TGF-β-Mediated Epithelial-Mesenchymal Transition and Cancer Metastasis.
A Perspective on the Development of TGF-β Inhibitors for Cancer Treatment.
The dynamic roles of TGF-β in cancer.
Characterization of Immunogenicity of Malignant Cells with Stemness in Intrahepatic Cholangiocarcinoma by Single-Cell RNA Sequencing.
The Oncogenic Potential of Mesenchymal Stem Cells in the Treatment of Cancer: Directions for Future Research.
Recent Advances in the Development of TGF-β Signaling Inhibitors for Anticancer Therapy.
IFNγ-induced stem-like state of cancer cells as a driver of metastatic progression following immunotherapy.
Transforming growth factor β signaling pathway: A promising therapeutic target for cancer.
Transforming Growth Factor-β Signaling in Immunity and Cancer.
Transforming growth factor-β signaling: Tumorigenesis and targeting for cancer therapy.
4-Acetylanthroquinonol B from antrodia cinnamomea enhances immune function of dendritic cells against liver cancer stem cells.
Different Shades of L1CAM in the Pathophysiology of Cancer Stem Cells.
[Mesenchymal stem/stromal cells. Its therapeutic potential in medicine].
How to conjugate the stemness marker ALDH1A1 with tumor angiogenesis, progression, and drug resistance.
Melanoma stem cells: targets for successful therapy?
Tumor-stroma interactions in tumorigenesis: lessons from stem cell biology.
[Mesenchymal stem cells and regenerative medicine in liver cirrhosis].
TLR4 interaction with LPS in glioma CD133+ cancer stem cells induces cell proliferation, resistance to chemotherapy and evasion from cytotoxic T lymphocyte-induced cytotoxicity.
Immune privilege of skin stem cells: What do we know and what can we learn?

Tubulin Isotypes: Emerging Roles in Defining Cancer Stem Cell Niche.
The Role of TGF- $\beta$ Signaling Pathways in Cancer and Its Potential as a Therapeutic Target.
B7-H1 enhances proliferation ability of gastric cancer stem-like cells as a receptor.
The role of transforming growth factor $\beta$ in upper gastrointestinal cancers: A systematic review.
Canonical and non-canonical WNT signaling in cancer stem cells and their niches: Cellular heterogeneity, omics reprogramming, targeted therapy and tumor plasticity (Review).
The Bright and the Dark Side of TGF- $\beta$ Signaling in Hepatocellular Carcinoma: Mechanisms, Dysregulation, and Therapeutic Implications.
Novel therapies emerging in oncology to target the TGF- $\beta$ pathway.
The novel role of SERPINB9 in cytotoxic protection of human mesenchymal stem cells.
MSCs as Tumor-Specific Vectors for the Delivery of Anticancer Agents-A Potential Therapeutic Strategy in Cancer Diseases: Perspectives for Quinazoline Derivatives.
Research advances in the role of gastric cancer-derived mesenchymal stem cells in tumor progression (Review).
PVT1 inhibition stimulates anti-tumor immunity, prevents metastasis, and depletes cancer stem cells in squamous cell carcinoma.
Hematopoiesis from human embryonic stem cells: overcoming the immune barrier in stem cell therapies.
TGF- $\beta$ : duality of function between tumor prevention and carcinogenesis.
Transforming growth factor-beta (TGF- $\beta$ ) in prostate cancer: A dual function mediator?
Cancer Stem Cells and Immunosuppressive Microenvironment in Glioma.
Transforming growth factor-beta: recent advances on its role in immune tolerance.
Multiple myeloma mesenchymal stem cells: characterization, origin, and tumor-promoting effects.
Tumor-Infiltrating Immunosuppressive Cells in Cancer-Cell Plasticity, Tumor Progression and Therapy Response.
Generation and characterization of human mesenchymal stem/stromal cells for cell therapy applications.
Small extracellular vesicle TGF- $\beta$ in cancer progression and immune evasion.
Mesenchymal stem cells express serine protease inhibitor to evade the host immune response.
The predictive value of transforming growth factor- $\beta$ in Wilms tumor immunopathogenesis.
Mechanisms navigating the TGF- $\beta$ pathway in prostate cancer.
Review: Targeting the Transforming Growth Factor-Beta Pathway in Ovarian Cancer.
Nanog signaling in cancer promotes stem-like phenotype and immune evasion.
Epithelial-Mesenchymal Transition Signaling and Prostate Cancer Stem Cells: Emerging Biomarkers and Opportunities for Precision Therapeutics.
Tumor initiation in human malignant melanoma and potential cancer therapies.
Identifying the Therapeutic Significance of Mesenchymal Stem Cells.
Impairment of NKG2D-Mediated Tumor Immunity by TGF- $\beta$ .
Rebuilding the Damaged Heart: Mesenchymal Stem Cells, Cell-Based Therapy, and Engineered Heart Tissue.
Insights into the Transforming Growth Factor- $\beta$ Signaling Pathway in Cutaneous Melanoma.
TGF- $\beta$ Signaling in Liver, Pancreas, and Gastrointestinal Diseases and Cancer.
Mesenchymal stem cells inhibit generation and function of both CD34+-derived and monocyte-derived dendritic cells.
Mesenchymal stem cells as a potential therapeutic tool to cure cognitive impairment caused by neuroinflammation.

Enhanced survival of hypoimmunogenic otic progenitors following intracochlear xenotransplantation: repercussions for stem cell therapy in hearing loss models.
Stem cells and targeted approaches to melanoma cure.
Efficient DNA Repair Mitigates Replication Stress Resulting in Less Immunogenic Cytosolic DNA in Radioresistant Breast Cancer Stem Cells.
Impact of proteolysis on cancer stem cell functions.
Mesenchymal stem cell treatment for hyperactive immune response in patients with COVID-19.
Effect of mesenchymal stem cells-derived exosomes on tumor microenvironment: Tumor progression versus tumor suppression.
IL-17A secreted from lymphatic endothelial cells promotes tumorigenesis by upregulation of PD-L1 in hepatoma stem cells.
TGF- $\beta$ 2 Reduces the Cell-Mediated Immunogenicity of Equine MHC-Mismatched Bone Marrow-Derived Mesenchymal Stem Cells Without Altering Immunomodulatory Properties.
Combination of MSC spheroids wrapped within autologous composite sheet dually protects against immune rejection and enhances stem cell transplantation efficacy.
Brain cancer stem cells: resilience through adaptive plasticity and hierarchical heterogeneity.
Role and effect of vein-transplanted human umbilical cord mesenchymal stem cells in the repair of diabetic foot ulcers in rats.
Manipulating TGF- $\beta$ signaling to optimize immunotherapy for cervical cancer.
Preoperative administration of polysaccharide Kureha and reduced plasma transforming growth factor- $\beta$ in patients with advanced gastric cancer: A randomized clinical trial.
Dormancy and cancer stem cells: An enigma for cancer therapeutic targeting.
Leukemic stem cell signatures in Acute myeloid leukemia- targeting the Guardians with novel approaches.
Tenascin-C Protects Cancer Stem-like Cells from Immune Surveillance by Arresting T-cell Activation.
Turning Cold Tumors Hot by Blocking TGF- $\beta$ .
Roles of lysine-specific demethylase 1 (LSD1) in homeostasis and diseases.
circFAT1 Promotes Cancer Stemness and Immune Evasion by Promoting STAT3 Activation.
A state of stochastic cancer stemness through the CDK1-SOX2 axis.
Treatment of COVID-19 by stage: any space left for mesenchymal stem cell therapy?
Microenvironment in neuroblastoma: isolation and characterization of tumor-derived mesenchymal stromal cells.
Epigenetic maintenance strategies after allogeneic stem cell transplantation in acute myeloid leukemia.
Hematopoietic stem cells: interplay with immunity.
Acute myeloid leukemia: therapeutic targeting of stem cells.
GDF-15: A Multifunctional Modulator and Potential Therapeutic Target in Cancer.
Integrin $\alpha\beta$ 8-expressing tumor cells evade host immunity by regulating TGF- $\beta$ activation in immune cells.
Interleukin-6 is overexpressed and augments invasiveness of human glioma stem cells in vitro.
The Blebbishield Emergency Program Overrides Chromosomal Instability and Phagocytosis Checkpoints in Cancer Stem Cells.
Emerging Role of Lymphocyte Antigen-6 Family of Genes in Cancer and Immune Cells.
Induced pluripotent stem cells display a distinct set of MHC I-associated peptides shared by human cancers.
Microenvironmental Features Driving Immune Evasion in Myelodysplastic Syndromes and Acute Myeloid Leukemia.

Immunosuppressant Drugs Mitigate Immune Responses Generated by Human Mesenchymal Stem Cells Transplanted into the Mouse Parenchyma.
Tumor-derived transforming growth factor- $\beta$ is critical for tumor progression and evasion from immune surveillance.
MiR-146b-5p overexpression attenuates stemness and radioresistance of glioma stem cells by targeting HuR/lincRNA-p21/ $\beta$ -catenin pathway.
A Kaposi's sarcoma-associated herpesvirus microRNA and its variants target the transforming growth factor $\beta$ pathway to promote cell survival.
B7-1 drives TGF- $\beta$ stimulated pancreatic carcinoma cell migration and expression of EMT target genes.
Immune Escape after Hematopoietic Stem Cell Transplantation (HSCT): From Mechanisms to Novel Therapies.
The effects of WW2/WW3 domains of Smurf2 molecule on TGF- $\beta$ signaling and arginase I gene expression.
The immunological role of mesenchymal stromal cells in patients with myelodysplastic syndrome.
Targeting FTO Suppresses Cancer Stem Cell Maintenance and Immune Evasion.
Oncogenic transformation tunes the cross-talk between mesenchymal stem cells and T lymphocytes.
API5 confers cancer stem cell-like properties through the FGF2-NANOG axis.
MSCs and inflammation: new insights into the potential association between ALCL and breast implants.
NDR1 activates CD47 transcription by increasing protein stability and nuclear location of ASCL1 to enhance cancer stem cell properties and evasion of phagocytosis in small cell lung cancer.
Blasts in context: the impact of the immune environment on acute myeloid leukemia prognosis and treatment.
Paroxysmal nocturnal hemoglobinuria: stem cells and clonality.
FGL2-wired macrophages secrete CXCL7 to regulate the stem-like functionality of glioma cells.
Proceedings from the National Cancer Institute's Second International Workshop on the Biology, Prevention, and Treatment of Relapse after Hematopoietic Stem Cell Transplantation: Part I. Biology of relapse after transplantation.
Overcoming Wnt- $\beta$ -catenin dependent anticancer therapy resistance in leukaemia stem cells.
Adaptive Immune Resistance Emerges from Tumor-Initiating Stem Cells.
Opposing roles and potential antagonistic mechanism between TGF- $\beta$ and BMP pathways: Implications for cancer progression.
A Century-long Journey From the Discovery of Insulin to the Implantation of Stem Cell-derived Islets.
An osteosarcoma zebrafish model implicates Mmp-19 and Ets-1 as well as reduced host immune response in angiogenesis and migration.
Glioblastoma Embryonic-like Stem Cells Exhibit Immune-Evasive Phenotype.
Wnt/ $\beta$ -catenin signalling in ovarian cancer: Insights into its hyperactivation and function in tumorigenesis.
Stemness and immune evasion conferred by the TDO2-AHR pathway are associated with liver metastasis of colon cancer.
Regulation of complement-dependent cytotoxicity by TGF- $\beta$ -induced epithelial-mesenchymal transition.
Galectin-3 in Prostate Cancer Stem-Like Cells Is Immunosuppressive and Drives Early Metastasis.
Vaccine and Cell-based Therapeutic Approaches in Acute Myeloid Leukemia.
B7-H1 is correlated with malignancy-grade gliomas but is not expressed exclusively on tumor stem-like cells.

The evolution of targeted therapy in pediatric AML: gemtuzumab ozogamicin, FLT3/IDH/BCL2 inhibitors, and other therapies.
Novel function of MDA-9/Syntenin (SDCBP) as a regulator of survival and stemness in glioma stem cells.
BCL7B, a SWI/SNF complex subunit, orchestrates cancer immunity and stemness.
Mesenchymal stem cell-mediated T cell suppression occurs through secreted galectins.
Wnt/beta-catenin pathway: modulating anticancer immune response.
Single-cell RNA-seq reveals developmental plasticity with coexisting oncogenic states and immune evasion programs in ETP-ALL.
$\beta$ -cell regeneration and differentiation: how close are we to the 'holy grail'?
BCR-ABL1-driven exosome-miR130b-3p-mediated gap-junction Cx43 MSC intercellular communications imply therapies of leukemic subclonal evolution.
Wnt signaling in cancer.
USP8 promotes cancer progression and extracellular vesicle-mediated CD8+ T cell exhaustion by deubiquitinating the TGF- $\beta$ receptor T $\beta$ RII.
A new chapter: hematopoietic stem cells are direct players in immunity.
Beneath the sword of Damocles: regenerative medicine and the shadow of immunogenicity.
Transcriptome Analysis of Mesenchymal Stem Cells from Multiple Myeloma Patients Reveals Downregulation of Genes Involved in Cell Cycle Progression, Immune Response, and Bone Metabolism.
Crosstalk between $\beta$ -Catenin and CCL2 Drives Migration of Monocytes towards Glioblastoma Cells.
HS-5 and HS-27A Stromal Cell Lines to Study Bone Marrow Mesenchymal Stromal Cell-Mediated Support to Cancer Development.
Macrophages in Oral Carcinomas: Relationship with Cancer Stem Cell Markers and PD-L1 Expression.
Circulating cells and exosomes in acute myelogenous leukemia and their role in disease progression and survival.
Hypoimmune induced pluripotent stem cell-derived cell therapeutics treat cardiovascular and pulmonary diseases in immunocompetent allogeneic mice.
Overcoming immunological barriers in regenerative medicine.
Tumor-initiating stem cell shapes its microenvironment into an immunosuppressive barrier and pro-tumorigenic niche.
Direct targeting of $\beta$ -catenin in the Wnt signaling pathway: Current progress and perspectives.
Suppression of Transposable Elements in Leukemic Stem Cells.
The immunometabolic landscape of the bone marrow microenvironment in acute myeloid leukemia.
Acute myeloid leukaemia and the immune system: implications for immunotherapy.
Acetylcholine promotes the self-renewal and immune escape of CD133+ thyroid cancer cells through activation of CD133-Akt pathway.
T11TS repress gliomagenic apoptosis of bone marrow hematopoietic stem cells.
Vascular endothelial growth factor inhibits the development of dendritic cells and dramatically affects the differentiation of multiple hematopoietic lineages in vivo.
New strategies in acute myelogenous leukemia: leukemogenesis and personalized medicine.
Immune Protection of Stem Cell-Derived Islet Cell Therapy for Treating Diabetes.
Molecular and phenotypic analyses of human embryonic stem cell-derived cardiomyocytes: opportunities and challenges for clinical translation.
Single-cell multiomics reveals increased plasticity, resistant populations, and stem-cell-like blasts in KMT2A-rearranged leukemia.
Stem Cell Membrane-Coated Nanogels for Highly Efficient In Vivo Tumor Targeted Drug Delivery.

Overexpression of Endogenous Retroviruses and Malignancy Markers in Neuroblastoma Cell Lines by Medium-Induced Microenvironmental Changes.
Embryonic stem cells: protecting pluripotency from alloreactivity.
Molecular insights into the role of electronic cigarettes in oral carcinogenesis.
Immunoregulatory functions and the therapeutic implications of GARP-TGF- $\beta$ in inflammation and cancer.
How do tumor stem cells actively escape from host immunosurveillance?
Modulation of Inflammatory Responses by Wnt/ $\beta$ -Catenin Signaling in Dendritic Cells: A Novel Immunotherapy Target for Autoimmunity and Cancer.
Immune impairments in multiple myeloma bone marrow mesenchymal stromal cells.
Seeing the good and bad in aplastic anemia: is autoimmunity in AA dysregulated or antineoplastic?
Suppression of tumor metastasis by blockade of transforming growth factor beta signaling in bone marrow cells through a retroviral-mediated gene therapy in mice.
Tumors promote altered maturation and early apoptosis of monocyte-derived dendritic cells.
Amphiphilic polyanhydride films promote neural stem cell adhesion and differentiation.
The sheep model of in utero gene therapy.
Immune escape and immunotherapy of acute myeloid leukemia.
Myeloid Sarcoma after Allogenic Stem Cell Transplantation for Acute Myeloid Leukemia: Successful Consolidation Treatment Approaches in Two Patients.
Differential roles of vascular endothelial growth factor receptors 1 and 2 in dendritic cell differentiation.
Microfluidic reprogramming to pluripotency of human somatic cells.
Interleukin-8: An evolving chemokine.
TSC22D3 as an immune-related prognostic biomarker for acute myeloid leukemia.
A Safety Checkpoint to Eliminate Cancer Risk of the Immune Evasive Cells Derived from Human Embryonic Stem Cells.
Autologous Stem Cell Transplantation for Myeloma: Cyto-reduction or an Immunotherapy?
Targeting the cluster of differentiation 47/signal-regulatory protein alpha axis in myeloid malignancies.
Generation of the human induced pluripotent stem cell line UKWNLi002-A from dermal fibroblasts of a woman with a heterozygous c.608 C>T (p.Thr203Met) mutation in exon 3 of the nerve growth factor gene potentially associated with hereditary sensory and autonomic neuropathy type 5.
Evaluation of $\beta$ -Catenin Inhibition of Axitinib and Nitazoxanide in Human Monocyte-Derived Dendritic Cells.
Role of TGF-beta in immune-evasion of cancer.
Fra-2 Expression in Osteoblasts Regulates Systemic Inflammation and Lung Injury through Osteopontin.
Biochemical mechanisms implemented by human acute myeloid leukemia cells to suppress host immune surveillance.
Transcriptomic Characterization of Postmolar Gestational Choriocarcinoma.
BCR-ABL Independent Mechanisms of Resistance in Chronic Myeloid Leukemia.
Amlexanox Downregulates S100A6 to Sensitize KMT2A/AFF1-Positive Acute Lymphoblastic Leukemia to TNF $\alpha$ Treatment.
"In medio stat virtus": Insights into hybrid E/M phenotype attitudes.
Human embryonic stem cell-extracts inhibit the differentiation and function of monocyte-derived dendritic cells.
Potential Targets to Mitigate Trauma- or Sepsis-Induced Immune Suppression.

SIRPα-αCD123 fusion antibodies targeting CD123 in conjunction with CD47 blockade enhance the clearance of AML-initiating cells.
Metastatic niche functions and therapeutic opportunities.
DUX4 expression in cancer induces a metastable early embryonic totipotent program.
EpCAM Inhibition Sensitizes Chemoresistant Leukemia to Immune Surveillance.
Silencing of retrotransposons by SETDB1 inhibits the interferon response in acute myeloid leukemia.
Multipotent fetal-derived Cdx2 cells from placenta regenerate the heart.
Schistosome infection-derived Hepatic Stellate Cells are cellular source of prostaglandin D <sub>2</sub> : role in TGF-β-stimulated VEGF production.
Biology and clinical management of hypoplastic MDS: MDS as a bone marrow failure syndrome.
MRD Monitoring by Multiparametric Flow Cytometry in AML: Is It Time to Incorporate Immune Parameters?
The immune landscape in BCR-ABL negative myeloproliferative neoplasms: inflammation, infections and opportunities for immunotherapy.
Effects of Polyphosphate on Leukocyte Function.
Senescent human hematopoietic progenitors show elevated expression of transposable elements and inflammatory genes.
A STAT3 decoy lures AML out of hiding.
Ocular immune privilege and ocular melanoma: parallel universes or immunological plagiarism?
It's Not What You Say But How You Say It: Targeting RNA Methylation in AML.
Clonality in immune aplastic anemia: Mechanisms of immune escape or malignant transformation.
Survival and maturation of human embryonic stem cell-derived cardiomyocytes in rat hearts.
Mutated clones driving leukemic transformation are already detectable at the single-cell level in CD34-positive cells in the chronic phase of primary myelofibrosis.
Mesenchymal Stromal Cells Are More Immunosuppressive In Vitro If They Are Derived from Endometriotic Lesions than from Eutopic Endometrium.
Cancer cells, on your histone marks, get SETDB1, silence retrotransposons, and go!
Microvesicles expressing adenosinergic ectoenzymes and their potential role in modulating bone marrow infiltration by neuroblastoma cells.
Staphylococcal superantigen-like 12 activates murine bone marrow derived mast cells.
Mutation-specific control of BCR-ABL T315I positive leukemia with a recombinant yeast-based therapeutic vaccine in a murine model.

Table A2-46, Cluster 45

Cluster 45 focuses on head and neck squamous cell carcinomas, especially immunotherapies to counter evasion of the immune system (300)
[Immunotherapies for head and neck squamous cell carcinomas].
[Immunotherapy: a new alternative treatment for head and neck squamous cell carcinoma].
The Double-Edged Sword-How Human Papillomaviruses Interact With Immunity in Head and Neck Cancer.
Protective mechanisms of head and neck squamous cell carcinomas from immune assault.
The emerging role of immunotherapy in head and neck squamous cell carcinoma (HNSCC): anti-tumor immunity and clinical applications.
Molecular biology and immunology of head and neck cancer.

The molecular mechanism of human papillomavirus-induced carcinogenesis in head and neck squamous cell carcinoma.
Promising systemic immunotherapies in head and neck squamous cell carcinoma.
Immune Checkpoints Pathways in Head and Neck Squamous Cell Carcinoma.
The head and neck cancer genome in the era of immunotherapy.
Molecular Aspects of Head and Neck Cancer Therapy.
Head and neck squamous cell carcinoma: Genomics and emerging biomarkers for immunomodulatory cancer treatments.
Current status and perspective of tumor immunotherapy for head and neck squamous cell carcinoma.
Heterogeneity of Head and Neck Squamous Cell Carcinoma Stem Cells.
Immunotherapy for Head and Neck Cancer.
Role of cytokines in head and neck squamous cell carcinoma.
The immune system and head and neck squamous cell carcinoma: from carcinogenesis to new therapeutic opportunities.
FAM3 Family as Prognostic Factors for Head and Neck Squamous Cell Carcinoma.
How Risk Factors Affect Head and Neck Squamous Cell Carcinoma (HNSCC) Tumor Immune Microenvironment (TIME): Their Influence on Immune Escape Mechanisms and Immunotherapy Strategy.
Immune Escape Mechanisms and Their Clinical Relevance in Head and Neck Squamous Cell Carcinoma.
Immunology and Immunotherapy of Head and Neck Cancer.
Immune deserts in head and neck squamous cell carcinoma: A review of challenges and opportunities for modulating the tumor immune microenvironment.
T-helper and T-regulatory cells modulation in head and neck squamous cell carcinoma.
Tumor microenvironment and immune evasion in head and neck squamous cell carcinoma.
RCAS-1 serum and tumor levels in head and neck squamous cell carcinoma.
Effects of checkpoint kinase 1 inhibition by prexasertib on the tumor immune microenvironment of head and neck squamous cell carcinoma.
Checkpoint immunotherapy in head and neck cancers.
At the Crossroads of Molecular Biology and Immunology: Molecular Pathways for Immunological Targeting of Head and Neck Squamous Cell Carcinoma.
Immunotherapy of head and neck cancer: Emerging clinical trials from a National Cancer Institute Head and Neck Cancer Steering Committee Planning Meeting.
Bioinformatic analysis identifies HPV-related tumor microenvironment remodeling prognostic biomarkers in head and neck squamous cell carcinoma.
HPV-induced oropharyngeal cancer, immune response and response to therapy.
The role of T-cells in head and neck squamous cell carcinoma: From immunity to immunotherapy.
Biological mechanisms of immune escape and implications for immunotherapy in head and neck squamous cell carcinoma.
The promise of immunotherapy in head and neck squamous cell carcinoma.
Costimulatory and coinhibitory immune checkpoint receptors in head and neck cancer: unleashing immune responses through therapeutic combinations.
Biology and immunology of cancer stem(-like) cells in head and neck cancer.
Cancer associated fibroblast derived gene signature determines cancer subtypes and prognostic model construction in head and neck squamous cell carcinomas.
Immuno-oncology in head and neck squamous cell carcinoma - a narrative review.



Causes and Consequences of HPV Integration in Head and Neck Squamous Cell Carcinomas: State of the Art.
The promise of immunotherapy in head and neck squamous cell carcinoma: combinatorial immunotherapy approaches.
Focus on TILs: prognostic significance of tumor infiltrating lymphocytes in head and neck cancers.
CircRNAs: Roles in regulating head and neck squamous cell carcinoma.
Immunotherapeutic Approaches to Head and Neck Cancer.
[The tumor microenvironment-relay station for prognosis and therapy response].
Immunotherapy for head and neck cancer: the future of treatment?
Immunotherapy: The future of cancer treatment.
Molecular and cellular processes underlying the hallmarks of head and neck cancer.
Immunotherapy in Head and Neck Squamous Cell Cancer.
Novel Treatment Options in Head and Neck Cancer.
Immunotherapy for head and neck cancer: latest developments and clinical potential.
Immune suppression in head and neck cancers: a review.
Immune escape mechanisms in head and neck squamous cell carcinoma and implication for new immunotherapy approach.
[Immunomodulation as innovative therapy for head and neck tumors : Current developments].
Immuno-oncology in head and neck squamous cell cancers: News from clinical trials, emerging predictive factors and unmet needs.
Toll-like receptor modulation in head and neck cancer.
CTLA4 blockade reduces immature myeloid cells in head and neck squamous cell carcinoma.
Prognostic capacity of Systemic Inflammation Response Index (SIRI) in patients with head and neck squamous cell carcinoma.
Involvement of HPV Infection in the Release of Macrophage Migration Inhibitory Factor in Head and Neck Squamous Cell Carcinoma.
Transforming growth factor- $\beta$ signaling in head and neck squamous cell carcinoma: Insights into cellular responses.
Integrating novel therapeutic monoclonal antibodies into the management of head and neck cancer.
[Clinical research on immune checkpoint and head and neck squamous cell carcinoma].
Immunogenetic Determinants of Susceptibility to Head and Neck Cancer in the Million Veteran Program Cohort.
Multomics data analyses to identify SLC25A17 as a novel biomarker to predict the prognosis and immune microenvironment in head and neck squamous cell carcinoma.
Peripheral Cytokine Levels Differ by HPV Status and Change Treatment-Dependently in Patients with Head and Neck Squamous Cell Carcinoma.
BASP1 is a prognostic biomarker associated with immunotherapeutic response in head and neck squamous cell carcinoma.
Stat3 orchestrates tumor development and progression: the Achilles' heel of head and neck cancers?
Tumor gene signatures that correlate with release of extracellular vesicles shape the immune landscape in head and neck squamous cell carcinoma.
Tadalafil augments tumor specific immunity in patients with head and neck squamous cell carcinoma.
USC-HN2, a new model cell line for recurrent oral cavity squamous cell carcinoma with immunosuppressive characteristics.
Genomic Landscapes of EBV-Associated Nasopharyngeal Carcinoma vs. HPV-Associated Head and Neck Cancer.

Mitigating SOX2-potentiated Immune Escape of Head and Neck Squamous Cell Carcinoma with a STING-inducing Nanosatellite Vaccine.
Chemokine-Cytokine Networks in the Head and Neck Tumor Microenvironment.
Circulating Tumour Cell Expression of Immune Markers as Prognostic and Therapeutic Biomarkers in Head and Neck Squamous Cell Carcinoma: A Systematic Review and Meta-Analysis.
A Neutrophil Extracellular Traps Signature Predicts the Clinical Outcomes and Immunotherapy Response in Head and Neck Squamous Cell Carcinoma.
Immunotherapy in head and neck cancer: evidence and perspectives.
14-3-3 $\zeta$ regulates immune response through Stat3 signaling in oral squamous cell carcinoma.
Targeted Therapy in Head and Neck Cancer: An Update on Current Clinical Developments in Epidermal Growth Factor Receptor-Targeted Therapy and Immunotherapies.
Impaired H3K36 methylation defines a subset of head and neck squamous cell carcinomas.
Distinct immune evasion in APOBEC-enriched, HPV-negative HNSCC.
Upregulated glycolysis correlates with tumor progression and immune evasion in head and neck squamous cell carcinoma.
Tailored immunotherapy for HPV positive head and neck squamous cell cancer.
Proteogenomic insights into the biology and treatment of HPV-negative head and neck squamous cell carcinoma.
The STAT3 pathway as a therapeutic target in head and neck cancer: Barriers and innovations.
HPV-Associated Head and Neck Cancer: Unique Features of Epidemiology and Clinical Management.
The role of granulocyte-macrophage colony-stimulating factor in head and neck cancer.
HPV: Molecular pathways and targets.
Identification of cuproptosis-related subtypes and characterization of the tumor microenvironment landscape in head and neck squamous cell carcinoma.
A novel immune signature predicts immunotherapy responsiveness and reveals the landscape of the tumor immune microenvironment in head and neck squamous cell carcinoma.
Immunomodulatory role of bitter melon extract in inhibition of head and neck squamous cell carcinoma growth.
Inhibition of SRC family kinases reduces myeloid-derived suppressor cells in head and neck cancer.
Clustering and machine learning-based integration identify cancer associated fibroblasts genes' signature in head and neck squamous cell carcinoma.
Plac1 Remodels the Tumor Immune Evasion Microenvironment and Predicts Therapeutic Response in Head and Neck Squamous Cell Carcinoma.
CD276 expression enables squamous cell carcinoma stem cells to evade immune surveillance.
Upregulated YTHDF1 associates with tumor immune microenvironment in head and neck squamous cell carcinomas.
Targeting LSD1 suppresses stem cell-like properties and sensitizes head and neck squamous cell carcinoma to PD-1 blockade.
Rationale for combined blockade of PD-1 and CTLA-4 in advanced head and neck squamous cell cancer—review of current data.
Tumor Immunity and Immunotherapy for HPV-Related Cancers.
Genomic Immune Evasion: Diagnostic and Therapeutic Opportunities in Head and Neck Squamous Cell Carcinomas.
Identification of a novel ceRNA network related to prognosis and immunity in HNSCC based on integrated bioinformatic investigation.
The prognostic impact of B7-H3 and B7-H4 in head and neck squamous cell carcinoma.

CD73 facilitates invadopodia formation and boosts malignancy of head and neck squamous cell carcinoma via the MAPK signaling pathway.
Down-regulation of the tumor suppressor miR-34a contributes to head and neck cancer by up-regulating the MET oncogene and modulating tumor immune evasion.
Expression of RCAS1 in esophageal squamous cell carcinoma is associated with a poor prognosis.
Oropharyngeal Carcinoma with a Special Focus on HPV-Related Squamous Cell Carcinoma.
Expression of RCAS1 and its function in human squamous cell carcinoma of the oral cavity.
Microbe-Mediated Activation of Toll-like Receptor 2 Drives PDL1 Expression in HNSCC.
The role of the tumor matrix environment in progression of head and neck cancer.
DNA-based immunotherapy for HPV-associated head and neck cancer.
Engineering Vaccines to Reprogram Immunity against Head and Neck Cancer.
CD200: association with cancer stem cell features and response to chemoradiation in head and neck squamous cell carcinoma.
Immunomodulatory aspects in the progression and treatment of oral malignancy.
Immune Evasion by Head and Neck Cancer: Foundations for Combination Therapy.
HPV16 drives cancer immune escape via NLRX1-mediated degradation of STING.
Expression of tumor-associated antigen RCAS1 and its possible involvement in immune evasion in oral squamous cell carcinoma.
Novel Immunotherapeutic Approaches in Head and Neck Cancer.
Immune responses against virus and tumor in cervical carcinogenesis: treatment strategies for avoiding the HPV-induced immune escape.
Molecular biology of anal squamous cell carcinoma: implications for future research and clinical intervention.
Enrichment and nLC-MS/MS Analysis of Head and Neck Cancer Mucinome Glycoproteins.
5'-Ectonucleotidase CD73/NT5E supports EGFR-mediated invasion of HPV-negative head and neck carcinoma cells.
Immune landscape and oncobiota in HPV-Associated Colorectal Cancer: an explorative study.
PD-L1 promotes head and neck squamous cell carcinoma cell growth through mTOR signaling.
The biology of combination immunotherapy in recurrent metastatic head and neck cancer.
[Correlation of RCAS1 expression to human papillomavirus 16 (HPV16) infection in cervical cancer].
RCAS1 expression in mobile tongue squamous cell carcinoma: an immunohistochemical study.
The role of interleukin 10 in human papilloma virus infection and progression to cervical carcinoma.
Defining the role of the JAK-STAT pathway in head and neck and thoracic malignancies: implications for future therapeutic approaches.
Human papillomavirus-related oropharyngeal cancer.
Immunosuppressive activity of cancer-associated fibroblasts in head and neck squamous cell carcinoma.
Apoptotic function of tumor-associated antigen RCAS1 in oral squamous cell carcinoma.
The Chemokines Initiating and Maintaining Immune Hot Phenotype Are Prognostic in ICB of HNSCC.
The Creation of the Suppressive Cancer Microenvironment in Patients with HPV-Positive Cervical Cancer.
The Roles of Programmed Cell Death Ligand-1/ Programmed Cell Death-1 (PD-L1/PD-1) in HPV-induced Cervical Cancer and Potential for their Use in Blockade Therapy.
The molecular genetics of laryngeal cancer.
Low levels of circulating invariant natural killer T cells predict poor clinical outcome in patients with head and neck squamous cell carcinoma.

HPV-transformed cells exhibit altered HMGB1-TLR4/MyD88-SARM1 signaling axis.
Genomic profiles and transcriptomic microenvironments in 2 patients with synchronous lung adenocarcinoma and lung squamous cell carcinoma: a case report.
Extrachromosomal DNA in HPV-Mediated Oropharyngeal Cancer Drives Diverse Oncogene Transcription.
Role of human papillomaviruses in carcinogenesis.
Putative Role of Circulating Human Papillomavirus DNA in the Development of Primary Squamous Cell Carcinoma of the Middle Rectum: A Case Report.
Significance of RCAS1 antigen in hepatocellular, cholangiocellular and pancreatic carcinomas.
Prognostic factors and new therapeutic approaches to cervical cancer.
Immunotherapeutic Approaches for the Treatment of HPV-Associated (Pre-)Cancer of the Cervix, Vulva and Penis.
Molecular Mechanism and Potential Targets for Blocking HPV-Induced Lesion Development.
The Changing Landscape of Systemic Treatment for Cervical Cancer: Rationale for Inhibition of the TGF- $\beta$ and PD-L1 Pathways.
Stress Keratin 17 Expression in Head and Neck Cancer Contributes to Immune Evasion and Resistance to Immune-Checkpoint Blockade.
The Blood Microenvironment Influences the Molecular Phenotypes of Circulating Tumor Cells in Head and Neck Squamous Cell Carcinoma.
Review of STAT3 (Signal Transducers and Activators of Transcription) in head and neck cancer.
A case report of complete pathological remission after chemotherapy in a patient with primary renal squamous cell carcinoma.
[The involvement of RCAS1 in creating a suppressive tumor microenvironment in patients with pharyngeal squamous cell carcinoma].
Management of locally advanced HPV-related oropharyngeal squamous cell carcinoma: where are we?
Modulation of antigen presenting cell functions during chronic HPV infection.
HPV upregulates MARCHF8 ubiquitin ligase and inhibits apoptosis by degrading the death receptors in head and neck cancer.
Immunosuppressive cytokine Interleukin-10 (IL-10) is up-regulated in high-grade CIN but not associated with high-risk human papillomavirus (HPV) at baseline, outcomes of HR-HPV infections or incident CIN in the LAMS cohort.
The papillomavirus E5 gene does not affect EGFR transcription and overall survival in cervical cancer.
Papillomavirus Immune Evasion Strategies Target the Infected Cell and the Local Immune System.
Immunopathogenesis of HPV-Associated Cancers and Prospects for Immunotherapy.
Recurrent HNSCC Harbor an Immunosuppressive Tumor Immune Microenvironment Suggesting Successful Tumor Immune Evasion.
Suppression of Antitumor Immune Responses by Human Papillomavirus through Epigenetic Downregulation of CXCL14.
Human Papillomaviruses-Associated Cancers: An Update of Current Knowledge.
The Role of Tumor Microenvironment in Invasion and Metastasis of Esophageal Squamous Cell Carcinoma.
STAT3 and Its Targeting Inhibitors in Oral Squamous Cell Carcinoma.
MicroRNAs in cervical cancer: evidences for a miRNA profile deregulated by HPV and its impact on radio-resistance.
Vaccines for cervical cancer.

High expression of tumor-associated antigen RCAS1 in pancreatic ductal adenocarcinoma is an unfavorable prognostic marker.
Expression and effects of high-mobility group box 1 in cervical cancer.
Activation of the Kynurenine Pathway in Human Malignancies Can Be Suppressed by the Cyclin-Dependent Kinase Inhibitor Dinaciclib.
The interplay between the vaginal microbiome and innate immunity in the focus of predictive, preventive, and personalized medical approach to combat HPV-induced cervical cancer.
Differential effect of GLUT1 overexpression on survival and tumor immune microenvironment of human papilloma virus type 16-positive and -negative cervical cancer.
Nimotuzumab for Patients With Inoperable Cancer of the Head and Neck.
Receptor-binding cancer antigen expressed on SiSo cells (RCAS1): a novel biomarker in the diagnosis and prognosis of human neoplasia.
Pretreatment peripheral blood leukocytes are independent predictors of survival in oral cavity cancer.
CMTM6 drives cisplatin resistance by regulating Wnt signaling through the ENO-1/AKT/GSK3 $\beta$ axis.
Novel therapeutic strategies to target RCAS1, which induces apoptosis via ectodomain shedding.
Immunosuppressive activity of proteases in cervical carcinoma.
Expression of RCAS1 in human gastric carcinoma: a potential mechanism of immune escape.
B7-H1-targeted immunotherapy for head and neck cancer.
A population-based case-control study of genetic variation in cytokine genes associated with risk of cervical and vulvar cancers.
Cancer-associated fibroblasts: An emerging target against esophageal squamous cell carcinoma.
RCAS1 expression: a potential prognostic marker for adenocarcinomas of the lung.
Increased RCAS1 expression is associated with advanced histopathological stage and poor prognosis in patients with gastric adenocarcinoma.
Administration of a vaccine composed of dendritic cells pulsed with premalignant oral lesion lysate to mice bearing carcinogen-induced premalignant oral lesions stimulates a protective immune response.
Immunoexpression of C4 Binding Protein in Oral Leukoplakia and Oral Squamous Cell Carcinoma.
Calcitonin gene-related peptide: A promising bridge between cancer development and cancer-associated pain in oral squamous cell carcinoma.
Clinico-pathological significance of RCAS1 expression in gliomas: a potential mechanism of tumor immune escape.
Identification of immunophenotypes in esophageal squamous cell carcinoma based on immune gene sets.
Evidence that the viral oncoproteins E6 and E7 of HPV induce the expression of a functional IL-2R on cervical cancer cells.
Analysis of the whole transcriptome from gingivo-buccal squamous cell carcinoma reveals deregulated immune landscape and suggests targets for immunotherapy.
Correlation of E6 and E7 levels in high-risk HPV16 type cervical lesions with CCL20 and Langerhans cells.
Induction of co-inhibitory molecule CTLA-4 by human papillomavirus E7 protein through downregulation of histone methyltransferase JHDM1B expression.
Genetic variations of IL-10: Identification of novel variations and evaluation of the impact of the SNPs/haplotype in the promoter region with the progression of Oral Squamous Cell Carcinoma in Indian population.
Genes and pathways monotonically dysregulated during progression from normal through leukoplakia to gingivo-buccal oral cancer.

Hypoxia-mediated activation of hypoxia-inducible factor-1 $\alpha$ in head and neck squamous cell carcinoma: A review.
Association between RCAS1 expression and microenvironmental immune cell death in uterine cervical cancer.
Decreased migration of myeloid dendritic cells through increased levels of C-reactive protein.
INPP5A/HLA-G1/IL-10/MMP-21 Axis in Progression of Esophageal Squamous Cell Carcinoma.
Manipulation of JAK/STAT Signalling by High-Risk HPVs: Potential Therapeutic Targets for HPV-Associated Malignancies.
Immune evasion in human papillomavirus-associated cervical cancer.
RCAS1 increases cell morphological changes in murine fibroblasts by reducing p38 phosphorylation.
Targeting the STAT pathway in head and neck cancer: recent advances and future prospects.
An individualized causal framework for learning intercellular communication networks that define microenvironments of individual tumors.
Head and Neck Carcinoma Immunotherapy: Facts and Hopes.
Expression of RCAS1 correlates with urothelial bladder cancer malignancy.
The analysis of receptor-binding cancer antigen expressed on SiSo cells (RCAS1) immunoreactivity within the microenvironment of the ovarian cancer lesion relative to the applied therapeutic strategy.
Evaluation of Elafin Immunohistochemical Expression as Marker of Cervical Cancer Severity.
Human papillomavirus E5 suppresses immunity via inhibition of the immunoproteasome and STING pathway.
Head and neck squamous cell carcinoma cell lines have an immunomodulatory effect on macrophages independent of hypoxia and toll-like receptor 9.
High-risk human papillomavirus E6 inhibits monocyte differentiation to Langerhans cells.
A Cocktail of Natural Compounds Holds Promise for New Immunotherapeutic Potential in Head and Neck Cancer.
Human Papillomaviruses and Skin Cancer.
Th9 cytokines curb cervical cancer progression and immune evasion.
Immunotherapy for cervical cancer: Can it do another lung cancer?
Early synergistic interactions between the HPV16-E7 oncoprotein and 17 $\beta$ -oestradiol for repressing the expression of Granzyme B in a cervical cancer model.
Receptor-Binding Cancer Antigen Expression in Thyroid Neoplasms: A Retrospective Study.
Galectin 3 expression in primary oral squamous cell carcinomas.
Comprehensive Analysis of the Effects of Genetic Ancestry and Genetic Characteristics on the Clinical Evolution of Oral Squamous Cell Carcinoma.
Receptor-binding cancer antigen expressed on SiSo cells (RCAS1) expression in human benign and malignant thyroid lesions.
Histone methylation antagonism drives tumor immune evasion in squamous cell carcinomas.
A comparative analysis of langerhans cell in oral epithelial dysplasia and oral squamous cell carcinoma using antibody CD-1a.
A Novel Role for the Soluble Isoform of CTLA-4 in Normal, Dysplastic and Neoplastic Oral and Oropharyngeal Epithelia.
Origin and immunoescape of uterine cervical cancer.
Leveraging innovative therapies with an evolving understanding of the molecular pathogenesis of penile squamous cell carcinoma.
Significant involvement of herpesvirus entry mediator in human esophageal squamous cell carcinoma.
Analysis of cell cycle arrest and apoptosis induced by RCAS1.

Expression profile of miRNAs computationally predicted to target PDL-1 in cervical tissues of different histology groups.
A pan-HER-targeted approach for recurrent or late-stage cervical cancer therapy: mechanisms, recent advances, and clinical prospects.
Association between B7-H1 and cervical cancer: B7-H1 impairs the immune response in human cervical cancer cells.
hrHPV E5 oncoprotein: immune evasion and related immunotherapies.
Multi-omics profiling identifies C1QA/B(+) macrophages with multiple immune checkpoints associated with esophageal squamous cell carcinoma (ESCC) liver metastasis.
Comparative analysis of RCAS1 level in neoplasms and placenta.
Macrophage migration inhibitory factor expression in cervical cancer.
Role of toll-like receptor 4 on the immune escape of human oral squamous cell carcinoma and resistance of cisplatin-induced apoptosis.
The association of RCAS1 serum concentration with the reversibility or irreversibility of the process of immune cytotoxic activity restriction during normal menstrual cycle, cancer relapse, and surgical treatment for various types of squamous cell carcinomas and adenocarcinomas.
Association of Fas-670 gene polymorphism with risk of cervical cancer in North Indian population.
Evaluation of the prognostic role of pSTAT3 expression in temporal bone squamous cell carcinoma.
SRSF10-mediated IL1RAP alternative splicing regulates cervical cancer oncogenesis via mIL1RAP-NF- $\kappa$ B-CD47 axis.
Serum metabolomic signatures of lymph node metastasis of esophageal squamous cell carcinoma.
Tumor cell phagocytosis (cannibalism) in lung cancer: possible biomarker for tumor immune escape and prognosis.
Expression of chemokine receptor CXCR4 in esophageal squamous cell and adenocarcinoma.
Cytoplasmic and membranous receptor-binding cancer antigens expressed on SiSo cells (RCAS1) immunoreactivity in epithelial ovarian cancer cells represent differing biological function of RCAS1.
Antigen peptide transporters are upregulated in squamous cell carcinoma of the oral tongue and show sex-specific associations with survival.
Deciphering nasopharyngeal carcinoma pathogenesis via proteomics.
Crosstalk between histone modification and DNA methylation orchestrates the epigenetic regulation of the costimulatory factors, Tim-3 and galectin-9, in cervical cancer.
Long Non-Cod-ing RNA Signature in Cervical Cancer.
Significance of Expression of Complement C4d in Esophageal Squamous Cell Carcinoma.
The role of gC1qR in regulating survival of human papillomavirus 16 oncogene-transfected cervical cancer cells.
Expression of CD44, CD44v9, ABCG2, CD24, Bmi-1 and ALDH1 in stage I and II oral squamous cell carcinoma and their association with clinicopathological factors.
The Polycomb proteins RING1B and EZH2 repress the tumoral pro-inflammatory function in metastasizing primary cutaneous squamous cell carcinoma.
Characterization of cluster of differentiation 47 expression and its potential as a therapeutic target in esophageal squamous cell cancer.
Dendritic cell subsets and immunological milieu in inflammatory human papilloma virus-related skin lesions.
The prognostic value of tumor-stromal ratio combined with TNM staging system in esophagus squamous cell carcinoma.
TIM-3 Participates in the Invasion and Metastasis of Nasopharyngeal Carcinoma via SMAD7/SMAD2/SNAIL1 Axis-Mediated Epithelial-Mesenchymal Transition.

Immune evasion before tumour invasion in early lung squamous carcinogenesis.
PTX3 gene activation in EGF-induced head and neck cancer cell metastasis.
Downregulation of LNMA5 orchestrates partial EMT and immune escape from macrophage phagocytosis to promote lymph node metastasis of cervical cancer.
Selective in situ protein expression profiles correlate with distinct phenotypes of basal cell carcinoma and squamous cell carcinoma of the skin.
High level of galectin-1 expression is a negative prognostic predictor of recurrence in laryngeal squamous cell carcinomas.
LINE-1 Retrotransposition Promotes the Development and Progression of Lung Squamous Cell Carcinoma by Disrupting the Tumor-Suppressor Gene FGGY.
Overexpression and interactions of interleukin-10, transforming growth factor beta, and vascular endothelial growth factor in esophageal squamous cell carcinoma.
Key Genetic Determinants Driving Esophageal Squamous Cell Carcinoma Initiation and Immune Evasion.
Using the R Package Spatstat to Assess Inhibitory Effects of Microregional Hypoxia on the Infiltration of Cancers of the Head and Neck Region by Cytotoxic T Lymphocytes.
Repression of Human Papillomavirus Oncogene Expression under Hypoxia Is Mediated by PI3K/mTORC2/AKT Signaling.
Epigenomic dysregulation-mediated alterations of key biological pathways and tumor immune evasion are hallmarks of gingivo-buccal oral cancer.
Protein profiling and identification of modulators regulated by the E7 oncogene in the C33A cell line by proteomics and genomics.
Immune Surveillance in Clinical Regression of Preinvasive Squamous Cell Lung Cancer.
Immunohistochemical evaluation of galectin-3 expression in oral squamous cell carcinoma, oral leukoplakia and normal mucosa.
Comprehensive Exome Analysis of Immunocompetent Metastatic Head and Neck Cancer Models Reveals Patient Relevant Landscapes.
TNPO1-Mediated Nuclear Import of FUBP1 Contributes to Tumor Immune Evasion by Increasing NRP1 Expression in Cervical Cancer.
Prognostic Significance of Serine-Phosphorylated STAT3 Expression in pT1-T2 Oral Tongue Carcinoma.
Meet the authors: Yinglu Li, Elizabeth Goldberg, and Xiao Chen.
Correction to: Clinical relevance of the tumor microenvironment and immune escape of oral squamous cell carcinoma.
Protecting Tumors by Preventing Human Papilloma Virus Antigen Presentation: Insights from Emerging Bioinformatics Algorithms.
Inactivation of interferon regulatory factor-1 tumor suppressor protein by HPV E7 oncoprotein. Implication for the E7-mediated immune evasion mechanism in cervical carcinogenesis.
Alterations of dendritic cell subsets in the peripheral circulation of patients with cervical carcinoma.
Cell-surface density of complement restriction factors (CD46, CD55, and CD59): oral squamous cell carcinoma versus other solid tumors.
Role of deltaNp63(pos)CD44v(pos) cells in the development of N-nitroso-tris-chloroethylurea-induced peripheral-type mouse lung squamous cell carcinomas.
Who should be treated with pembrolizumab in addition to standard of care in advanced cervical cancer?
Elevated pleural fluid RCAS1 is a diagnostic marker and outcome predictor in lung cancer patients.



Case Report: Combined Intra-Lesional IL-2 and Topical Imiquimod Safely and Effectively Clears Multi-Focal, High Grade Cutaneous Squamous Cell Cancer in a Combined Liver and Kidney Transplant Patient.
HPV-16 E5 oncoprotein upregulates lipid raft components caveolin-1 and ganglioside GM1 at the plasma membrane of cervical cells.
The upregulated expression of RFC4 and GMPS mediated by DNA copy number alteration is associated with the early diagnosis and immune escape of ESCC based on a bioinformatic analysis.
Down modulation of IL-18 expression by human papillomavirus type 16 E6 oncogene via binding to IL-18.
CD200 upregulation in vascular endothelium surrounding cutaneous squamous cell carcinoma.
Immunoregulatory molecule B7-H1 (CD274) contributes to skin carcinogenesis.
HPV11E7 inhibits IMQ-induced chemokine and colony-stimulating factor production in keratinocytes.
Evorpacept alone and in combination with pembrolizumab or trastuzumab in patients with advanced solid tumours (ASPEN-01): a first-in-human, open-label, multicentre, phase 1 dose-escalation and dose-expansion study.
Expression of the complement regulatory proteins decay accelerating factor (DAF, CD55), membrane cofactor protein (MCP, CD46) and CD59 in the normal human uterine cervix and in premalignant and malignant cervical disease.
Porphyromonas gingivalis Promotes Immuno-evasion of Oral Cancer by Protecting Cancer from Macrophage Attack.
IDO1 Expression Increased After Neoadjuvant Therapy Predicts Poor Pathologic Response and Prognosis in Esophageal Squamous Cell Carcinoma.
Immunosurveillance and Regression in the Context of Squamous Pulmonary Premalignancy.
Fusobacterium nucleatum and oral cancer: a critical review.
Ultraviolet light downregulates CD95 ligand and TRAIL receptor expression facilitating actinic keratosis and squamous cell carcinoma formation.
A new option for the treatment of condyloma acuminatum in the male urethra: Multimodal ultrasound image-guided scraping and photodynamic therapy (USP).
Effects of Cisplatin Combined with Metformin on Proliferation and Apoptosis of Nasopharyngeal Carcinoma Cells.
Falkner's needling technique for the treatment of warts: Minimum investment, maximum benefit.
Expression of selected regulatory molecules on the CD83+ monocyte-derived dendritic cells generated from patients with laryngeal cancer and their clinical significance.
Cell cannibalism by malignant neoplastic cells: three cases in dogs and a literature review.
Rediscovery of NF- $\kappa$ B signaling in nasopharyngeal carcinoma: How genetic defects of NF- $\kappa$ B pathway interplay with EBV in driving oncogenesis?
iTRAQ-based analysis for the identification of MARCH8 targets in human esophageal squamous cell carcinoma.
USF1 regulated circPRDM4 modulates tumorigenesis and immune escape in chemoresistant cervical cancer.
Loss of Cxcr4 in Endometriosis Reduces Proliferation and Lesion Number while Increasing Intraepithelial Lymphocyte Infiltration.

Table A2-47, Cluster 46

Cluster 46 focuses on PD-L1 expression in advanced cancers, especially its contribution to immune escape (842)
Significance of PD-L1 clones and C-MET expression in hepatocellular carcinoma.
Butein suppresses PD-L1 expression via downregulating STAT1 in non-small cell lung cancer.
A mechanistic systems pharmacology modeling platform to investigate the effect of PD-L1 expression heterogeneity and dynamics on the efficacy of PD-1 and PD-L1 blocking antibodies in cancer.
Temporally Distinct PD-L1 Expression by Tumor and Host Cells Contributes to Immune Escape.
Frequencies and expression levels of programmed death ligand 1 (PD-L1) in circulating tumor RNA (ctRNA) in various cancer types.
Cancer-associated oxidoreductase ERO1- $\alpha$ promotes immune escape through up-regulation of PD-L1 in human breast cancer.
Generation, secretion and degradation of cancer immunotherapy target PD-L1.
Tumor Suppressor microRNAs Contribute to the Regulation of PD-L1 Expression in Malignant Pleural Mesothelioma.
Regulation of PD-L1: a novel role of pro-survival signalling in cancer.
Expression of Programmed Death-Ligand 1 and Programmed Death-1 in Patients with Extramammary Paget's Disease.
PD-L1 expression in advanced NSCLC: Insights into risk stratification and treatment selection from a systematic literature review.
Regulation of PD-L1 expression on murine tumor-associated monocytes and macrophages by locally produced TNF- $\alpha$ .
Programmed Death Ligand 1 Indicates Pre-Existing Adaptive Immune Response by Tumor-Infiltrating CD8(+) T Cells in Non-Small Cell Lung Cancer.
USP2 promotes tumor immune evasion via deubiquitination and stabilization of PD-L1.
Immunohistochemical expression of programmed death-ligand 1 and CD8 in glioblastomas.
Expression of programmed cell death-ligand 1 and its correlation with clinical outcomes in gliomas.
Upregulation of PD-L1 expression in breast cancer cells through the formation of 3D multicellular cancer aggregates under different chemical and mechanical conditions.
Deubiquitination and Stabilization of PD-L1 by USP21.
Expression of Programmed Cell Death Ligand 1 and Associated Lymphocyte Infiltration in Olfactory Neuroblastoma.
Programmed death ligand-1 and MET co-expression is a poor prognostic factor in gastric cancers after resection.
THADA drives Golgi residency and upregulation of PD-L1 in cancer cells and provides promising target for immunotherapy.
PD-L1 Expression in Extramammary Paget Disease: A Case Series.
Association Between PD-L1 and HPV Status and the Prognostic Value of PD-L1 in Oropharyngeal Squamous Cell Carcinoma.
Sensitive detection of PD-L1 expression on circulating epithelial tumor cells (CETCs) could be a potential biomarker to select patients for treatment with PD-1/PD-L1 inhibitors in early and metastatic solid tumors.
Programmed Death-Ligand 1 Expression in Breast Cancer Patients: Clinicopathological Associations from a Single-Institution Study.
Membranous and Cytoplasmic Expression of PD-L1 in Ovarian Cancer Cells.
Inhibiting PD-L1 palmitoylation enhances T-cell immune responses against tumours.
Highly Activated PD-1/PD-L1 Pathway in Gastric Cancer with PD-L1 Expression.

Combined Radiotherapy and Anti-PD-L1 Antibody Synergistically Enhances Antitumor Effect in Non-Small Cell Lung Cancer.
PD-L1 expression in papillary renal cell carcinoma.
Tumor PD-L1 expression is correlated with increased TILs and poor prognosis in penile squamous cell carcinoma.
The DPY30-H3K4me3 Axis-Mediated PD-L1 Expression in Melanoma.
The "Inside" Story on Tumor-Expressed PD-L1.
Evaluation of PD-L1 expression on vortex-isolated circulating tumor cells in metastatic lung cancer.
Tumor cells versus host immune cells: whose PD-L1 contributes to PD-1/PD-L1 blockade mediated cancer immunotherapy?
Programmed Death Ligand 1 Expression Among 700 Consecutive Endometrial Cancers: Strong Association With Mismatch Repair Protein Deficiency.
The Role of Oncogenes and Redox Signaling in the Regulation of PD-L1 in Cancer.
Analysis of expression of programmed cell death 1 ligand 1 (PD-L1) in malignant pleural mesothelioma (MPM).
[Regulatory Mechanisms of PD-L1 Expression and Its Role in Immune Evasion].
Canagliflozin primes antitumor immunity by triggering PD-L1 degradation in endocytic recycling.
Expression patterns of programmed death-1 and programmed death-1 ligand-1 on T cells in gastric cancer.
Dysregulation of PD-L1 by UFMylation imparts tumor immune evasion and identified as a potential therapeutic target.
Interferon- $\beta$ signal may up-regulate PD-L1 expression through IRF9-dependent and independent pathways in lung cancer cells.
PD-L1 expression is increased in a subset of basal type breast cancer cells.
The nuclear transportation of PD-L1 and the function in tumor immunity and progression.
Demethylase JMJD2D induces PD-L1 expression to promote colorectal cancer immune escape by enhancing IFNGR1-STAT3-IRF1 signaling.
RAS-Mitogen-Activated Protein Kinase Signal Is Required for Enhanced PD-L1 Expression in Human Lung Cancers.
Circ_0000052/miR-382-3p axis induces PD-L1 expression and regulates cell proliferation and immune evasion in head and neck squamous cell carcinoma.
Exosomal PD-L1 promotes tumor growth through immune escape in non-small cell lung cancer.
WNT signaling modulates PD-L1 expression in the stem cell compartment of triple-negative breast cancer.
Non-cytomembrane PD-L1: An atypical target for cancer.
A humanized antibody for imaging immune checkpoint ligand PD-L1 expression in tumors.
MLL3 enhances the transcription of PD-L1 and regulates anti-tumor immunity.
Tumor-Intrinsic PD-L1 Signaling in Cancer Initiation, Development and Treatment: Beyond Immune Evasion.
c-Met up-regulates the expression of PD-L1 through MAPK/NF- $\kappa$ Bp65 pathway.
Absence of PD-L1 on tumor cells is associated with reduced MHC I expression and PD-L1 expression increases in recurrent serous ovarian cancer.
Regulation of PD-L1 expression in the tumor microenvironment.
Sorting nexin 6 interacts with Cullin3 and regulates programmed death ligand 1 expression.
KDM1A Promotes Immunosuppression in Hepatocellular Carcinoma by Regulating PD-L1 through Demethylating MEF2D.

Regulation of programmed death-ligand 1 expression in response to DNA damage in cancer cells: Implications for precision medicine.
Programmed death-ligand 1 is a promising blood marker for predicting tumor progression and prognosis in patients with gastric cancer.
Light-controlled elimination of PD-L1+ cells.
Critical role of PD-L1 expression on non-tumor cells rather than on tumor cells for effective anti-PD-L1 immunotherapy in a transplantable mouse hematopoietic tumor model.
FAM83A drives PD-L1 expression via ERK signaling and FAM83A/PD-L1 co-expression correlates with poor prognosis in lung adenocarcinoma.
Overall survival and PD-L1 expression in metastasized malignant melanoma.
Prognostic value of PD -L1 expression in patients with primary solid tumors.
Immune regulation of canine tumour and macrophage PD-L1 expression.
Association of PD-L1 expression and PD-L1 gene polymorphism with poor prognosis in lung adenocarcinoma and squamous cell carcinoma.
The opportunities and challenges in immunotherapy: Insights from the regulation of PD-L1 in cancer cells.
Glucose metabolism involved in PD-L1-mediated immune escape in the malignant kidney tumour microenvironment.
Interaction of PD-L1 on tumor cells with PD-1 on tumor-specific T cells as a mechanism of immune evasion: implications for tumor immunotherapy.
Determination of PD-L1 expression in effusions from mesothelioma by immuno-cytochemical staining.
Combining protein arginine methyltransferase inhibitor and anti-programmed death-ligand-1 inhibits pancreatic cancer progression.
Identification of the Cell-Intrinsic and -Extrinsic Pathways Downstream of EGFR and IFN $\gamma$ That Induce PD-L1 Expression in Head and Neck Cancer.
Current methods and emerging approaches for detection of programmed death ligand 1.
Recent advances of molecular mechanisms of regulating PD-L1 expression in melanoma.
The miR-3127-5p/p-STAT3 axis up-regulates PD-L1 inducing chemoresistance in non-small-cell lung cancer.
Circulating PD-L1 in NSCLC patients and the correlation between the level of PD-L1 expression and the clinical characteristics.
Increased expression of PD-L1 in endometrial cancer stem-like cells is regulated by hypoxia.
Atorvastatin Attenuates Programmed Death Ligand-1 (PD-L1) Induction in Human Hepatocellular Carcinoma Cells.
PD-L1 Expression and Intratumoral Heterogeneity Across Breast Cancer Subtypes and Stages: An Assessment of 245 Primary and 40 Metastatic Tumors.
Beyond Cancer: Regulation and Function of PD-L1 in Health and Immune-Related Diseases.
PD-L1 Tumor Cell Expression in Upper Tract Urothelial Carcinomas is Associated With Higher Pathologic Stage.
PD-L1 (SP142) expression in neoplastic cells predicts a poor prognosis for patients with intravascular large B-cell lymphoma treated with rituximab-based multi-agent chemotherapy.
Differential Expression of Immune-Regulatory Genes Associated with PD-L1 Display in Melanoma: Implications for PD-1 Pathway Blockade.
A pooled analysis of the prognostic value of PD-L1 in melanoma: evidence from 1062 patients.
Synthesis and Biologic Evaluation of a Novel (18)F-Labeled Adnectin as a PET Radioligand for Imaging PD-L1 Expression.
Hydroxyproline metabolism enhances IFN- $\gamma$ -induced PD-L1 expression and inhibits autophagic flux.

[Radiotherapy and PD-L1 Expression].
Foci of Programmed Cell Death-Ligand 1 (PD-L1)-positive Tumor Areas With Tumor-infiltrating Leukocytes (TILs) Evocative of a PD-1/PD-L1-related Adaptive Immune Resistance are Frequent in Merkel Cell Carcinoma.
NPM1 upregulates the transcription of PD-L1 and suppresses T cell activity in triple-negative breast cancer.
Interleukin 6 regulates the expression of programmed cell death ligand 1 in thyroid cancer.
What Do We Have to Know about PD-L1 Expression in Prostate Cancer? A Systematic Literature Review. Part 5: Epigenetic Regulation of PD-L1.
PD-L1 expression in medulloblastoma: an evaluation by subgroup.
Relationship between PD-L1 expression, CD8+ T-cell infiltration and prognosis in intrahepatic cholangiocarcinoma patients.
Nasopharyngeal cancer cell-derived exosomal PD-L1 inhibits CD8+ T-cell activity and promotes immune escape.
High expression of PD-L1 in lung cancer may contribute to poor prognosis and tumor cells immune escape through suppressing tumor infiltrating dendritic cells maturation.
High PD-L1 expression is associated with stage IV disease and poorer overall survival in 186 cases of small cell lung cancers.
PD-L1 degradation pathway and immunotherapy for cancer.
Expression of Programmed Death Ligand 1 in Breast Cancer in Mexican Women.
ICSBP-induced PD-L1 enhances osteosarcoma cell growth.
Interleukin-22 promotes PD-L1 expression via STAT3 in colon cancer cells.
Anti-PD-L1 Antibody Enhances T Cell Immune Responses and Reduces Resistance of Breast Cancer Cells to Radiotherapy.
Alpha5 nicotinic acetylcholine receptor mediated immune escape of lung adenocarcinoma via STAT3/Jab1-PD-L1 signalling.
Mutant APC promotes tumor immune evasion via PD-L1 in colorectal cancer.
PD-L1 expression in colorectal cancer is associated with microsatellite instability, BRAF mutation, medullary morphology and cytotoxic tumor-infiltrating lymphocytes.
Current insight into the regulation of PD-L1 in cancer.
Lipopolysaccharide facilitates immune escape of hepatocellular carcinoma cells via m6A modification of lncRNA MIR155HG to upregulate PD-L1 expression.
Targeting tumor-intrinsic PD-L1 suppresses the progression and aggressiveness of head and neck cancer by inhibiting GSK3β-dependent Snail degradation.
Cancer-Associated Fibroblasts Promote the Upregulation of PD-L1 Expression Through Akt Phosphorylation in Colorectal Cancer.
Long noncoding RNA HITT coordinates with RGS2 to inhibit PD-L1 translation in T cell immunity.
Marked Global DNA Hypomethylation Is Associated with Constitutive PD-L1 Expression in Melanoma.
Expression of STAT1 is positively correlated with PD-L1 in human ovarian cancer.
High PD-L1 expression is associated with unfavorable clinical features in myelodysplastic neoplasms.
[The Expression and Biological Significance of PD-L1 on Lung Cancer Cell Lines.].
Neddylation inhibition upregulates PD-L1 expression and enhances the efficacy of immune checkpoint blockade in glioblastoma.
[Expression of Programmed Death Ligand-1 (PD-L1) in Human Acute Leukemia and Its Clinical Significance].
Evaluation of PD-L1 Expression and Associated Tumor-Infiltrating Lymphocytes in Laryngeal Squamous Cell Carcinoma.

Pan-cancer analysis of copy number changes in programmed death-ligand 1 (PD-L1, CD274) - associations with gene expression, mutational load, and survival.
PD-L1 expression in tumor tissue and peripheral blood of patients with oral squamous cell carcinoma.
PD-L1 Expression Is Associated with Tumor FOXP3(+) Regulatory T-Cell Infiltration of Breast Cancer and Poor Prognosis of Patient.
Prognostic implications of PD-L1 expression in patients with soft tissue sarcoma.
PD-L1 expression associated with worse survival outcome in malignant pleural mesothelioma.
Loss of VGLL4 suppresses tumor PD-L1 expression and immune evasion.
Tumor PD-L1 Induction by Resveratrol/Piceatannol May Function as a Search, Enhance, and Engage ("SEE") Signal to Facilitate the Elimination of "Cold, Non-Responsive" Low PD-L1-Expressing Tumors by PD-L1 Blockade.
Role of tumor microenvironment in the regulation of PD-L1: A novel role in resistance to cancer immunotherapy.
TOPK mediates immune evasion of renal cell carcinoma via upregulating the expression of PD-L1.
PD-L1 expression of the residual tumor serves as a prognostic marker in local advanced breast cancer after neoadjuvant chemotherapy.
Immunohistochemical expression of PD-L1 and MDR1 in breast tumors: association with clinicopathological parameters and treatment outcome.
Association of programmed cell death ligand 1 and circulating lymphocytes with risk of venous thromboembolism in patients with glioma.
Aberrant PD-L1 expression through 3'-UTR disruption in multiple cancers.
PD-L1 in squamous cell carcinoma of the oral tongue shows gender-specific association with prognosis.
PD-L1 is a direct target of cancer-FOXP3 in pancreatic ductal adenocarcinoma (PDAC), and combined immunotherapy with antibodies against PD-L1 and CCL5 is effective in the treatment of PDAC.
CMTM6 maintains the expression of PD-L1 and regulates anti-tumour immunity.
PD-L1 Expression in Clear Cell Renal Cell Carcinoma: An Analysis of Nephrectomy and Sites of Metastases.
COX2/mPGES1/PGE2 pathway regulates PD-L1 expression in tumor-associated macrophages and myeloid-derived suppressor cells.
Research progress on the intrinsic non-immune function of PD-L1 in tumors (Review).
PD-L1 Acts as a Promising Immune Marker to Predict the Response to Neoadjuvant Chemotherapy in Breast Cancer Patients.
PD-L1 Expression and Combined Status of PD-L1/PD-1-Positive Tumor Infiltrating Mononuclear Cell Density Predict Prognosis in Glioblastoma Patients.
Characterization of PD-L1 Expression and Associated T-cell Infiltrates in Metastatic Melanoma Samples from Variable Anatomic Sites.
Interferon-γ-induced PD-L1 surface expression on human oral squamous carcinoma via PKD2 signal pathway.
Recent Advancements in the Mechanisms Underlying Resistance to PD-1/PD-L1 Blockade Immunotherapy.
Programmed death ligand 1 (PD-L1) expression influences the immune-tolerogenic microenvironment in antiretroviral therapy-refractory Kaposi's sarcoma: A pilot study.
PD-L1-Mediated Immunosuppression in Oral Squamous Cell Carcinoma: Relationship With Macrophage Infiltration and Epithelial to Mesenchymal Transition Markers.

Clinical implication of programmed cell death-1 ligand-1 expression in tonsillar squamous cell carcinoma in association with intratumoral heterogeneity, human papillomavirus, and epithelial-to-mesenchymal transition.
Cooperativity of HOXA5 and STAT3 Is Critical for HDAC8 Inhibition-Mediated Transcriptional Activation of PD-L1 in Human Melanoma Cells.
PD-L1 on tumor cells is sufficient for immune evasion in immunogenic tumors and inhibits CD8 T cell cytotoxicity.
METTL3/IGF2BP3 axis inhibits tumor immune surveillance by upregulating N(6)-methyladenosine modification of PD-L1 mRNA in breast cancer.
CPI-203 improves the efficacy of anti-PD-1 therapy by inhibiting the induced PD-L1 overexpression in liver cancer.
PD-L1 Expression Is a Prognostic Factor in Patients with Thoracic Esophageal Cancer Treated Without Adjuvant Chemotherapy.
TERC suppresses PD-L1 expression by downregulating RNA binding protein HuR.
Platelet PD-L1 suppresses anti-cancer immune cell activity in PD-L1 negative tumors.
Programmed cell death 1 ligand 1 (PD-L1) expression is associated with poor prognosis of malignant pleural mesothelioma patients with good performance status.
HOXA11-AS1 Promotes PD-L1-Mediated Immune Escape and Metastasis of Hypopharyngeal Carcinoma by Facilitating PTBP1 and FOSL1 Association.
PD-L1 translocation to the plasma membrane enables tumor immune evasion through MIB2 ubiquitination.
Identification of vitamin B6 as a PD-L1 suppressor and an adjuvant for cancer immunotherapy.
Mechanisms Controlling PD-L1 Expression in Cancer.
Brazilein inhibits epithelial-mesenchymal transition (EMT) and programmed death ligand 1 (PD-L1) expression in breast cancer cells.
Programmed Death Ligand 1 (PD-L1) Expression and CD8 + Tumor-infiltrating Lymphocyte-based Tumor Immune Microenvironment Classification in Gynecologic Carcinosarcoma: Prognostic Impact and Implications for Therapy.
Programmed Death Ligand 1 (PD-L1) Tumor Expression Is Associated with a Better Prognosis and Diabetic Disease in Triple Negative Breast Cancer Patients.
Comprehensive Immunohistochemical Study of Programmed Cell Death Ligand 1 (PD-L1): Analysis in 5536 Cases Revealed Consistent Expression in Trophoblastic Tumors.
Biochanin A Suppresses Tumor Progression and PD-L1 Expression via Inhibiting ZEB1 Expression in Colorectal Cancer.
The PD-L1 interactome demonstrates bidirectional signaling coordinating immune suppression and cancer progression in HNSCC.
PD-L1 diagnostic tests: a systematic literature review of scoring algorithms and test-validation metrics.
Programmed death ligand-1 and CD8 tumor-infiltrating lymphocytes (TILs) as prognostic predictors in ovarian high-grade serous carcinoma (HGSC).
Identification and characterization of an alternative cancer-derived PD-L1 splice variant.
BRD4 inhibition suppresses PD-L1 expression in triple-negative breast cancer.
FGFR2 Promotes Expression of PD-L1 in Colorectal Cancer via the JAK/STAT3 Signaling Pathway.
Sequential Tracking of PD-L1 Expression and RAD50 Induction in Circulating Tumor and Stromal Cells of Lung Cancer Patients Undergoing Radiotherapy.
Recent Advancement of PD-L1 Detection Technologies and Clinical Applications in the Era of Precision Cancer Therapy.

αvβ3-integrin regulates PD-L1 expression and is involved in cancer immune evasion.
DENR controls JAK2 translation to induce PD-L1 expression for tumor immune evasion.
Prognostic effect of different PD-L1 expression patterns in squamous cell carcinoma and adenocarcinoma of the cervix.
IFN-γ induces PD-L1 through p38/JNK/ERK signaling pathways and counteracts the tumor promoting effect mediated by PD-L1 in Glioblastoma.
Syntenin1/MDA-9 (SDCBP) induces immune evasion in triple-negative breast cancer by upregulating PD-L1.
Dihydropyridine Calcium Channel Blockers Suppress the Transcription of PD-L1 by Inhibiting the Activation of STAT1.
Tumor cells PD-L1 expression as a favorable prognosis factor in nasopharyngeal carcinoma patients with pre-existing intratumor-infiltrating lymphocytes.
PD-L1 gene alterations identify a subset of diffuse large B-cell lymphoma harboring a T-cell-inflamed phenotype.
Integrin-Linked Kinase Expression Characterizes the Immunosuppressive Tumor Microenvironment in Colorectal Cancer and Regulates PD-L1 Expression and Immune Cell Cytotoxicity.
Promoting anti-tumor immunity by targeting TMUB1 to modulate PD-L1 polyubiquitination and glycosylation.
Regulatory mechanisms of PD-L1 expression in cancer cells.
Membrane-Associated RING-CH 8 Functions as a Novel PD-L1 E3 Ligase to Mediate PD-L1 Degradation Induced by EGFR Inhibitors.
Oncogenic RAS Signaling Promotes Tumor Immuno-resistance by Stabilizing PD-L1 mRNA.
Radiation-induced PD-L1 expression in tumor and its microenvironment facilitates cancer-immune escape: a narrative review.
BRD4-IRF1 axis regulates chemoradiotherapy-induced PD-L1 expression and immune evasion in non-small cell lung cancer.
Juxtacrine Signaling Inhibits Antitumor Immunity by Upregulating PD-L1 Expression.
CD137 ligand feedback upregulates PD-L1 expression on lung cancer via T cell production of IFN-γ.
Analysis of PD-L1 expression in trophoblastic tissues and tumors.
Associations among pretreatment tumor necrosis and the expression of HIF-1α and PD-L1 in advanced oral squamous cell carcinoma and the prognostic impact thereof.
Contribution of PD-L1 to oncogenesis of lymphoma and its RNAi-based targeting therapy.
PD-L1 Expression in Mastocytosis.
Dickkopf-1 drives tumor immune evasion by inducing PD-L1 expression in hepatocellular carcinoma.
Cyclic adenosine monophosphate/phosphodiesterase 4 pathway associated with immune infiltration and PD-L1 expression in lung adenocarcinoma cells.
S100A14: A novel negative regulator of cancer stemness and immune evasion by inhibiting STAT3-mediated programmed death-ligand 1 expression in colorectal cancer.
[Regulation of PD-L1 posttranslational modification and its application progress in tumor immunotherapy].
Tumor programmed cell death ligand 1 expression correlates with nodal metastasis in patients with cutaneous squamous cell carcinoma of the head and neck.
Immunoblotting Analysis of Intracellular PD-L1 Levels in Interferon-γ-Treated Ovarian Cancer Cells Stably Transfected with Bcl3 shRNA.
The IFN-γ/PD-L1 axis between T cells and tumor microenvironment: hints for glioma anti-PD-1/PD-L1 therapy.



What Do We Have to Know about PD-L1 Expression in Prostate Cancer? A Systematic Literature Review. Part 7: PD-L1 Expression in Liquid Biopsy.
Gastric cancer mesenchymal stem cells derived IL-8 induces PD-L1 expression in gastric cancer cells via STAT3/mTOR-c-Myc signal axis.
Epigenetically modified AP-2 $\alpha$ by DNA methyltransferase facilitates glioma immune evasion by upregulating PD-L1 expression.
Cytidine Deaminase APOBEC3A Regulates PD-L1 Expression in Cancer Cells in a JNK/c-JUN-Dependent Manner.
Prognostic Value of Programmed Death Ligand 1 and Programmed Death 1 Expression in Thymic Carcinoma.
Transforming growth factor beta orchestrates PD-L1 enrichment in tumor-derived exosomes and mediates CD8 T-cell dysfunction regulating early phosphorylation of TCR signalome in breast cancer.
Expression of tumor-associated macrophages and PD-L1 in patients with hepatocellular carcinoma and construction of a prognostic model.
LRPPRC facilitates tumor progression and immune evasion through upregulation of m(6)A modification of PD-L1 mRNA in hepatocellular carcinoma.
Obesity and metabolic syndrome related macrophage promotes PD-L1 expression in TNBC through IL6/JAK/STAT pathway and can be reversed by telmisartan.
PD-L1.
microRNA-378a-3p regulates the progression of hepatocellular carcinoma by regulating PD-L1 and STAT3.
Regulation of post-translational modification of PD-L1 and advances in tumor immunotherapy.
PD-L1 amplification is associated with an immune cell rich phenotype in squamous cell cancer of the lung.
The roles of programmed death ligand 1 in virus-associated cancers.
PD-L1 immunohistochemistry: Clones, cutoffs, and controversies.
A polymorphism in the promoter region of PD-L1 serves as a binding-site for SP1 and is associated with PD-L1 overexpression and increased occurrence of gastric cancer.
A Novel Anti-PD-L1 Vaccine for Cancer Immunotherapy and Immunoprevention.
Programmed Death-Ligand 1 (PD-L1) Positivity and Factors Associated with Poor Prognosis in Patients with Gastric Cancer: An Umbrella Meta-Analysis.
Programmed Death Ligand 1: A Poor Prognostic Marker in Endometrial Carcinoma.
Expression of Programmed Death Ligand 1 in Penile Cancer is of Prognostic Value and Associated with HPV Status.
ARIH1 signaling promotes anti-tumor immunity by targeting PD-L1 for proteasomal degradation.
Clinical significance of programmed death 1/programmed death ligand 1 pathway in gastric neuroendocrine carcinomas.
Clinical Implications of Exosomal PD-L1 in Cancer Immunotherapy.
Biochemical Aspects of PD-L1 Regulation in Cancer Immunotherapy.
Chemotherapy induces tumor immune evasion by upregulation of programmed cell death ligand 1 expression in bone marrow stromal cells.
Fructose-1,6-bisphosphatase loss modulates STAT3-dependent expression of PD-L1 and cancer immunity.
Level of circulating PD-L1 expression in patients with advanced gastric cancer and its clinical implications.
Inhibition of PAI-1 Blocks PD-L1 Endocytosis and Improves the Response of Melanoma Cells to Immune Checkpoint Blockade.

Identification, binding, and structural characterization of single domain anti-PD-L1 antibodies inhibitory of immune regulatory proteins PD-1 and CD80.
Disruption of RBMS3 suppresses PD-L1 and enhances antitumor immune activities and therapeutic effects of auranofin against triple-negative breast cancer.
Tumor cell PD-L1 expression is a strong predictor of unfavorable prognosis in immune checkpoint therapy-naive clear cell renal cell cancer.
Radiotherapy Upregulates Programmed Death Ligand-1 through the Pathways Downstream of Epidermal Growth Factor Receptor in Glioma.
Targeting the YB-1/PD-L1 Axis to Enhance Chemotherapy and Antitumor Immunity.
Phosphofructokinase 1 platelet isoform induces PD-L1 expression to promote glioblastoma immune evasion.
Cyclin D-CDK4 kinase destabilizes PD-L1 via cullin 3-SPOP to control cancer immune surveillance.
Human papillomavirus 16 E6/E7 contributes to immune escape and progression of cervical cancer by regulating miR-142-5p/PD-L1 axis.
Pentamethylquercetin Inhibits Hepatocellular Carcinoma Progression and Adipocytes-induced PD-L1 Expression via IFN- $\gamma$ Signaling.
Regulation of PD-L1 expression by matrix stiffness in lung cancer cells.
Selective Suppression of Cell Growth and Programmed Cell Death-Ligand 1 Expression in HT1080 Fibrosarcoma Cells by Low Molecular Weight Fucoidan Extract.
Increase of cells expressing PD-L1 in bovine leukemia virus infection and enhancement of anti-viral immune responses in vitro via PD-L1 blockade.
Expression of PD-L1 on canine tumor cells and enhancement of IFN- $\gamma$ production from tumor-infiltrating cells by PD-L1 blockade.
Chrysin inhibits hepatocellular carcinoma progression through suppressing programmed death ligand 1 expression.
Glycosylation and stabilization of programmed death ligand-1 suppresses T-cell activity.
Proinflammatory S100A8 Induces PD-L1 Expression in Macrophages, Mediating Tumor Immune Escape.
IFN- $\gamma$ Promotes Epithelial-Mesenchymal Transition and the Expression of PD-L1 in Pancreatic Cancer.
PD-L1 checkpoint inhibition and anti-CTLA-4 whole tumor cell vaccination counter adaptive immune resistance: A mouse neuroblastoma model that mimics human disease.
Extracellular acidity in tumor tissue upregulates programmed cell death protein 1 expression on tumor cells via proton-sensing G protein-coupled receptors.
PD-L1 Expression and Immune Escape in Melanoma Resistance to MAPK Inhibitors.
PD-L1 Expression Is Associated with FOXP3+ Regulatory T-Cell Infiltration of Soft Tissue Sarcoma and Poor Patient Prognosis.
Histone deacetylase 2 knockout suppresses immune escape of triple-negative breast cancer cells via downregulating PD-L1 expression.
M1 Macrophages Induce PD-L1 Expression in Hepatocellular Carcinoma Cells Through IL-1 $\beta$ Signaling.
Programmed death-ligand 1 and its soluble form are highly expressed in nasal natural killer/T-cell lymphoma: a potential rationale for immunotherapy.
Orphan nuclear receptor TLX promotes immunosuppression via its transcriptional activation of PD-L1 in glioma.
A mechanism of hypoxia-mediated escape from adaptive immunity in cancer cells.
Regulation of PD-L1: Emerging Routes for Targeting Tumor Immune Evasion.
miRNA-148a-3p Regulates Immunosuppression in DNA Mismatch Repair-Deficient Colorectal Cancer by Targeting PD-L1.

An inhibitor of programmed death ligand 1 enhances natural killer cell-mediated immunity against malignant melanoma cells.
PD-1 Independent Role of PD-L1 in Triple-Negative Breast Cancer Progression.
PD-1 and PD-L1 expression on TILs in peritoneal metastases compared to ovarian tumor tissues and its associations with clinical outcome.
PD-L1 expression is regulated by both DNA methylation and NF- $\kappa$ B during EMT signaling in non-small cell lung carcinoma.
PD-L1 targeting high-affinity NK (t-haNK) cells induce direct antitumor effects and target suppressive MDSC populations.
High PD-L1 expression on immune cells, but not on tumor cells, is a favorable prognostic factor in urothelial carcinoma.
Role of PD-L1 expression as a biomarker for GEP neuroendocrine neoplasm grading.
Macrophage migration inhibitory factor-CD74 interaction regulates the expression of programmed cell death ligand 1 in melanoma cells.
Increased PD-L1 expression in erlotinib-resistant NSCLC cells with MET gene amplification is reversed upon MET-TKI treatment.
Regulation of programmed death ligand 1 (PD-L1) expression by TNF-related apoptosis-inducing ligand (TRAIL) in triple-negative breast cancer cells.
Regulation of programmed death ligand 1 expression by interferon- $\gamma$ and tumour necrosis factor- $\alpha$ in canine tumour cell lines.
Lentivirus-mediated RNA interference targeting programmed death receptor ligand 1 increases the immunologic anti-tumor effect of dendritic cell vaccination against pancreatic cancer in SCID-hu mice.
Increased PD-L1 Expression in Human Skin Acutely and Chronically Exposed to UV Irradiation.
Immune evasion mediated by PD-L1 on glioblastoma-derived extracellular vesicles.
Blocking the PD-1/PD-L1 pathway in glioma: a potential new treatment strategy.
Atorvastatin Enhances the Efficacy of Immune Checkpoint Therapy and Suppresses the Cellular and Extracellular Vesicle PD-L1.
Expression of immune checkpoints (PD-L1 and IDO) and tumour-infiltrating lymphocytes in breast cancer.
Control of PD-L1 expression by miR-140/142/340/383 and oncogenic activation of the OCT4-miR-18a pathway in cervical cancer.
MET Receptor Tyrosine Kinase Regulates the Expression of Co-Stimulatory and Co-Inhibitory Molecules in Tumor Cells and Contributes to PD-L1-Mediated Suppression of Immune Cell Function.
Salmonella enterica Serovar Typhimurium Increases Functional PD-L1 Synergistically with Gamma Interferon in Intestinal Epithelial Cells via Salmonella Pathogenicity Island 2.
Clinical Significance of Program Death Ligand-1 and Indoleamine-2,3-Dioxygenase Expression in Colorectal Carcinoma.
miR-4759 suppresses breast cancer through immune checkpoint blockade.
Programmed cell death ligand 1 (PD-L1) expression is not a predominant feature in Ewing sarcomas.
Glycolytic enzyme HK2 promotes PD-L1 expression and breast cancer cell immune evasion.
Classifying Cancers Based on T-cell Infiltration and PD-L1.
Colorectal Cancer-Derived Small Extracellular Vesicles Promote Tumor Immune Evasion by Upregulating PD-L1 Expression in Tumor-Associated Macrophages.
A Tumor-Specific Super-Enhancer Drives Immune Evasion by Guiding Synchronous Expression of PD-L1 and PD-L2.
Extracellular vesicles containing PD-L1 contribute to CD8+ T-cell immune suppression and predict poor outcomes in small cell lung cancer.

POU2F1 induces the immune escape in lung cancer by up-regulating PD-L1.
Variability in Immunohistochemical Detection of Programmed Death Ligand 1 (PD-L1) in Cancer Tissue Types.
Immune gene expression profiles in high-grade urothelial carcinoma of the bladder: a NanoString study.
New Dancing Couple: PD-L1 and MicroRNA.
YAP regulates PD-L1 expression in human NSCLC cells.
PD-L1 expression is associated with epithelial-to-mesenchymal transition in adenocarcinoma of the lung.
SPOP mutations promote tumor immune escape in endometrial cancer via the IRF1-PD-L1 axis.
Silencing of RAB42 down-regulated PD-L1 expression to inhibit the immune escape of hepatocellular carcinoma cells through inhibiting the E2F signaling pathway.
O-GlcNAcylation promotes tumor immune evasion by inhibiting PD-L1 lysosomal degradation.
A novel model of controlling PD-L1 expression in ALK(+) anaplastic large cell lymphoma revealed by CRISPR screening.
TIMELESS upregulates PD-L1 expression and exerts an immunosuppressive role in breast cancer.
Development and characterization of single domain monoclonal antibody against programmed cell death ligand-1; as a cancer inhibitor candidate.
lncRNA EGFR-AS1 facilitates leiomyosarcoma progression and immune escape via the EGFR-MYC-PD-L1 axis.
The prevalence and clinicopathological features of programmed death-ligand 1 (PD-L1) expression: a pooled analysis of literatures.
PD-L1 and PD-L2 expression correlated genes in non-small-cell lung cancer.
Baicalin and baicalin promote antitumor immunity by suppressing PD-L1 expression in hepatocellular carcinoma cells.
Hepatic myofibroblasts exert immunosuppressive effects independent of the immune checkpoint regulator PD-L1 in liver metastasis of pancreatic ductal adenocarcinoma.
Epstein-Barr virus LMP1 enhances levels of large extracellular vesicle-associated PD-L1.
Gastric cancer derived mesenchymal stem cells promoted DNA repair and cisplatin resistance through up-regulating PD-L1/Rad51 in gastric cancer.
High expression of AMAP1, an ARF6 effector, is associated with elevated levels of PD-L1 and fibrosis of pancreatic cancer.
Dexamethasone suppresses immune evasion by inducing GR/STAT3 mediated downregulation of PD-L1 and IDO1 pathways.
Prognostic value and clinicopathological roles of the tumor immune microenvironment in salivary duct carcinoma.
Targeting Lin28 axis enhances glypican-3-CAR T cell efficacy against hepatic tumor initiating cell population.
Combined vaccine-immune-checkpoint inhibition constitutes a promising strategy for treatment of dMMR tumors.
PD-L1 expression is associated with ALK positivity and STAT3 activation, but not outcome in patients with systemic anaplastic large cell lymphoma.
Lynch syndrome-associated epithelial ovarian cancer and its immunological profile.
Targeting CXCL12 from FAP-expressing carcinoma-associated fibroblasts synergizes with anti-PD-L1 immunotherapy in pancreatic cancer.
MiR-10b-5p Impairs TET2-Mediated Inhibition of PD-L1 Transcription Thus Promoting Immune Evasion and Tumor Progression in Glioblastoma.

Mechanism of immune escape mediated by receptor tyrosine kinase KIT in thyroid cancer.
Engineering of a novel subnanomolar affinity fibronectin III domain binder targeting human programmed death-ligand 1.
Anlotinib suppresses proliferation, migration, and immune escape of gastric cancer cells by activating the cGAS-STING/IFN- $\beta$ pathway.
Distinct tumor immune microenvironments in primary and metastatic lesions in gastric cancer patients.
INPP4B inhibits glioma cell proliferation and immune escape via inhibition of the PI3K/AKT signaling pathway.
CMTM6 shapes antitumor T cell response through modulating protein expression of CD58 and PD-L1.
Zinc Status Impacts the Epidermal Growth Factor Receptor and Downstream Protein Expression in A549 Cells.
$\beta$ -Catenin regulates tumor-derived PD-L1.
GM-CSF mediates immune evasion via upregulation of PD-L1 expression in extranodal natural killer/T cell lymphoma.
FGFBP1 as a potential biomarker predicting bacillus Calmette-Guérin response in bladder cancer.
Involvement of PD-L1-mediated tumor-intrinsic signaling and immune suppression in tumorigenic effect of $\alpha$ -tocopherol.
Docosahexaenoic acid may inhibit immune evasion of colorectal cancer cells through targeting immune checkpoint and immunomodulator genes and their controlling microRNAs.
Intravascular Large B-Cell Lymphoma Genomic Profile Is Characterized by Alterations in Genes Regulating NF- $\kappa$ B and Immune Checkpoints.
The Effects of PI3K/Akt/mTOR Signaling Pathway Inhibitors on the Expression of Immune Checkpoint Ligands in Acute Myeloid Leukemia Cell Line.
Phosphatidylserine Sensing by TAM Receptors Regulates AKT-Dependent Chemoresistance and PD-L1 Expression.
Aggressive tumor microenvironment of solid predominant lung adenocarcinoma subtype harboring with epidermal growth factor receptor mutations.
Beyond the Driver Mutation: Immunotherapies in Gastrointestinal Stromal Tumors.
Age-related EBV-associated B-cell lymphoproliferative disorders and other EBV + lymphoproliferative diseases: New insights into immune escape and immunodeficiency through staining with anti-PD-L1 antibody clone SP142.
LncRNA SNHG1 regulates immune escape of renal cell carcinoma by targeting miR-129-3p to activate STAT3 and PD-L1.
Tumor-Derived PD-L1(+) Exosomes with Natural Inflammation Tropism for Psoriasis-Targeted Treatment.
The folate cycle enzyme MTHFD2 induces cancer immune evasion through PD-L1 up-regulation.
Prostate cancer C5a receptor expression and augmentation of cancer cell proliferation, invasion, and PD-L1 expression by C5a.
ATXN2-Mediated PI3K/AKT Activation Confers Gastric Cancer Chemoresistance and Attenuates CD8(+) T Cell Cytotoxicity.
ALKBH5 promotes PD-L1-mediated immune escape through m6A modification of ZDHHC3 in glioma.
Synthesis and Evaluation of (177)Lu-DOTA-PD-L1-i and (225)Ac-HEHA-PD-L1-i as Potential Radiopharmaceuticals for Tumor Microenvironment-Targeted Radiotherapy.
Unexpected PD-L1 immune evasion mechanism in TNBC, ovarian, and other solid tumors by DR5 agonist antibodies.

Intestine-Specific Homeobox Gene ISX Integrates IL6 Signaling, Tryptophan Catabolism, and Immune Suppression.
Increased Nuclear Transporter KPNA2 Contributes to Tumor Immune Evasion by Enhancing PD-L1 Expression in PDAC.
Circular RNA hsa_circ_0001598 promotes programmed death-ligand-1-mediated immune escape and trastuzumab resistance via sponging miR-1184 in breast cancer cells.
Loss of Lkb1 and Pten leads to lung squamous cell carcinoma with elevated PD-L1 expression.
The diverse pancreatic tumor cell-intrinsic response to IFN $\gamma$ is determined by epigenetic heterogeneity.
Lymphocyte-depleted classic Hodgkin lymphoma with primary extranodal disease: Two cases that highlight the combination of immunodeficiency and immune escape in the pathogenesis.
The loss of epithelial Smad4 drives immune evasion via CXCL1 while displaying vulnerability to combinatorial immunotherapy in gastric cancer.
Elimination of tumor by CD47/PD-L1 dual-targeting fusion protein that engages innate and adaptive immune responses.
Translation control of the immune checkpoint in cancer and its therapeutic targeting.
IL6 Induces mtDNA Leakage to Affect the Immune Escape of Endometrial Carcinoma via cGAS-STING.
Expression of ALCAM (CD166) and PD-L1 (CD274) independently predicts shorter survival in malignant pleural mesothelioma.
CircRNA VIM silence synergizes with sevoflurane to inhibit immune escape and multiple oncogenic activities of esophageal cancer by simultaneously regulating miR-124/PD-L1 axis.
Targeting WD repeat domain 5 enhances chemosensitivity and inhibits proliferation and programmed death-ligand 1 expression in bladder cancer.
CD73 Promotes Chronic Lymphocytic Leukemia.
Mitochondrial Lon-induced mtDNA leakage contributes to PD-L1-mediated immunoescape via STING-IFN signaling and extracellular vesicles.
Helicobacter pylori Modulated Host Immunity in Gastric Cancer Patients With S-1 Adjuvant Chemotherapy.
Dual FGFR and VEGFR inhibition synergistically restrain hexokinase 2-dependent lymphangiogenesis and immune escape in intrahepatic cholangiocarcinoma.
Chinese herbal compound SanHuang decoction reverses axitinib resistance in ccRCC through regulating immune cell infiltration by affecting ADAMTS18 expression.
Indoleamine 2,3-dioxygenase in endometrial cancer: a targetable mechanism of immune resistance in mismatch repair-deficient and intact endometrial carcinomas.
A new antisarcoma strategy: multisubtype heat shock protein/peptide immunotherapy combined with PD-L1 immunological checkpoint inhibitors.
Correction to: Long non-coding RNA HOTTIP enhances IL-6 expression to potentiate immune escape of ovarian cancer cells by upregulating the expression of PD-L1 in neutrophils.
Upregulation of PD-L1 by SPP1 mediates macrophage polarization and facilitates immune escape in lung adenocarcinoma.
Clinical Implications of Immune Checkpoints and the RANK/RANK-L Signaling Pathway in High-Grade Canine Mast Cell Tumors.
Microbial hydrogen "manufactory" for enhanced gas therapy and self-activated immunotherapy via reduced immune escape.
Prognostic relevance of tumor-infiltrating lymphocytes and immune checkpoints in pediatric medulloblastoma.

Clinicopathologic profile, immunophenotype, and genotype of CD274 (PD-L1)-positive colorectal carcinomas.
Treatment with decitabine induces the expression of stemness markers, PD-L1 and NY-ESO-1 in colorectal cancer: potential for combined chemoimmunotherapy.
Clinical, Genetic and Innate Immunity Characteristics of Melanoma in Organ Transplant Recipients.
Prodrug-Based Versatile Nanomedicine with Simultaneous Physical and Physiological Tumor Penetration for Enhanced Cancer Chemo-Immunotherapy.
Fibroblast-derived LPP as a biomarker for treatment response and therapeutic target in gastric cancer.
Identification of CircRNA signature associated with tumor immune infiltration to predict therapeutic efficacy of immunotherapy.
Expression of immune checkpoints on circulating tumor cells in men with metastatic prostate cancer.
SUMOylated IL-33 in the nucleus stabilizes the transcription factor IRF1 in hepatocellular carcinoma cells to promote immune escape.
Immune-evasive human islet-like organoids ameliorate diabetes.
Association of increased B7 protein expression by infiltrating immune cells with progression of gastric carcinogenesis.
Immune checkpoint molecules soluble program death ligand 1 and galectin-9 are increased in pregnancy.
Lactate induces PD-L1 in HRAS(G12V)-positive oropharyngeal squamous cell carcinoma.
Therapeutic Co-targeting of WEE1 and ATM Downregulates PD-L1 Expression in Pancreatic Cancer.
CD47 is associated with the up-regulation of the PD-1 oncogenic signaling pathway.
Upregulation of ARNTL2 is associated with poor survival and immune infiltration in clear cell renal cell carcinoma.
Deubiquitinase ubiquitin-specific peptidase 10 maintains cysteine rich angiogenic inducer 61 expression via Yes1 associated transcriptional regulator to augment immune escape and metastasis of pancreatic adenocarcinoma.
WD repeat domain 5 promotes chemoresistance and Programmed Death-Ligand 1 expression in prostate cancer.
HOTAIR Up-Regulation Activates NF- $\kappa$ B to Induce Immunoescape in Gliomas.
Aryl Hydrocarbon Receptor Signaling Controls CD155 Expression on Macrophages and Mediates Tumor Immunosuppression.
Avelumab and other recent advances in Merkel cell carcinoma.
A Case Report of Metachronous Multiple Adenosquamous Carcinoma of the Colon Over-expressing PD-L1 and a Literature Review.
PD-L1, LAG3, and HLA-DR are increasingly expressed during smoldering myeloma progression.
Potentiated effects of lactate receptor GPR81 on immune microenvironment in breast cancer.
Treatment by PI3K/mTOR Inhibitor BEZ235 Combined with TLR-7/8 Agonist Interfere with Immune Evasion Mechanisms of WEHI-3 Mouse Leukemia Cells.
Role of PD-1/PD-L1-mediated tumour immune escape mechanism and microsatellite instability in the BCG failure of high-grade urothelial carcinomas.
c-Myb facilitates immune escape of esophageal adenocarcinoma cells through the miR-145-5p/SPOP/PD-L1 axis.
A self-amplifying nanodrug to manipulate the Janus-faced nature of ferroptosis for tumor therapy.
B7x-from bench to bedside.
[Immunotherapy Targeting Tumor-Associated Carbohydrate Antigens in Deficient Mismatch Repair Colorectal Cancer].
Deciphering Mechanisms of UVR-Induced Tumoral Immune Checkpoint Regulation against Melanoma.

LINC01119 encapsulated by cancer-associated adipocytes-derived exosomes promotes M2 polarization of macrophages to induce immune escape in ovarian cancer in a 3D co-culture cell-based model.
RNF182 induces p65 ubiquitination to affect PDL1 transcription and suppress immune evasion in lung adenocarcinoma.
Immunobiology of Uveal Melanoma: State of the Art and Therapeutic Targets.
Loss of PTEN causes SHP2 activation, making lung cancer cells unresponsive to IFN- $\gamma$ .
CMTM6 attenuates cisplatin-induced cell death in OSCC by regulating AKT/c-Myc-driven ribosome biogenesis.
Long Noncoding RNA Hotair Promotes the Progression and Immune Escape in Laryngeal Squamous Cell Carcinoma through MicroRNA-30a/GRP78/PD-L1 Axis.
Increased cFLIP expression in thymic epithelial tumors blocks autophagy via NF- $\kappa$ B signalling.
Preclinical Evaluation of the Multiple Tyrosine Kinases Inhibitor Anlotinib in Leukemia Stem Cells.
Anaplastic lymphoma kinase (ALK)-induced malignancies: novel mechanisms of cell transformation and potential therapeutic approaches.

Table A2-48, Cluster 47

Cluster 47 focuses on fungal pathogens, especially the interactions of <i>Candida albicans</i> with innate host defense and its escape strategies (359)
Inhibition of Respiration of <i>Candida albicans</i> by Small Molecules Increases Phagocytosis Efficacy by Macrophages.
I want to break free - macrophage strategies to recognize and kill <i>Candida albicans</i> , and fungal counter-strategies to escape.
Interactions of fungal pathogens with phagocytes.
High-Throughput Screening Identifies Genes Required for <i>Candida albicans</i> Induction of Macrophage Pyroptosis.
Revisiting the Vital Drivers and Mechanisms of $\beta$ -Glucan Masking in Human Fungal Pathogen, <i>Candida albicans</i> .
Fungal mechanisms of intracellular survival: what can we learn from bacterial pathogens?
Global analysis of fungal morphology exposes mechanisms of host cell escape.
Impact of the Environment upon the <i>Candida albicans</i> Cell Wall and Resultant Effects upon Immune Surveillance.
Interaction of <i>Candida albicans</i> with host cells: virulence factors, host defense, escape strategies, and the microbiota.
Interplay between <i>Candida albicans</i> and the mammalian innate host defense.
<i>Candida albicans</i> hypha formation and mannan masking of $\beta$ -glucan inhibit macrophage phagosome maturation.
Transcriptional Control of Drug Resistance, Virulence and Immune System Evasion in Pathogenic Fungi: A Cross-Species Comparison.
The advances in the regulation of immune microenvironment by <i>Candida albicans</i> and macrophage cross-talk.
Is There a Relationship Between Mating and Pathogenesis in Two Human Fungal Pathogens, <i>Candida albicans</i> and <i>Candida glabrata</i> ?
Complement and innate immune evasion strategies of the human pathogenic fungus <i>Candida albicans</i> .



Two unlike cousins: <i>Candida albicans</i> and <i>C. glabrata</i> infection strategies.
Epitope Shaving Promotes Fungal Immune Evasion.
<i>Candida</i> survival strategies.
Immune escape of the human facultative pathogenic yeast <i>Candida albicans</i> : the many faces of the <i>Candida</i> Pra1 protein.
Neutrophil Attack Triggers Extracellular Trap-Dependent <i>Candida</i> Cell Wall Remodeling and Altered Immune Recognition.
Insect fungal pathogens secrete a cell wall-associated glucanase that acts to help avoid recognition by the host immune system.
Immune Recognition of Fungal Polysaccharides.
A virtual infection model quantifies innate effector mechanisms and <i>Candida albicans</i> immune escape in human blood.
Beyond the wall: <i>Candida albicans</i> secret(e)s to survive.
A novel immune evasion strategy of <i>Candida albicans</i> : proteolytic cleavage of a salivary antimicrobial peptide.
[Progress on the role of Toll-like receptors in <i>Candida albicans</i> infections].
<i>Candida albicans</i> escapes from mouse neutrophils.
Host-pathogen interactions and the pathological consequences of acute systemic <i>Candida albicans</i> infections in mice.
Mnn10 Maintains Pathogenicity in <i>Candida albicans</i> by Extending $\alpha$ -1,6-Mannose Backbone to Evade Host Dectin-1 Mediated Antifungal Immunity.
A <i>Candida albicans</i> regulator of disseminated infection operates primarily as a repressor and governs cell surface remodeling.
In vitro and in vivo Characterization of Host-Pathogen Interactions of the L3881 <i>Candida albicans</i> Clinical Isolate.
The pathogen <i>Candida albicans</i> hijacks pyroptosis for escape from macrophages.
<i>Candida albicans</i> Quorum Sensing Molecules Stimulate Mouse Macrophage Migration.
Friendly fungi: symbiosis with commensal <i>Candida albicans</i> .
The macrophage-derived protein PTMA induces filamentation of the human fungal pathogen <i>Candida albicans</i> .
Predictive Virtual Infection Modeling of Fungal Immune Evasion in Human Whole Blood.
Dectin-1 escape by fungal dimorphism.
Role of the innate immune system in host defence against fungal infections.
Infection-associated genes of <i>Candida albicans</i> .
Analysis of the <i>Candida albicans</i> - specific T-cell response and oropharyngeal <i>Candida</i> colonization in a cohort of HIV-1-infected patients.
Divergent Approaches to Virulence in <i>C. albicans</i> and <i>C. glabrata</i> : Two Sides of the Same Coin.
<i>Candida</i> spp. and phagocytosis: multiple evasion mechanisms.
Characterization of a <i>Candida albicans</i> isolate from a recurrent cervical lymphadenitis patient.
The glycosylphosphatidylinositol-anchored protease Sap9 modulates the interaction of <i>Candida albicans</i> with human neutrophils.
Analysis of Volatile Molecules Present in the Secretome of the Fungal Pathogen <i>Candida glabrata</i> .
Integrated inference and evaluation of host-fungi interaction networks.
Lactate signalling regulates fungal $\beta$ -glucan masking and immune evasion.
<i>Candida albicans</i> cell surface superoxide dismutases degrade host-derived reactive oxygen species to escape innate immune surveillance.

Live <i>Candida albicans</i> suppresses production of reactive oxygen species in phagocytes.
Estrogen promotes innate immune evasion of <i>Candida albicans</i> through inactivation of the alternative complement system.
Post-transcriptional control of fungal cell wall synthesis.
Mechanisms of immune evasion in fungal pathogens.
The Dual Function of the Fungal Toxin Candidalysin during <i>Candida albicans</i> -Macrophage Interaction and Virulence.
Fungal morphogenetic changes inside the mammalian host.
<i>Candida albicans</i> Chitin Increases Arginase-1 Activity in Human Macrophages, with an Impact on Macrophage Antimicrobial Functions.
Morphology Changes in Human Fungal Pathogens upon Interaction with the Host.
<i>Candida albicans</i> PPG1, a serine/threonine phosphatase, plays a vital role in central carbon metabolisms under filament-inducing conditions: A multi-omics approach.
<i>Candida albicans</i> Shields the Periodontal Killer <i>Porphyromonas gingivalis</i> from Recognition by the Host Immune System and Supports the Bacterial Infection of Gingival Tissue.
Fungal Strategies to Evade the Host Immune Recognition.
The Interactions Between <i>Candida albicans</i> and Mucosal Immunity.
Fungal strategies for overcoming host innate immune response.
Dynamic Fungal Cell Wall Architecture in Stress Adaptation and Immune Evasion.
Role of phospholipases in fungal fitness, pathogenicity, and drug development - lessons from <i>Cryptococcus neoformans</i> .
Polysaccharides Cell Wall Architecture of Mucorales.
Evasion of Immune Surveillance in Low Oxygen Environments Enhances <i>Candida albicans</i> Virulence.
Fungal infections: Pathogenesis, antifungals and alternate treatment approaches.
<i>Candida albicans</i> dimorphism as a therapeutic target.
Rab14 regulates maturation of macrophage phagosomes containing the fungal pathogen <i>Candida albicans</i> and outcome of the host-pathogen interaction.
Flavodoxin-Like Proteins Protect <i>Candida albicans</i> from Oxidative Stress and Promote Virulence.
Non- <i>albicans</i> <i>Candida</i> Species: Immune Response, Evasion Mechanisms, and New Plant-Derived Alternative Therapies.
Genetic Susceptibility to Fungal Infections and Links to Human Ancestry.
<i>Candida auris</i> Cell Wall Mannosylation Contributes to Neutrophil Evasion through Pathways Divergent from <i>Candida albicans</i> and <i>Candida glabrata</i> .
The Endoplasmic Reticulum-Mitochondrion Tether ERMES Orchestrates Fungal Immune Evasion, Illuminating Inflammasome Responses to Hyphal Signals.
Dectin-1 is required for miR155 upregulation in murine macrophages in response to <i>Candida albicans</i> .
The secreted <i>Candida albicans</i> protein Pra1 disrupts host defense by broadly targeting and blocking complement C3 and C3 activation fragments.
ATAC-Seq Identifies Chromatin Landscapes Linked to the Regulation of Oxidative Stress in the Human Fungal Pathogen <i>Candida albicans</i> .
Thriving within the host: <i>Candida</i> spp. interactions with phagocytic cells.
Hypoxia Promotes Immune Evasion by Triggering $\beta$ -Glucan Masking on the <i>Candida albicans</i> Cell Surface via Mitochondrial and cAMP-Protein Kinase A Signaling.
Host-fungal interactions: key players of antifungal immunity.
<i>Aspergillus fumigatus</i> cytochrome c impacts conidial survival during sterilizing immunity.

Rim Pathway-Mediated Alterations in the Fungal Cell Wall Influence Immune Recognition and Inflammation.
Dynamic, morphotype-specific <i>Candida albicans</i> beta-glucan exposure during infection and drug treatment.
Soluble Enolase 1 of <i>Candida albicans</i> and <i>Aspergillus fumigatus</i> Stimulates Human and Mouse B Cells and Monocytes.
Mixed Fungal Biofilms: From Mycobiota to Devices, a New Challenge on Clinical Practice.
Mechanism of <i>Candida</i> pathogenesis: revisiting the vital drivers.
Cell biology of <i>Candida albicans</i> -host interactions.
Non-canonical signalling mediates changes in fungal cell wall PAMPs that drive immune evasion.
Nature of $\beta$ -1,3-Glucan-Exposing Features on <i>Candida albicans</i> Cell Wall and Their Modulation.
Fungal biotin homeostasis is essential for immune evasion after macrophage phagocytosis and virulence.
Fungal morphogenesis and host invasion.
Insect Pathogenic Fungi: Genomics, Molecular Interactions, and Genetic Improvements.
Fungal factors involved in host immune evasion, modulation and exploitation during infection.
Pattern recognition receptors in antifungal immunity.
Adapting to survive: How <i>Candida</i> overcomes host-imposed constraints during human colonization.
The fungal peptide toxin Candidalysin activates the NLRP3 inflammasome and causes cytolysis in mononuclear phagocytes.
Development of vaccines for <i>Candida albicans</i> : fighting a skilled transformer.
The ER-mitochondria encounter structure contributes to hyphal growth, mitochondrial morphology and virulence of the pathogenic mold <i>Aspergillus fumigatus</i> .
Extracellular proteinases of <i>Candida</i> species pathogenic yeasts.
Fungal Mimicry of a Mammalian Aminopeptidase Disables Innate Immunity and Promotes Pathogenicity.
<i>Candida</i> species distribution, genotyping and virulence factors of <i>Candida albicans</i> isolated from the oral cavity of kidney transplant recipients of two geographic regions of Brazil.
Systems biology of host-fungus interactions: turning complexity into simplicity.
Endosomal sensing of fungi: current understanding and emerging concepts.
Emerging Fungal Pathogen <i>Candida auris</i> Evades Neutrophil Attack.
Defects in intracellular trafficking of fungal cell wall synthases lead to aberrant host immune recognition.
Histologic findings of <i>Massospora cicadina</i> infection in periodical cicadas ( <i>Magicicada septendecim</i> ).
What Are the Functions of Chitin Deacetylases in <i>Aspergillus fumigatus</i> ?
Glucan Unmasking Identifies Regulators of Temperature-Induced Translatome Reprogramming in <i>C. neoformans</i> .
How Environmental Fungi Cause a Range of Clinical Outcomes in Susceptible Hosts.
Host-Pathogen Interactions Mediated by MDR Transporters in Fungi: As Pleiotropic as it Gets!
The lipid language of plant-fungal interactions.
Contribution of Aspartic Proteases in <i>Candida</i> Virulence. Protease Inhibitors against <i>Candida</i> Infections.
Fungi in the cystic fibrosis lung: bystanders or pathogens?
Genomic and functional analyses unveil the response to hyphal wall stress in <i>Candida albicans</i> cells lacking $\beta$ (1,3)-glucan remodeling.
Opportunistic yeast pathogen <i>Candida</i> spp.: Secreted and membrane-bound virulence factors.

Conserved and Divergent Features of pH Sensing in Major Fungal Pathogens.
Magnesium impairs <i>Candida albicans</i> immune evasion by reduced hyphal damage, enhanced $\beta$ -glucan exposure and altered vacuole homeostasis.
Cek1 regulates $\beta(1,3)$ -glucan exposure through calcineurin effectors in <i>Candida albicans</i> .
Transcriptomic analyses reveal comprehensive responses of insect hemocytes to mycopathogen <i>Beauveria bassiana</i> , and fungal virulence-related cell wall protein assists pathogen to evade host cellular defense.
Phenotypic profiles of virulence in different <i>Candida</i> species isolated from vulvovaginal infections.
The potential impact of antifungal drug resistance mechanisms on the host immune response to <i>Candida</i> .
Innate antifungal immunity: the key role of phagocytes.
<i>Mucor circinelloides</i> Thrives inside the Phagosome through an Atf-Mediated Germination Pathway.
Multifunctional role of a fungal pathogen-secreted laccase 2 in evasion of insect immune defense.
Basic principles of the virulence of <i>Cryptococcus</i> .
Secreted aspartyl proteases family: a perspective review on the regulation of fungal pathogenesis.
"Under Pressure" - How fungi evade, exploit, and modulate cells of the innate immune system.
<i>Cryptococcus</i> and Phagocytes: Complex Interactions that Influence Disease Outcome.
Innate Pulmonary Phagocytes and Their Interactions with Pathogenic <i>Cryptococcus</i> Species.
Fungal-induced cell cycle impairment, chromosome instability and apoptosis via differential activation of NF- $\kappa$ B.
Exposure of Keratinocytes to <i>Candida Albicans</i> in the Context of Atopic Milieu Induces Changes in the Surface Glycosylation Pattern of Small Extracellular Vesicles to Enhance Their Propensity to Interact With Inhibitory Siglec Receptors.
Identification of Botanicals that Unmask $\beta$ -Glucan from the Cell Surface of an Opportunistic Fungal Pathogen.
Innate immune recognition of microbial cell wall components and microbial strategies to evade such recognitions.
Genes and molecules involved in <i>Aspergillus fumigatus</i> virulence.
Aspergillosis and stem cell transplantation: An overview of experimental pathogenesis studies.
Investigating Metabolic and Molecular Ecological Evolution of Opportunistic Pulmonary Fungal Coinfections: Protocol for a Laboratory-Based Cross-Sectional Study.
Innate immune evasion strategies against Cryptococcal meningitis caused by <i>Cryptococcus neoformans</i> .
<i>Candida glabrata</i> Antifungal Resistance and Virulence Factors, a Perfect Pathogenic Combination.
The histone chaperone HIR maintains chromatin states to control nitrogen assimilation and fungal virulence.
Functional Characterization of Secreted Aspartyl Proteases in <i>Candida parapsilosis</i> .
Caspofungin-induced $\beta(1,3)$ -glucan exposure in <i>Candida albicans</i> is driven by increased chitin levels.
Enolase 1 of <i>Candida albicans</i> binds human CD4(+) T cells and modulates naïve and memory responses.
Enemy of the (immunosuppressed) state: an update on the pathogenesis of <i>Aspergillus fumigatus</i> infection.
<i>Cryptococcus</i> interactions with macrophages: evasion and manipulation of the phagosome by a fungal pathogen.
A Strategy of On-Demand Immune Activation for Antifungal Treatment Using Near-Infrared Responsive Conjugated Polymer Nanoparticles.

The Paralogous Transcription Factors Stp1 and Stp2 of <i>Candida albicans</i> Have Distinct Functions in Nutrient Acquisition and Host Interaction.
Fungal co-expression network analyses identify pathogen gene modules associated with host insect invasion.
Enzymatic Mechanisms Involved in Evasion of Fungi to the Oxidative Stress: Focus on <i>Scedosporium apiospermum</i> .
Thoughts on the evolution of Core Environmental Responses in yeasts.
Peptidylarginine Deiminase of <i>Porphyromonas gingivalis</i> Modulates the Interactions between <i>Candida albicans</i> Biofilm and Human Plasminogen and High-Molecular-Mass Kininogen.
Proteome Analysis Reveals the Conidial Surface Protein CcpA Essential for Virulence of the Pathogenic Fungus <i>Aspergillus fumigatus</i> .
Proteinase and phospholipase activity as virulence factors in <i>Candida</i> species isolated from blood.
The F-Box Protein Fbp1 Shapes the Immunogenic Potential of <i>Cryptococcus neoformans</i> .
A novel role for a glycosylphosphatidylinositol-anchored aspartyl protease, CgYps1, in the regulation of pH homeostasis in <i>Candida glabrata</i> .
Thermotolerance in the pathogen <i>Cryptococcus neoformans</i> is linked to antigen masking via mRNA decay-dependent reprogramming.
Immune evasion, stress resistance, and efficient nutrient acquisition are crucial for intracellular survival of <i>Candida glabrata</i> within macrophages.
An evolutionarily diverged mitochondrial protein controls biofilm growth and virulence in <i>Candida albicans</i> .
Bst1 is required for <i>Candida albicans</i> infecting host via facilitating cell wall anchorage of Glycosylphosphatidyl inositol anchored proteins.
Depletion of Extracellular Chemokines by <i>Aspergillus</i> Melanin.
Identifying <i>Candida albicans</i> Gene Networks Involved in Pathogenicity.
Exploring the intersection of <i>Aspergillus fumigatus</i> biofilms, infections, immune response and antifungal resistance.
Analysis of the cellular <i>Aspergillus fumigatus</i> proteome that reacts with sera from rabbits developing an acquired immunity after experimental aspergillosis.
<i>Cryptococcus neoformans</i> Csn1201 Is Associated With Pulmonary Immune Responses and Disseminated Infection.
Direct Binding of the pH-Regulated Protein 1 (Pra1) from <i>Candida albicans</i> Inhibits Cytokine Secretion by Mouse CD4(+) T Cells.
The <i>Cryptococcus neoformans</i> Rim101 transcription factor directly regulates genes required for adaptation to the host.
<i>Cryptococcus neoformans</i> Rim101 is associated with cell wall remodeling and evasion of the host immune responses.
Mannan Molecular Substructures Control Nanoscale Glucan Exposure in <i>Candida</i> .
Memory in Fungal Pathogens Promotes Immune Evasion, Colonisation, and Infection.
<i>Candida glabrata</i> persistence in mice does not depend on host immunosuppression and is unaffected by fungal amino acid auxotrophy.
The cause and effect of <i>Cryptococcus</i> interactions with the host.
The pathogenicity of <i>Aspergillus fumigatus</i> , drug resistance, and nanoparticle delivery.
Killed <i>Candida albicans</i> yeasts and hyphae inhibit gamma interferon release by murine natural killer cells.
Paracoccidioides-host Interaction: An Overview on Recent Advances in the Paracoccidioidomycosis.

Aspergillus fumigatus versus Genus Aspergillus: Conservation, Adaptive Evolution and Specific Virulence Genes.
Fungal extracellular vesicles: modulating host-pathogen interactions by both the fungus and the host.
A lysin motif effector subverts chitin-triggered immunity to facilitate arbuscular mycorrhizal symbiosis.
The facultative intracellular pathogen Candida glabrata subverts macrophage cytokine production and phagolysosome maturation.
MrpacC regulates sporulation, insect cuticle penetration and immune evasion in Metarhizium robertsii.
Melanization of Candida auris Is Associated with Alteration of Extracellular pH.
Type I Interferons Ameliorate Zinc Intoxication of Candida glabrata by Macrophages and Promote Fungal Immune Evasion.
Fungal biosynthesis of the bibenzoquinone oosporein to evade insect immunity.
The two-component histidine kinases DrkA and SlnA are required for in vivo growth in the human pathogen Penicillium marneffeii.
Divergent LysM effectors contribute to the virulence of Beauveria bassiana by evasion of insect immune defenses.
MripacC regulates blastosphere budding and influences virulence of the pathogenic fungus Metarhizium rileyi.
RAD51-WSS1-dependent genetic pathways are essential for DNA-protein crosslink repair and pathogenesis in Candida albicans.
A phylogenetic approach to explore the Aspergillus fumigatus conidial surface-associated proteome and its role in pathogenesis.
Members of chitin synthase family in Metarhizium acridum differentially affect fungal growth, stress tolerances, cell wall integrity and virulence.
The SPS amino acid sensor mediates nutrient acquisition and immune evasion in Candida albicans.
The fungal pathogen Cryptococcus neoformans manipulates macrophage phagosome maturation.
Interference of Aspergillus fumigatus with the immune response.
The escape of Candida albicans from macrophages is enabled by the fungal toxin candidalysin and two host cell death pathways.
Multiple Disguises for the Same Party: The Concepts of Morphogenesis and Phenotypic Variations in Cryptococcus neoformans.
No evidence for immune priming in ants exposed to a fungal pathogen.
Connecting Cryptococcal Meningitis and Gut Microbiome.
Possible role of secreted proteinases in Candida albicans infections.
The transcription factor Ndt80 is a repressor of Candida parapsilosis virulence attributes.
Trojan Horse Transit Contributes to Blood-Brain Barrier Crossing of a Eukaryotic Pathogen.
[Biochemical analysis and application of molecular display technology on Candida albicans for diagnosing and preventing candidiasis].
Molecular interactions between entomopathogenic fungi (Hypocreales) and their insect host: Perspectives from stressful cuticle and hemolymph battlefields and the potential of dual RNA sequencing for future studies.
Induction of signal transduction pathways related to the pathogenicity of Cryptococcus neoformans in the host environment.
Type I Interferon Response Dysregulates Host Iron Homeostasis and Enhances Candida glabrata Infection.

Persistent <i>Cryptococcus neoformans</i> pulmonary infection in the rat is associated with intracellular parasitism, decreased inducible nitric oxide synthase expression, and altered antibody responsiveness to cryptococcal polysaccharide.
Understanding <i>Aspergillus fumigatus</i> galactosaminogalactan biosynthesis: A few questions remain.
Intracellular survival of <i>Candida glabrata</i> in macrophages: immune evasion and persistence.
Surface $\alpha$ -1,3-glucan facilitates fungal stealth infection by interfering with innate immunity in plants.
Comparative genomic and transcriptomic analysis of <i>wangiella dermatitidis</i> , a major cause of phaeohyphomycosis and a model black yeast human pathogen.
<i>Beauveria bassiana</i> ribotoxin inhibits insect immunity responses to facilitate infection via host translational blockage.
Genome-Wide Response to Drugs and Stress in the Pathogenic Yeast <i>Candida glabrata</i> .
Molecular Insights into Pathogenesis and Infection with <i>Aspergillus Fumigatus</i> .
Impact of genetic background on allele selection in a highly mutable <i>Candida albicans</i> gene, PNG2.
<i>Cryptococcus neoformans</i> Evades Pulmonary Immunity by Modulating Xylose Precursor Transport.
Chemical organization of the cell wall polysaccharide core of <i>Malassezia restricta</i> .
Intracellular infection of tick cell lines by the entomopathogenic fungus <i>Metarhizium anisopliae</i> .
MIF-like domain containing protein orchestrates cellular differentiation and virulence in the fungal pathogen <i>Magnaporthe oryzae</i> .
Could the Lung Be a Gateway for Amphotericin B to Attack the Army of Fungi?
MaPacC, a pH-responsive transcription factor, negatively regulates thermotolerance and contributes to conidiation and virulence in <i>Metarhizium acridum</i> .
Interaction of pathogenic yeasts with phagocytes: survival, persistence and escape.
Mechanisms of cryptococcal virulence and persistence.
Posttranslational, translational, and transcriptional responses to nitric oxide stress in <i>Cryptococcus neoformans</i> : implications for virulence.
Moonlighting proteins in medically relevant fungi.
Immuno-evasive <i>Aspergillus</i> virulence factors.
Anti-Immune Strategies of Pathogenic Fungi.
The Pathogenesis of <i>Aspergillus fumigatus</i> , Host Defense Mechanisms, and the Development of AFMP4 Antigen as a Vaccine.
Disruption of an adenylate-forming reductase required for conidiation, increases virulence of the insect pathogenic fungus <i>Metarhizium acridum</i> by enhancing cuticle invasion.
Plasma Gelsolin Enhances Phagocytosis of <i>Candida auris</i> by Human Neutrophils through Scavenger Receptor Class B.
Global gene deletion analysis exploring yeast filamentous growth.
$\beta$ -1,3-Glucan/CR3/SYK pathway-dependent LC3B-II accumulation enhanced the fungicidal activity in human neutrophils.
Fungal cell gigantism during mammalian infection.
Stimuli that induce production of <i>Candida albicans</i> extracellular aspartyl proteinase.
Common strategies for antigenic variation by bacterial, fungal and protozoan pathogens.
Plant defense mechanisms are activated during biotrophic and necrotrophic development of <i>Colletotricum graminicola</i> in maize.
Fungal Dimorphism and Virulence: Molecular Mechanisms for Temperature Adaptation, Immune Evasion, and In Vivo Survival.
Short-Term Probiotic Administration Increases Fecal-Anti <i>Candida</i> Activity in Healthy Subjects.

IL-27 Induced by Select <i>Candida</i> spp. via TLR7/NOD2 Signaling and IFN- $\beta$ Production Inhibits Fungal Clearance.
The novel Dbl homology/BAR domain protein, MsgA, of <i>Talaromyces marneffei</i> regulates yeast morphogenesis during growth inside host cells.
The human fungal pathogen <i>Cryptococcus neoformans</i> escapes macrophages by a phagosome emptying mechanism that is inhibited by Arp2/3 complex-mediated actin polymerisation.
Ribosomal Protein L40e Fused With a Ubiquitin Moiety Is Essential for the Vegetative Growth, Morphological Homeostasis, Cell Cycle Progression, and Pathogenicity of <i>Cryptococcus neoformans</i> .
Eosinophils subvert host resistance to an intracellular pathogen by instigating non-protective IL-4 in CCR2(-/-) mice.
Immunological Aspects of Chytridiomycosis.
Proteomic analysis of <i>Sporothrix schenckii</i> cell wall reveals proteins involved in oxidative stress response induced by menadione.
Anticipatory Stress Responses and Immune Evasion in Fungal Pathogens.
Isolation and characterization of senescent <i>Cryptococcus neoformans</i> and implications for phenotypic switching and pathogenesis in chronic cryptococcosis.
A bacterial endosymbiont of the fungus <i>Rhizopus microsporus</i> drives phagocyte evasion and opportunistic virulence.
Clinical and pathological characterization of Central Nervous System cryptococcosis in an experimental mouse model of stereotaxic intracerebral infection.
The Host Immune Response to <i>Scedosporium/Lomentospora</i> .
A 32-kilodalton hydrolase plays an important role in <i>Paracoccidioides brasiliensis</i> adherence to host cells and influences pathogenicity.
Phagocytosis of melanized <i>Aspergillus</i> conidia by macrophages exerts cytoprotective effects by sustained PI3K/Akt signalling.
The temporal dynamics of differential gene expression in <i>Aspergillus fumigatus</i> interacting with human immature dendritic cells in vitro.
How to invade a susceptible host: cellular aspects of aspergillosis.
Inhibition of neutrophil function following exposure to the <i>Aspergillus fumigatus</i> toxin fumagillin.
The Outcome of the <i>Cryptococcus neoformans</i> -Macrophage Interaction Depends on Phagolysosomal Membrane Integrity.
Mass Spectrometry-Based Proteomic and Immunoproteomic Analyses of the <i>Candida albicans</i> Hyphal Secretome Reveal Diagnostic Biomarker Candidates for Invasive Candidiasis.
Host-Pathogen Molecular Factors Contribute to the Pathogenesis of <i>Rhizopus</i> spp. in Diabetes Mellitus.
<i>Ustilago maydis</i> effectors and their impact on virulence.
Recent Advances in the Pathogenesis of Mucormycoses.
A secretory phospholipase A2 of a fungal pathogen contributes to lipid droplet homeostasis, assimilation of insect-derived lipids, and repression of host immune responses.
<i>Cryptococcus</i> escapes host immunity: What do we know?
Protein glycosylation during infection by plant pathogenic fungi.
MaNCP1, a C2H2 Zinc Finger Protein, Governs the Conidiation Pattern Shift through Regulating the Reductive Pathway for Nitric Oxide Synthesis in the Filamentous Fungus <i>Metarhizium acridum</i> .
Two glutathione peroxidases in the fungal pathogen <i>Cryptococcus neoformans</i> are expressed in the presence of specific substrates.
Chitosan Mediates Germling Adhesion in <i>Magnaporthe oryzae</i> and Is Required for Surface Sensing and Germling Morphogenesis.



Transcriptional profiling of <i>Scenedosporium apiospermum</i> enzymatic antioxidant gene battery unravels the involvement of thioredoxin reductases against chemical and phagocytic cells oxidative stress.
Gliotoxin Suppresses Macrophage Immune Function by Subverting Phosphatidylinositol 3,4,5-Trisphosphate Homeostasis.
<i>Paracoccidioides</i> Spp.: Virulence Factors and Immune-Evasion Strategies.
Proteomics of <i>Aspergillus fumigatus</i> Conidia-containing Phagolysosomes Identifies Processes Governing Immune Evasion.
Genes involved in <i>Beauveria bassiana</i> infection to <i>Galleria mellonella</i> .
Biotinylated Surfome Profiling Identifies Potential Biomarkers for Diagnosis and Therapy of <i>Aspergillus fumigatus</i> Infection.
Phagosomal F-Actin Retention by <i>Cryptococcus gattii</i> Induces Dendritic Cell Immunoparalysis.
Cordycepin, a metabolite of <i>Cordyceps militaris</i> , reduces immune-related gene expression in insects.
Uptake of the fluorescent probe FM4-64 by hyphae and haemolymph-derived in vivo hyphal bodies of the entomopathogenic fungus <i>Beauveria bassiana</i> .
Secretome analysis of <i>Aspergillus fumigatus</i> reveals Asp-hemolysin as a major secreted protein.
Human lysozyme has fungicidal activity against nasal fungi.
Promising whole-cell vaccines against cryptococcosis.
The potential management of oral candidiasis using anti-biofilm therapies.
The Trojan Horse Model in <i>Paracoccidioides</i> : A Fantastic Pathway to Survive Infecting Human Cells.
An Intracellular Arrangement of <i>Histoplasma capsulatum</i> Yeast-Aggregates Generates Nuclear Damage to the Cultured Murine Alveolar Macrophages.
Uptake of <i>Aspergillus fumigatus</i> Conidia by phagocytic and nonphagocytic cells in vitro: quantitation using strains expressing green fluorescent protein.
White-opaque switching of <i>Candida albicans</i> allows immune evasion in an environment-dependent fashion.
Single-Cell Analysis of Fungal Uptake in Cultured Airway Epithelial Cells Using Differential Fluorescent Staining and Imaging Flow Cytometry.
Clinical Manifestations and Treatment of Blastomycosis.
Subterranean termite prophylactic secretions and external antifungal defenses.
Proteomic Analysis of <i>Sporothrix schenckii</i> Exposed to Oxidative Stress Induced by Hydrogen Peroxide.
<i>Talaromyces marneffe</i> Mp1p Is a Virulence Factor that Binds and Sequesters a Key Proinflammatory Lipid to Dampen Host Innate Immune Response.
Neutrophil Extracellular Traps Go Viral.
How eukaryotic filamentous pathogens evade plant recognition.
Influenza Suppresses Neutrophil Recruitment to the Lung and Exacerbates Secondary Invasive Pulmonary Aspergillosis.
Comparative assessment of immune evasion mechanisms in human whole-blood infection assays by a systems biology approach.
GPI7-mediated glycosylphosphatidylinositol anchoring regulates appressorial penetration and immune evasion during infection of <i>Magnaporthe oryzae</i> .
A collagenous protective coat enables <i>Metarhizium anisopliae</i> to evade insect immune responses.
A decade after the emergence of <i>Candida auris</i> : what do we know?
Oxidative and nitrosative stress on phagocytes' function: from effective defense to immunity evasion mechanisms.
<i>Paracoccidioidomycosis</i> : eco-epidemiology, taxonomy and clinical and therapeutic issues.

Magnaporthe oryzae pathotype Triticum (MoT) can act as a heterologous expression system for fungal effectors with high transcript abundance in wheat.
Phenotypic switching of Cryptococcus neoformans can influence the outcome of the human immune response.
Mycophenolic Acid as a Promising Fungal Dimorphism Inhibitor to Control Sugar Cane Disease Caused by Sporisorium scitamineum.
A Genetic Screen for Saccharomyces cerevisiae Mutants That Fail to Enter Quiescence.
Unraveling the susceptibility of paracoccidioidomycosis: Insights towards the pathogen-immune interplay and immunogenetics.
Cryptococcus gattii is killed by dendritic cells, but evades adaptive immunity by failing to induce dendritic cell maturation.
Virulence Factors in Sporothrix schenckii, One of the Causative Agents of Sporotrichosis.
Cryptococcus neoformans UGT1 encodes a UDP-Galactose/UDP-GalNAc transporter.
Differential expression of Toll-like receptors in chronic hyperplastic candidosis.
Growth substrates and caleosin-mediated functions affect conidial virulence in the insect pathogenic fungus Beauveria bassiana.
Calcineurin A Is Essential in the Regulation of Asexual Development, Stress Responses and Pathogenesis in Talaromyces marneffei.
Talaromyces marneffei Infection: Virulence, Intracellular Lifestyle and Host Defense Mechanisms.
Preventing phagocytosis takes more than a sweet disposition.
The Aspergillus fumigatus conidial melanin production is regulated by the bifunctional bHLH DevR and MADS-box RlmA transcription factors.
Secreted protein MoHrip2 is required for full virulence of Magnaporthe oryzae and modulation of rice immunity.
Evasion of Innate Immune Responses by the Highly Virulent Cryptococcus gattii by Altering Capsule Glucuronoxylomannan Structure.
N-glycosylation of effector proteins by an $\alpha$ -1,3-mannosyltransferase is required for the rice blast fungus to evade host innate immunity.
Dynamic virulence-related regions of the plant pathogenic fungus Verticillium dahliae display enhanced sequence conservation.
Talaromyces marneffei promotes M2-like polarization of human macrophages by downregulating SOCS3 expression and activating the TLR9 pathway.
Cryptococcus gattii evades CD11b-mediated fungal recognition by coating itself with capsular polysaccharides.
MrSVP, a secreted virulence-associated protein, contributes to thermotolerance and virulence of the entomopathogenic fungus Metarhizium robertsii.
Histoplasma capsulatum modulates the immune response, affects proliferation and differentiation, and induces apoptosis of mesenchymal stromal cells.
Th1-Th2 cytokine kinetics in the bronchoalveolar lavage fluid of mice infected with Cryptococcus neoformans of different virulences.
Comparative transcriptome analysis unveils the adaptative mechanisms of Scedosporium apiospermum to the microenvironment encountered in the lungs of patients with cystic fibrosis.
Pathobiology of Pneumocystis pneumonia: life cycle, cell wall and cell signal transduction.
Expression of Cytokine Profiles in Human THP-1 Cells during Phase Transition of Talaromyces marneffei.
Cryptococcus gattii Capsule Blocks Surface Recognition Required for Dendritic Cell Maturation Independent of Internalization and Antigen Processing.

The fungal pathogen <i>Magnaporthe oryzae</i> suppresses innate immunity by modulating a host potassium channel.
Immunology of oral candidiasis.
Dissecting the Molecular Interactions between Wheat and the Fungal Pathogen <i>Zymoseptoria tritici</i> .
Oxidative stress response in <i>Paracoccidioides brasiliensis</i> : assessing catalase and cytochrome c peroxidase.
Is Antifungal Resistance a Cause for Treatment Failure in Dermatophytosis: A Study Focused on <i>Tinea Corporis</i> and <i>Cruris</i> from a Tertiary Centre?
Immune evasion or avoidance: fungal skin infection linked to reduced defence peptides in Australian green-eyed treefrogs, <i>Litoria serrata</i> .
Pathogenesis and treatment of chronic rhinosinusitis.
<i>Aspergillus</i> Hydrophobins: Physicochemical Properties, Biochemical Properties, and Functions in Solid Polymer Degradation.
Molecular antifungal defenses in subterranean termites: RNA interference reveals in vivo roles of termicins and GNBPs against a naturally encountered pathogen.
Comparative transcriptome analysis reveals significant differences in gene expression between appressoria and hyphae in <i>Colletotrichum gloeosporioides</i> .
Mechanism of <i>Metarhizium rileyi</i> evading cellular immune responses in <i>Helicoverpa armigera</i> .
Systems biology of infectious diseases: a focus on fungal infections.
Transcriptional profile of oil palm pathogen, <i>Ganoderma boninense</i> , reveals activation of lignin degradation machinery and possible evasion of host immune response.
Cross-talk between the Ras GTPase and the Hog1 survival pathways in response to nitrosative stress in <i>Paracoccidioides brasiliensis</i> .
Transcriptional response of <i>Aspergillus fumigatus</i> to copper and the role of the Cu chaperones.
When Defenses Fail: <i>Atelopus zeteki</i> Skin Secretions Increase Growth of the Pathogen <i>Batrachochytrium dendrobatidis</i> .
Pneumocystis: immune recognition and evasion.
A Critical Review on Communication Mechanism within Plant-Endophytic Fungi Interactions to Cope with Biotic and Abiotic Stresses.
Role of PLB1 in pulmonary inflammation and cryptococcal eicosanoid production.
The Immunologic Response to <i>Trichophyton Rubrum</i> in Lower Extremity Fungal Infections.
Malassezia-Can it be Ignored?
Endophytic <i>Beauveria bassiana</i> Induces Oxidative Stress and Enhances the Growth of <i>Fusarium oxysporum</i> -Infected Tomato Plants.
Increased susceptibility against <i>Cryptococcus neoformans</i> of lupus mouse models (pristane-induction and FcγRIIb deficiency) is associated with activated macrophage, regardless of genetic background.
<i>Histoplasma capsulatum</i> proteome response to decreased iron availability.
Microglial immune response is impaired against the neurotropic fungus <i>Lomentospora prolificans</i> .
Where Does the Peanut Smut Pathogen, <i>Thecaphora frezii</i> , Fit in the Spectrum of Smut Diseases?
Organization of chromosome ends in the rice blast fungus, <i>Magnaporthe oryzae</i> .
Proteins Potentially Involved in Immune Evasion Strategies in <i>Sporothrix brasiliensis</i> Elucidated by Ultra-High-Resolution Mass Spectrometry.
The State of the Art in Transcriptomics and Proteomics of Clinically Relevant <i>Sporothrix</i> Species.
<i>Paracoccidioides brasiliensis</i> interacts with dermal dendritic cells and keratinocytes in human skin and oral mucosa lesions.
The impact of the absence of Toll-like receptor-2 during <i>Sporothrix brasiliensis</i> infection.

43 kDa Glycoprotein (gp43) from <i>Paracoccidioides brasiliensis</i> Induced IL-17A and PGE2 Production by Human Polymorphonuclear Neutrophils: Involvement of TLR2 and TLR4.
Metabolites Involved in Immune Evasion by <i>Batrachochytrium dendrobatidis</i> Include the Polyamine Spermidine.
Nucleotide sequence-based analysis for determining the molecular epidemiology of <i>Penicillium marneffei</i> .
Valley fever: danger lurking in a dust cloud.
Do immune system changes at metamorphosis predict vulnerability to chytridiomycosis? An update.
Lymphocyte Inhibition by the Salamander-Killing Chytrid Fungus, <i>Batrachochytrium salamandrivorans</i> .
The hemolymph microbiome of insects.
Chronic dermatophytosis: what is special about <i>Trichophyton rubrum</i> ?
Gene Duplication and Mutation in the Emergence of a Novel Aggressive Allele of the AVR-Pik Effector in the Rice Blast Fungus.
Improved pharmacokinetic and biodistribution of 5-fluorouracil loaded biomimetic nanoerythrocytes decorated nanocarriers for liver cancer treatment.

Table A2-49, Cluster 48

Cluster 48 focuses on Type 1 interferon, emphasizing viral pathogen-induced mechanisms to antagonize mammalian interferon signaling pathways (729)
Type I Interferon Induced and Antagonized by Foot-and-Mouth Disease Virus.
Viral pathogen-induced mechanisms to antagonize mammalian interferon (IFN) signaling pathway.
Innate immune interferon responses to human immunodeficiency virus-1 infection.
MicroRNA-218 inhibits type I interferon production and facilitates virus immune evasion via targeting RIG-I.
Evasion mechanisms of the type I interferons responses by influenza A virus.
Hepatitis E Virus Methyltransferase Inhibits Type I Interferon Induction by Targeting RIG-I.
Porcine bocavirus NP1 protein suppresses type I IFN production by interfering with IRF3 DNA-binding activity.
The Molecular Mechanism of Herpes Simplex Virus 1 UL31 in Antagonizing the Activity of IFN- $\beta$ .
Signaling through RIG-I and type I interferon receptor: Immune activation by Newcastle disease virus in man versus immune evasion by Ebola virus (Review).
RIG-I-mediated antiviral signaling is inhibited in HIV-1 infection by a protease-mediated sequestration of RIG-I.
Interplay between interferon-mediated innate immunity and porcine reproductive and respiratory syndrome virus.
The 3C protein of enterovirus 71 inhibits retinoid acid-inducible gene I-mediated interferon regulatory factor 3 activation and type I interferon responses.
Caspase-Mediated Regulation and Cellular Heterogeneity of the cGAS/STING Pathway in Kaposi's Sarcoma-Associated Herpesvirus Infection.
Herpes Simplex Virus 1 UL36USP Antagonizes Type I Interferon-Mediated Antiviral Innate Immunity.
Induction and evasion of type I interferon responses by influenza viruses.
The Role of Type I IFNs in Influenza: Antiviral Superheroes or Immunopathogenic Villains?
Anti-chicken type I IFN countermeasures by major avian RNA viruses.
The US3 Kinase of Herpes Simplex Virus Phosphorylates the RNA Sensor RIG-I To Suppress Innate Immunity.

Influenza C virus NS1 protein counteracts RIG-I-mediated IFN signalling.
The type I interferon response during viral infections: a "SWOT" analysis.
Influenza A virus NS1 resembles a TRAF3-interacting motif to target the RNA sensing-TRAF3-type I IFN axis and impair antiviral innate immunity.
IL-36 promotes anti-viral immunity by boosting sensitivity to IFN- $\alpha/\beta$ in IRF1 dependent and independent manners.
Downregulation of microRNA miR-526a by enterovirus inhibits RIG-I-dependent innate immune response.
Proteolysis of MDA5 and IPS-1 is not required for inhibition of the type I IFN response by poliovirus.
Cleavage of 14-3-3 $\epsilon$ by the enteroviral 3C protease dampens RIG-I-mediated antiviral signaling.
Negative regulation of cytoplasmic RNA-mediated antiviral signaling.
Evasion and subversion of interferon-mediated antiviral immunity by Kaposi's sarcoma-associated herpesvirus: an overview.
A selective contribution of the RIG-I-like receptor pathway to type I interferon responses activated by cytosolic DNA.
Enterovirus 71 inhibits cellular type I interferon signaling by inhibiting host RIG-I ubiquitination.
Influenza A Virus PA Antagonizes Interferon- $\beta$ by Interacting with Interferon Regulatory Factor 3.
The Human Papillomavirus E6 Oncoprotein Targets USP15 and TRIM25 To Suppress RIG-I-Mediated Innate Immune Signaling.
Induction and control of the type I interferon pathway by Bluetongue virus.
Chicken anemia virus VP1 negatively regulates type I interferon via targeting interferon regulatory factor 7 of the DNA-sensing pathway.
Hemagglutinin of Influenza A Virus Antagonizes Type I Interferon (IFN) Responses by Inducing Degradation of Type I IFN Receptor 1.
On taking the STING out of immune activation.
Innate immunity evasion by Dengue virus.
EBV MicroRNA BART16 Suppresses Type I IFN Signaling.
Enterovirus 71 blocks selectively type I interferon production through the 3C viral protein in mice.
How Dengue Virus Circumvents Innate Immunity.
Type I IFN--a blunt spear in fighting HIV-1 infection.
An unbiased genetic screen reveals the polygenic nature of the influenza virus anti-interferon response.
Dengue virus regulates type I interferon signalling in a strain-dependent manner in human cell lines.
Mycobacterium tuberculosis MmsA (Rv0753c) Interacts with STING and Blunts the Type I Interferon Response.
HIV blocks Type I IFN signaling through disruption of STAT1 phosphorylation.
The C-Terminal Effector Domain of Non-Structural Protein 1 of Influenza A Virus Blocks IFN- $\beta$ Production by Targeting TNF Receptor-Associated Factor 3.
Induction and control of the type I interferon pathway by Bluetongue virus.
Caspase-Dependent Suppression of Type I Interferon Signaling Promotes Kaposi's Sarcoma-Associated Herpesvirus Lytic Replication.
Vaccinia Virus Protein C6 Inhibits Type I IFN Signalling in the Nucleus and Binds to the Transactivation Domain of STAT2.
Disruption of type I interferon signaling by the nonstructural protein of severe fever with thrombocytopenia syndrome virus via the hijacking of STAT2 and STAT1 into inclusion bodies.
West Nile Virus NS1 Antagonizes Interferon Beta Production by Targeting RIG-I and MDA5.

Rotavirus nonstructural protein 1 antagonizes innate immune response by interacting with retinoic acid inducible gene I.
Viral tricks to grid-lock the type I interferon system.
Feline Infectious Peritonitis Virus Nsp5 Inhibits Type I Interferon Production by Cleaving NEMO at Multiple Sites.
Histone deacetylase 4 promotes type I interferon signaling, restricts DNA viruses, and is degraded via vaccinia virus protein C6.
Early IFN type I response: Learning from microbial evasion strategies.
The Cytomegalovirus M35 Protein Directly Binds to the Interferon- $\beta$ Enhancer and Modulates Transcription of Ifnb1 and Other IRF3-Driven Genes.
Dengue Virus Subverts Host Innate Immunity by Targeting Adaptor Protein MAVS.
HIV Blocks Interferon Induction in Human Dendritic Cells and Macrophages by Dysregulation of TBK1.
Dengue Virus NS Proteins Inhibit RIG-I/MAVS Signaling by Blocking TBK1/IRF3 Phosphorylation: Dengue Virus Serotype 1 NS4A Is a Unique Interferon-Regulating Virulence Determinant.
Hazara Orthonairovirus Nucleoprotein Antagonizes Type I Interferon Production by Inhibition of RIG-I Ubiquitination.
Porcine epidemic diarrhea virus nucleocapsid protein antagonizes beta interferon production by sequestering the interaction between IRF3 and TBK1.
Interferon induction by RNA viruses and antagonism by viral pathogens.
A RIG-I-like receptor directs antiviral responses to a bunyavirus and is antagonized by virus-induced blockade of TRIM25-mediated ubiquitination.
The Interplay Between Coronavirus and Type I IFN Response.
A Tug of War: DNA-Sensing Antiviral Innate Immunity and Herpes Simplex Virus Type I Infection.
Evasion of host antiviral innate immunity by HSV-1, an update.
Human microRNA-30 inhibits influenza virus infection by suppressing the expression of SOCS1, SOCS3, and NEDD4.
Inhibition of DNA-Sensing Pathway by Marek's Disease Virus VP23 Protein through Suppression of Interferon Regulatory Factor 7 Activation.
Targeting Type I Interferon Induction and Signaling: How Zika Virus Escapes from Host Innate Immunity.
Herpes simplex virus 1 targets IRF7 via ICP0 to limit type I IFN induction.
Pseudorabies Virus dUTPase UL50 Induces Lysosomal Degradation of Type I Interferon Receptor 1 and Antagonizes the Alpha Interferon Response.
Inhibition of the type I interferon response in human dendritic cells by dengue virus infection requires a catalytically active NS2B3 complex.
Bovine coronavirus nucleocapsid suppresses IFN- $\beta$ production by inhibiting RIG-I-like receptors pathway in host cells.
A Type I Interferon and IL-10 Induced by Orientia tsutsugamushi Infection Suppresses Antigen-Specific T Cells and Their Memory Responses.
Zika virus elicits inflammation to evade antiviral response by cleaving cGAS via NS1-caspase-1 axis.
Evasion of I Interferon-Mediated Innate Immunity by Pseudorabies Virus.
Autocrine activation of the IFN signaling pathway may promote immune escape in glioblastoma.
Viral evasion of the interferon response at a glance.
Mechanisms of type III interferon expression.
RNA Sensors as a Mechanism of Innate Immune Evasion among SARSCoV2, HIV and Nipah Viruses.
Type I Interferons Direct Gammaherpesvirus Host Colonization.

Human Cytomegalovirus UL23 Attenuates Signal Transducer and Activator of Transcription 1 Phosphorylation and Type I Interferon Response.
Caprine MAVS Is a RIG-I Interacting Type I Interferon Inducer Downregulated by Peste des Petits Ruminants Virus Infection.
Coxsackievirus counters the host innate immune response by blocking type III interferon expression.
Reprogramming viral immune evasion for a rational design of next-generation vaccines for RNA viruses.
Hepatitis C Virus Infection Is Inhibited by a Noncanonical Antiviral Signaling Pathway Targeted by NS3-NS4A.
"Toll-free" pathways for production of type I interferons.
Avian bornaviruses escape recognition by the innate immune system.
HIV-1, interferon and the interferon regulatory factor system: an interplay between induction, antiviral responses and viral evasion.
Inhibition of antiviral innate immunity by birnavirus VP3 protein via blockage of viral double-stranded RNA binding to the host cytoplasmic RNA detector MDA5.
RIG-I-mediated antiviral responses to single-stranded RNA bearing 5'-phosphates.
Beyond Good and Evil: Molecular Mechanisms of Type I and III IFN Functions.
Mechanism of inhibiting type I interferon induction by hepatitis B virus X protein.
Viral dedication to vigorous destruction of interferon receptors.
Pseudorabies Virus DNA Polymerase Processivity Factor UL42 Inhibits Type I IFN Response by Preventing ISGF3-ISRE Interaction.
Insights into antiviral innate immunity revealed by studying hepatitis C virus.
Hepatitis E Virus: How It Escapes Host Innate Immunity.
Encephalomyocarditis virus 3C protease attenuates type I interferon production through disrupting the TANK-TBK1-IKK $\epsilon$ -IRF3 complex.
Innate Immune Sensing of Influenza A Virus.
Negative Regulation of RIG-I by Tim-3 Promotes H1N1 Infection.
The game between host antiviral innate immunity and immune evasion strategies of senecavirus A - A cell biological perspective.
Inhibition of Type III Interferon Expression in Intestinal Epithelial Cells-A Strategy Used by Coxsackie B Virus to Evade the Host's Innate Immune Response at the Primary Site of Infection?
Role of metapneumoviral glycoproteins in the evasion of the host cell innate immune response.
Seneca Valley Virus Suppresses Host Type I Interferon Production by Targeting Adaptor Proteins MAVS, TRIF, and TANK for Cleavage.
$\beta$ -Catenin Is Required for the cGAS/STING Signaling Pathway but Antagonized by the Herpes Simplex Virus 1 US3 Protein.
Hepatitis A and hepatitis C viruses: divergent infection outcomes marked by similarities in induction and evasion of interferon responses.
Zika Virus Antagonizes Type I Interferon Responses during Infection of Human Dendritic Cells.
A critical function for type I interferons in cancer immunoediting.
Type III Interferon-Mediated Signaling Is Critical for Controlling Live Attenuated Yellow Fever Virus Infection In Vivo.
Zika Virus Non-structural Protein 4A Blocks the RLR-MAVS Signaling.
Induction of Siglec-G by RNA viruses inhibits the innate immune response by promoting RIG-I degradation.
Filoviral immune evasion mechanisms.

MAPK Phosphatase 5 Expression Induced by Influenza and Other RNA Virus Infection Negatively Regulates IRF3 Activation and Type I Interferon Response.
The Swine IFN System in Viral Infections: Major Advances and Translational Prospects.
Transmissible gastroenteritis virus does not suppress IFN- $\beta$ induction but is sensitive to IFN in IPEC-J2 cells.
A Rhesus Rhadinovirus Viral Interferon (IFN) Regulatory Factor Is Virion Associated and Inhibits the Early IFN Antiviral Response.
Small-molecule inhibitors of ubiquitin-specific protease 7 enhance type-I interferon antiviral efficacy by destabilizing SOCS1.
Evasion of the mucosal innate immune system by herpes simplex virus type 2.
Host Innate Immunity against Hepatitis E Virus and Viral Evasion Mechanisms.
Lymphocytic Choriomeningitis Virus Differentially Affects the Virus-Induced Type I Interferon Response and Mitochondrial Apoptosis Mediated by RIG-I/MAVS.
TOB1 attenuates IRF3-directed antiviral responses by recruiting HDAC8 to specifically suppress IFN- $\beta$ expression.
Human metapneumovirus M2-2 protein inhibits RIG-I signaling by preventing TRIM25-mediated RIG-I ubiquitination.
Overcoming NS1-mediated immune antagonism involves both interferon-dependent and independent mechanisms.
Molecular Mechanisms of Foot-and-Mouth Disease Virus Targeting the Host Antiviral Response.
Casein kinase II controls TBK1/IRF3 activation in IFN response against viral infection.
Enterovirus 2Apro targets MDA5 and MAVS in infected cells.
The interaction between human enteroviruses and type I IFN signaling pathway.
Human Beta Papillomavirus Type 8 E1 and E2 Proteins Suppress the Activation of the RIG-I-Like Receptor MDA5.
LncNSPL facilitates influenza A viral immune escape by restricting TRIM25-mediated K63-linked RIG-I ubiquitination.
Foot-and-Mouth Disease Virus Viroporin 2B Antagonizes RIG-I-Mediated Antiviral Effects by Inhibition of Its Protein Expression.
Zika Virus Proteins NS2A and NS4A Are Major Antagonists that Reduce IFN- $\beta$ Promoter Activity Induced by the MDA5/RIG-I Signaling Pathway.
Cleavage of TANK-Binding Kinase 1 by HIV-1 Protease Triggers Viral Innate Immune Evasion.
Type I interferon inhibition and dendritic cell activation during gammaherpesvirus respiratory infection.
MiR-202-5p Inhibits RIG-I-Dependent Innate Immune Responses to RGNNV Infection by Targeting TRIM25 to Mediate RIG-I Ubiquitination.
MOV10 Provides Antiviral Activity against RNA Viruses by Enhancing RIG-I-MAVS-Independent IFN Induction.
To interfere and to anti-interfere: the interplay between hepatitis C virus and interferon.
Ten Strategies of Interferon Evasion by Viruses.
Viruses evade the immune system through type I interferon-mediated STAT2-dependent, but STAT1-independent, signaling.
Decoding type I and III interferon signalling during viral infection.
Topoisomerase II Inhibitors Induce DNA Damage-Dependent Interferon Responses Circumventing Ebola Virus Immune Evasion.
Seneca Valley virus 2C and 3C inhibit type I interferon production by inducing the degradation of RIG-I.



DENV inhibits type I IFN production in infected cells by cleaving human STING.
The Coronavirus Transmissible Gastroenteritis Virus Evades the Type I Interferon Response through IRE1 $\alpha$ -Mediated Manipulation of the MicroRNA miR-30a-5p/SOCS1/3 Axis.
Influenza A virus NS1 induces degradation of sphingosine 1-phosphate lyase to obstruct the host innate immune response.
Dengue Virus 2 NS2B Targets MAVS and IKK $\epsilon$ to Evade the Antiviral Innate Immune Response.
RNAs Containing Modified Nucleotides Fail To Trigger RIG-I Conformational Changes for Innate Immune Signaling.
Myxoma virus selectively disrupts type I interferon signaling in primary human fibroblasts by blocking the activation of the Janus kinase Tyk2.
Ebolavirus Species-Specific Interferon Antagonism Mediated by VP24.
La Piedad Michoacán Mexico Virus V protein antagonizes type I interferon response by binding STAT2 protein and preventing STATs nuclear translocation.
Porcine deltacoronavirus nucleocapsid protein antagonizes IFN- $\beta$ production by impairing dsRNA and PACT binding to RIG-I.
Hepatitis C Virus Evasion from RIG-I-Dependent Hepatic Innate Immunity.
Equine herpesvirus-1 suppresses type-I interferon induction in equine endothelial cells.
Japanese Encephalitis Virus NS5 Inhibits Type I Interferon (IFN) Production by Blocking the Nuclear Translocation of IFN Regulatory Factor 3 and NF- $\kappa$ B.
Double Plant Homeodomain Fingers 2 (DPF2) Promotes the Immune Escape of Influenza Virus by Suppressing Beta Interferon Production.
Innate immune response and viral interference strategies developed by human herpesviruses.
Nonstructural Protein of Severe Fever with Thrombocytopenia Syndrome Phlebovirus Inhibits TBK1 to Evade Interferon-Mediated Response.
Interferon-independent STING signaling promotes resistance to HSV-1 in vivo.
HSV-1 ICP27 targets the TBK1-activated STING signalsome to inhibit virus-induced type I IFN expression.
Adaptive Immune Responses to Zika Virus Are Important for Controlling Virus Infection and Preventing Infection in Brain and Testes.
Regulation of MAVS Expression and Signaling Function in the Antiviral Innate Immune Response.
Herpes Simplex Virus 1 Tegument Protein UL46 Inhibits TANK-Binding Kinase 1-Mediated Signaling.
The human cytomegalovirus tegument protein pp65 (pUL83): a key player in innate immune evasion.
Mismatches in the Influenza A Virus RNA Panhandle Prevent Retinoic Acid-Inducible Gene I (RIG-I) Sensing by Impairing RNA/RIG-I Complex Formation.
Enterovirus 71 protease 2Apro targets MAVS to inhibit anti-viral type I interferon responses.
The MAVS Immune Recognition Pathway in Viral Infection and Sepsis.
Interplay between Intrinsic and Innate Immunity during HIV Infection.
Type I Interferon Signaling to Dendritic Cells Limits Murid Herpesvirus 4 Spread from the Olfactory Epithelium.
TRIM proteins: another class of viral victims.
Evasion of the STING DNA-Sensing Pathway by VP11/12 of Herpes Simplex Virus 1.
The papain-like protease of porcine epidemic diarrhea virus negatively regulates type I interferon pathway by acting as a viral deubiquitinase.
Paramyxovirus accessory proteins as interferon antagonists.
Mechanism of Rhinovirus Immunity and Asthma.
Structural basis for concerted recruitment and activation of IRF-3 by innate immune adaptor proteins.

Duck plague virus UL41 protein inhibits RIG-I/MDA5-mediated duck IFN- $\beta$ production via mRNA degradation activity.
HIV Triggers a cGAS-Dependent, Vpu- and Vpr-Regulated Type I Interferon Response in CD4(+) T Cells.
The virion host shutoff protein of herpes simplex virus 1 blocks the replication-independent activation of NF- $\kappa$ B in dendritic cells in the absence of type I interferon signaling.
Innate immune escape by Dengue and West Nile viruses.
Porcine epidemic diarrhea virus E protein suppresses RIG-I signaling-mediated interferon- $\beta$ production.
West Nile virus noncoding subgenomic RNA contributes to viral evasion of the type I interferon-mediated antiviral response.
Pleiotropic Roles of Type 1 Interferons in Antiviral Immune Responses.
Respiratory syncytial virus proteins modulate suppressors of cytokine signaling 1 and 3 and the type I interferon response to infection by a toll-like receptor pathway.
Innate immune evasion strategies of DNA and RNA viruses.
Latent Membrane Protein 1 of Epstein-Barr Virus Promotes RIG-I Degradation Mediated by Proteasome Pathway.
Parainfluenza Virus 3 Blocks Antiviral Mediators Downstream of the Interferon Lambda Receptor by Modulating Stat1 Phosphorylation.
Screening interferon antagonists from accessory proteins encoded by P gene for immune escape of Caprine parainfluenza virus 3.
Structural insights into the distinctive RNA recognition and therapeutic potentials of RIG-I-like receptors.
Adenovirus evasion of interferon-mediated innate immunity by direct antagonism of a cellular histone posttranslational modification.
Enterovirus 71-induced has-miR-21 contributes to evasion of host immune system by targeting MyD88 and IRAK1.
KSHV-encoded viral interferon regulatory factor 4 (vIRF4) interacts with IRF7 and inhibits interferon alpha production.
Innate immune evasion by filoviruses.
The Protein Kinase Receptor Modulates the Innate Immune Response against Tacaribe Virus.
Viral Evasion of RIG-I-Like Receptor-Mediated Immunity through Dysregulation of Ubiquitination and ISGylation.
Feline Herpesvirus 1 US3 Blocks the Type I Interferon Signal Pathway by Targeting Interferon Regulatory Factor 3 Dimerization in a Kinase-Independent Manner.
Defective Influenza A Virus RNA Products Mediate MAVS-Dependent Upregulation of Human Leukocyte Antigen Class I Proteins.
Antagonism of type I interferon by flaviviruses.
Ebolavirus VP35 suppresses IFN production from conventional but not plasmacytoid dendritic cells.
Herpes Simplex Virus 1 Serine Protease VP24 Blocks the DNA-Sensing Signal Pathway by Abrogating Activation of Interferon Regulatory Factor 3.
Foreign RNA induces the degradation of mitochondrial antiviral signaling protein (MAVS): the role of intracellular antiviral factors.
DHAV 3CD targets IRF7 and RIG-I proteins to block the type I interferon upstream signaling pathway.
Replication of Human Norovirus RNA in Mammalian Cells Reveals Lack of Interferon Response.
Rhesus Macaque Rhadinovirus Encodes a Viral Interferon Regulatory Factor To Disrupt Promyelocytic Leukemia Nuclear Bodies and Antagonize Type I Interferon Signaling.

KAP1-Mediated Epigenetic Suppression in Anti-RNA Viral Responses by Direct Targeting RIG-I and MDA5.
Myxoma Virus dsRNA Binding Protein M029 Inhibits the Type I IFN-Induced Antiviral State in a Highly Species-Specific Fashion.
The cGAS-STING Defense Pathway and Its Counteraction by Viruses.
Antiviral Properties of ISG15.
Measles virus suppresses RIG-I-like receptor activation in dendritic cells via DC-SIGN-mediated inhibition of PP1 phosphatases.
Interferons and viruses: an evolutionary arms race of molecular interactions.
Pseudorabies Virus Tegument Protein UL13 Suppresses RLR-Mediated Antiviral Innate Immunity through Regulating Receptor Transcription.
Cleavage of interferon regulatory factor 7 by enterovirus 71 3C suppresses cellular responses.
Inhibition of type I interferon production via suppressing IKK-gamma expression: a new strategy for counteracting host antiviral defense by influenza A viruses?
Zika virus antagonizes interferon response in patients and disrupts RIG-I-MAVS interaction through its CARD-TM domains.
Processing of genome 5' termini as a strategy of negative-strand RNA viruses to avoid RIG-I-dependent interferon induction.
Participation of interferon type I during canine parvovirus infection.
Herpes simplex virus type 2 virion host shutoff protein suppresses innate dsRNA antiviral pathways in human vaginal epithelial cells.
Herpes Simplex Virus 1 Tegument Protein VP22 Abrogates cGAS/STING-Mediated Antiviral Innate Immunity.
Collateral Damage during Dengue Virus Infection: Making Sense of DNA by cGAS.
Strategies for immune evasion by human tumor viruses.
Small Hero with Great Powers: Vaccinia Virus E3 Protein and Evasion of the Type I IFN Response.
An arms race: innate antiviral responses and counteracting viral strategies.
Grass Carp Reovirus (GCRV) Giving Its All to Suppress IFN Production by Countering MAVS Signaling Transduction.
HCV-induced miR-21 contributes to evasion of host immune system by targeting MyD88 and IRAK1.
Threonine 80 phosphorylation of non-structural protein 1 regulates the replication of influenza A virus by reducing the binding affinity with RIG-I.
Camouflage and interception: how pathogens evade detection by intracellular nucleic acid sensors.
Vpu mediates depletion of interferon regulatory factor 3 during HIV infection by a lysosome-dependent mechanism.
Species-Specific Deamidation of RIG-I Reveals Collaborative Action between Viral and Cellular Deamidases in HSV-1 Lytic Replication.
The FDA-Approved Oral Drug Nitazoxanide Amplifies Host Antiviral Responses and Inhibits Ebola Virus.
Differential expression of type I interferon mRNA and protein levels induced by virulent Marek's disease virus infection in chickens.
Encephalomyocarditis Virus Abrogates the Interferon Beta Signaling Pathway via Its Structural Protein VP2.
Porcine deltacoronavirus (PDCoV) infection suppresses RIG-I-mediated interferon- $\beta$ production.
Porcine Epidemic Diarrhea Virus Infection Inhibits Interferon Signaling by Targeted Degradation of STAT1.
Foot-and-mouth disease virus non-structural protein 3A inhibits the interferon- $\beta$ signaling pathway.

Human Papillomavirus E7 Oncoprotein Subverts Host Innate Immunity via SUV39H1-Mediated Epigenetic Silencing of Immune Sensor Genes.
Interferon-alpha production by swine dendritic cells is inhibited during acute infection with foot-and-mouth disease virus.
Spring Viremia of Carp Virus N Protein Suppresses Fish IFN $\phi$ 1 Production by Targeting the Mitochondrial Antiviral Signaling Protein.
Social evolution of innate immunity evasion in a virus.
Negative regulation of TBK1-mediated antiviral immunity.
One Step Ahead: Herpesviruses Light the Way to Understanding Interferon-Stimulated Genes (ISGs).
The Matrix Protein of Nipah Virus Targets the E3-Ubiquitin Ligase TRIM6 to Inhibit the IKK $\epsilon$ Kinase-Mediated Type-I IFN Antiviral Response.
Vpu-deficient HIV strains stimulate innate immune signaling responses in target cells.
Immune evasion of porcine enteric coronaviruses and viral modulation of antiviral innate signaling.
Foot-and-mouth disease virus non-structural protein 2B negatively regulates the RLR-mediated IFN- $\beta$ induction.
Suppression of cGAS- and RIG-I-mediated innate immune signaling by Epstein-Barr virus deubiquitinase BPLF1.
N(6)-Methyladenosine modification of hepatitis B and C viral RNAs attenuates host innate immunity via RIG-I signaling.
Kisspeptin/GPR54 signaling restricts antiviral innate immune response through regulating calcineurin phosphatase activity.
A de novo transcriptome analysis shows that modulation of the JAK-STAT signaling pathway by salmonid alphavirus subtype 3 favors virus replication in macrophage/dendritic-like TO-cells.
HIV-1 evades innate immune recognition through specific cofactor recruitment.
HSV1 VP1-2 deubiquitinates STING to block type I interferon expression and promote brain infection.
Zika virus NS2A inhibits interferon signaling by degradation of STAT1 and STAT2.
Modulation of innate immune signaling by nonstructural protein 1 (nsp1) in the family Arteriviridae.
Differences in the ability to suppress interferon $\beta$ production between allele A and allele B NS1 proteins from H10 influenza A viruses.
Attack, parry and riposte: molecular fencing between the innate immune system and human herpesviruses.
A Kaposi's sarcoma-associated herpesviral protein inhibits virus-mediated induction of type I interferon by blocking IRF-7 phosphorylation and nuclear accumulation.
Viral Inhibition of PRR-Mediated Innate Immune Response: Learning from KSHV Evasion Strategies.
Pestiviruses infection: Interferon-virus mutual regulation.
Porcine Deltacoronavirus Accessory Protein NS6 Antagonizes Interferon Beta Production by Interfering with the Binding of RIG-I/MDA5 to Double-Stranded RNA.
Duck Tembusu Virus Infection Promotes the Expression of Duck Interferon-Induced Protein 35 to Counteract RIG-I Antiviral Signaling in Duck Embryo Fibroblasts.
HIV-1 Vif suppresses antiviral immunity by targeting STING.
RIG-I-like receptor regulation in virus infection and immunity.
Vaccinia virus blocks Stat1-dependent and Stat1-independent gene expression induced by type I and type II interferons.
Influenza virus protein PB1-F2 inhibits the induction of type I interferon by binding to MAVS and decreasing mitochondrial membrane potential.
Identification of a Natural Viral RNA Motif That Optimizes Sensing of Viral RNA by RIG-I.
Japanese Encephalitis Virus NS1' Protein Antagonizes Interferon Beta Production.

Evidence against the Human Metapneumovirus G, SH, and M2-2 Proteins as Bona Fide Interferon Antagonists.
Single point mutations in the helicase domain of the NS3 protein enhance dengue virus replicative capacity in human monocyte-derived dendritic cells and circumvent the type I interferon response.
Nuclear localization of Zika virus NS5 contributes to suppression of type I interferon production and response.
PPM1G restricts innate immune signaling mediated by STING and MAVS and is hijacked by KSHV for immune evasion.
ERK inhibition aids IFN- $\beta$ promoter activation during EV71 infection by blocking CRYAB degradation in SH-SY5Y cells.
The X proteins of bornaviruses interfere with type I interferon signalling.
Interplay of measles virus with early induced cytokines reveals different wild type phenotypes.
Evasion of Host Antiviral Innate Immunity by Paramyxovirus Accessory Proteins.
Grass Carp Reovirus Nonstructural Proteins Avoid Host Antiviral Immune Response by Targeting the RLR Signaling Pathway.
Foot-and-Mouth Disease Virus Evades Innate Immune Response by 3C-Targeting of MDA5.
HSV-1-encoded ICPO degrades the host deubiquitinase BRCC36 to antagonize interferon antiviral response.
Seneca Valley Virus 3C protease negatively regulates the type I interferon pathway by acting as a viral deubiquitinase.
Heartland virus antagonizes type I and III interferon antiviral signaling by inhibiting phosphorylation and nuclear translocation of STAT2 and STAT1.
Structural basis for m7G recognition and 2'-O-methyl discrimination in capped RNAs by the innate immune receptor RIG-I.
MicroRNA-373 facilitates HSV-1 replication through suppression of type I IFN response by targeting IRF1.
Herpesviral infection and Toll-like receptor 2.
The Strategy of Picornavirus Evading Host Antiviral Responses: Non-structural Proteins Suppress the Production of IFNs.
Molecular basis for ebolavirus VP35 suppression of human dendritic cell maturation.
Evasion of interferon-mediated immune response by arteriviruses.
HBV confers innate immune evasion through triggering HAT1/acetylation of H4K5/H4K12/miR-181a-5p or KPNA2/cGAS-STING/IFN-I signaling.
OASL-a new player in controlling antiviral innate immunity.
Enterovirus 71 Proteins 2A and 3D Antagonize the Antiviral Activity of Gamma Interferon via Signaling Attenuation.
Differential Antagonism of Human Innate Immune Responses by Tick-Borne Phlebovirus Nonstructural Proteins.
The interplay between viruses and innate immune signaling: recent insights and therapeutic opportunities.
Strategies of highly pathogenic RNA viruses to block dsRNA detection by RIG-I-like receptors: hide, mask, hit.
The Small t Antigen of JC Virus Antagonizes RIG-I-Mediated Innate Immunity by Inhibiting TRIM25's RNA Binding Ability.
Peste des petits ruminants virus non-structural C protein inhibits the induction of interferon- $\beta$ by potentially interacting with MAVS and RIG-I.
PGRMC1 Exerts Its Function of Anti-Influenza Virus in the Central Nervous System.

Viral pseudo-enzymes activate RIG-I via deamidation to evade cytokine production.
HSV-2 inhibits type-I interferon signaling via multiple complementary and compensatory STAT2-associated mechanisms.
Human bocavirus NP1 inhibits IFN- $\beta$ production by blocking association of IFN regulatory factor 3 with IFNB promoter.
Identification of multiple RIG-I-specific pathogen associated molecular patterns within the West Nile virus genome and antigenome.
Toxoplasma gondii Inhibits gamma interferon (IFN- $\gamma$ )- and IFN- $\beta$ -induced host cell STAT1 transcriptional activity by increasing the association of STAT1 with DNA.
Feline Panleucopenia Virus NS2 Suppresses the Host IFN- $\beta$ Induction by Disrupting the Interaction between TBK1 and STING.
Zika Virus NS3 Mimics a Cellular 14-3-3-Binding Motif to Antagonize RIG-I- and MDA5-Mediated Innate Immunity.
Interferon response and viral evasion by members of the family rhabdoviridae.
Grass carp reovirus VP56 represses interferon production by degrading phosphorylated IRF7.
Modulation of Innate Immune Signaling Pathways by Herpesviruses.
The autophagy-related degradation of MDA5 by Tembusu virus nonstructural 2B disrupts IFN $\beta$ production.
Viral IFN-regulatory factors inhibit activation-induced cell death via two positive regulatory IFN-regulatory factor 1-dependent domains in the CD95 ligand promoter.
Interplay of foot and mouth disease virus with cell-mediated and humoral immunity of host.
Hepatitis C virus NS4B induces the degradation of TRIF to inhibit TLR3-mediated interferon signaling pathway.
DNA tumor virus oncogenes antagonize the cGAS-STING DNA-sensing pathway.
The Measles Virus V Protein Binding Site to STAT2 Overlaps That of IRF9.
Cytokine response in mouse bone marrow derived macrophages after infection with pathogenic and non-pathogenic Rift Valley fever virus.
Transcription of interferon stimulated genes in response to Porcine rubulavirus infection in vitro.
Structural basis for dsRNA recognition and interferon antagonism by Ebola VP35.
Strategies to Target ISG15 and USP18 Toward Therapeutic Applications.
Coronavirus virulence genes with main focus on SARS-CoV envelope gene.
Expression of interferon gamma in the brain of cats with natural Borna disease virus infection.
Acute bovine viral diarrhea virus infection inhibits expression of interferon tau-stimulated genes in bovine endometrium.
Antagonism of dsRNA-Induced Innate Immune Pathways by NS4a and NS4b Accessory Proteins during MERS Coronavirus Infection.
Dugbe virus ovarian tumour domain interferes with ubiquitin/ISG15-regulated innate immune cell signalling.
Coronavirus membrane-associated papain-like proteases induce autophagy through interacting with Beclin1 to negatively regulate antiviral innate immunity.
Dissociation of a MAVS/IPS-1/VISA/Cardif-IKKepsilon molecular complex from the mitochondrial outer membrane by hepatitis C virus NS3-4A proteolytic cleavage.
Regulation of the interferon-inducible 2'-5'-oligoadenylate synthetases by adenovirus VA(I) RNA. [EV71 Infection and Innate Antiviral Innate Immunity].
$\gamma$ HV68 vGAT: a viral pseudoenzyme pimping for PAMPs.
Singapore grouper iridovirus VP122 targets grouper STING to evade the interferon immune response.
A Secreted Viral Nonstructural Protein Determines Intestinal Norovirus Pathogenesis.

Overlapping Patterns of Rapid Evolution in the Nucleic Acid Sensors cGAS and OAS1 Suggest a Common Mechanism of Pathogen Antagonism and Escape.
Innate immune sensor LGP2 is cleaved by the Leader protease of foot-and-mouth disease virus.
Innate Immune Response to Cytoplasmic DNA: Mechanisms and Diseases.
Coronaviral PLpro proteases and the immunomodulatory roles of conjugated versus free Interferon Stimulated Gene product-15 (ISG15).
The Glycoprotein and Nucleocapsid Protein of Hantaviruses Manipulate Autophagy Flux to Restrain Host Innate Immune Responses.
SPI-2/CrmA inhibits IFN- $\beta$ induction by targeting TBK1/IKK $\epsilon$ .
Antiviral and Inflammatory Cellular Signaling Associated with Enterovirus 71 Infection.
dsRNA binding characterization of full length recombinant wild type and mutants Zaire ebolavirus VP35.
Arm race between Rift Valley fever virus and host.
Molecular Pathogenesis and Immune Evasion of Vesicular Stomatitis New Jersey Virus Inferred from Genes Expression Changes in Infected Porcine Macrophages.
Insights into the homo-oligomerization properties of N-terminal coiled-coil domain of Ebola virus VP35 protein.
Enabled interferon signaling evasion in an immune-competent transgenic mouse model of parainfluenza virus 5 infection.
Inhibition of Antiviral Innate Immunity by Foot-and-Mouth Disease Virus L(pro) through Interaction with the N-Terminal Domain of Swine RNase L.
Measles virus V protein inhibits NLRP3 inflammasome-mediated interleukin-1 $\beta$ secretion.
Innate immune responses to duck Tembusu virus infection.
Middle East Respiratory Coronavirus Accessory Protein 4a Inhibits PKR-Mediated Antiviral Stress Responses.
Structure-dependent modulation of alpha interferon production by porcine circovirus 2 oligodeoxyribonucleotide and CpG DNAs in porcine peripheral blood mononuclear cells.
The hepatitis B e antigen (HBeAg) targets and suppresses activation of the toll-like receptor signaling pathway.
Dissociation of paramyxovirus interferon evasion activities: universal and virus-specific requirements for conserved V protein amino acids in MDA5 interference.
CD95 Signaling Inhibits B Cell Receptor-Mediated Gammaherpesvirus Replication in Apoptosis-Resistant B Lymphoma Cells.
Immune-related gene expression in the kidneys and spleens of goslings infected with goose nephritic astrovirus.
HSV-1-induced SOCS-1 expression in keratinocytes: use of a SOCS-1 antagonist to block a novel mechanism of viral immune evasion.
Functional motifs responsible for human metapneumovirus M2-2-mediated innate immune evasion.
3C Protease of Enterovirus D68 Inhibits Cellular Defense Mediated by Interferon Regulatory Factor 7.
Marburg virus evades interferon responses by a mechanism distinct from ebola virus.
Negative Regulatory Role of the Spring Viremia of Carp Virus Matrix Protein in the Host Interferon Response by Targeting the MAVS/TRAF3 Signaling Axis.
Interferon Regulatory Factor 3-Mediated Signaling Limits Middle-East Respiratory Syndrome (MERS) Coronavirus Propagation in Cells from an Insectivorous Bat.
Unique resistance of I/LnJ mice to a retrovirus is due to sustained interferon gamma-dependent production of virus-neutralizing antibodies.

Epigenetic silencing of IRF1 dysregulates type III interferon responses to respiratory virus infection in epithelial to mesenchymal transition.
Infectious hematopoietic necrosis virus N protein suppresses fish IFN1 production by targeting the MITA.
The Ambystoma tigrinum virus (ATV) RNase III gene can modulate host PKR activation and interferon production.
Amino Acid Polymorphisms on the Brazilian Strain of Yellow Fever Virus Methyltransferase Are Related to the Host's Immune Evasion Mediated by Type I Interferon.
TRAF3IP3 Is Cleaved by EV71 3C Protease and Exhibits Antiviral Activity.
Singapore grouper iridovirus infection counteracts poly I:C induced antiviral immune response in vitro.
Structural and Functional Characterization of the Phosphoprotein Central Domain of Spring Viremia of Carp Virus.
DDX56 cooperates with FMDV 3A to enhance FMDV replication by inhibiting the phosphorylation of IRF3.
Fifty Shades of E(rns): Innate Immune Evasion by the Viral Endonucleases of All Pestivirus Species.
IFN $\beta$ -dependent increases in STAT1, STAT2, and IRF9 mediate resistance to viruses and DNA damage.
Enterovirus 71 3C inhibits cytokine expression through cleavage of the TAK1/TAB1/TAB2/TAB3 complex.
Swine acute diarrhoea syndrome coronavirus (SADS-CoV) Nsp5 antagonizes type I interferon signaling by cleaving DCP1A.
Amelioration of Beta Interferon Inhibition by NS4B Contributes to Attenuating Tembusu Virus Virulence in Ducks.
PCR array profiling of antiviral genes in human embryonic kidney cells expressing human coronavirus OC43 structural and accessory proteins.
Molecular characterization, transcriptional regulation of sea perch Moloney leukemia virus 10 and its antiviral function against VHSV.
The plasmacytoid dendritic cell as the Swiss army knife of the immune system: molecular regulation of its multifaceted functions.
Immunity to fish rhabdoviruses.
Viral and metazoan poxins are cGAMP-specific nucleases that restrict cGAS-STING signalling.
Human metapneumovirus M2-2 protein inhibits innate immune response in monocyte-derived dendritic cells.
Structures of diverse poxin cGAMP nucleases reveal a widespread role for cGAS-STING evasion in host-pathogen conflict.
Staying on message: design principles for controlling nonspecific responses to siRNA.
Enterovirus 71 antagonizes the antiviral activity of host STAT3 and IL-6R with partial dependence on virus-induced miR-124.
Suppression of immunostimulatory siRNA-driven innate immune activation by 2'-modified RNAs.
Abortigenic but Not Neurotropic Equine Herpes Virus 1 Modulates the Interferon Antiviral Defense.
Positively Charged Amino Acids in the Pestiviral E(rns) Control Cell Entry, Endoribonuclease Activity and Innate Immune Evasion.
Innate immune evasion by alphaviruses.
Structural and Biological Basis of Alphacoronavirus nsp1 Associated with Host Proliferation and Immune Evasion.
Toll-like receptor 9 in plasmacytoid dendritic cells fails to detect parvoviruses.
Tyrosine 110 in the measles virus phosphoprotein is required to block STAT1 phosphorylation.



Backbone and Ile- $\delta$ 1, Leu, Val methyl 1H, 13C and 15N NMR chemical shift assignments for human interferon-stimulated gene 15 protein.
Simian varicella virus inhibits the interferon gamma signalling pathway.
Tim-3/Galectin-9 signaling pathway is involved in the cytokine changes in mice with alveolar echinococcosis.
Experimental Evidence of Intrinsic Disorder and Amyloid Formation by the Henipavirus W Proteins.
HIV-1 transactivator protein induction of suppressor of cytokine signaling-2 contributes to dysregulation of IFN[gamma] signaling.
Viral tegument proteins restrict cGAS-DNA phase separation to mediate immune evasion.
Tousled-Like Kinases Suppress Innate Immune Signaling Triggered by Alternative Lengthening of Telomeres.
Virology: Poxins Soothe the STING.
Convergent mutations in phage virion assembly proteins enable evasion of Type I CBASS immunity.
Foot-and-Mouth Disease Virus Inhibits RIP2 Protein Expression to Promote Viral Replication.
Tuft-cell-intrinsic and -extrinsic mediators of norovirus tropism regulate viral immunity.
Essential roles of RNA cap-proximal ribose methylation in mammalian embryonic development and fertility.
Peroxisomes are platforms for cytomegalovirus' evasion from the cellular immune response.
MicroRNA-302a suppresses influenza A virus-stimulated interferon regulatory factor-5 expression and cytokine storm induction.
Comparative proteome analyses of host protein expression in response to Enterovirus 71 and Coxsackievirus A16 infections.
Foot-and-Mouth Disease Virus Induces Porcine Gasdermin E-Mediated Pyroptosis through the Protease Activity of 3C(pro).
pol-miR-194a of Japanese flounder ( <i>Paralichthys olivaceus</i> ) suppresses type I interferon response and facilitates <i>Edwardsiella tarda</i> infection.
Induction of Immune Responses and Immune Evasion by Human Bocavirus.
Identification and characterization of bovine programmed death-ligand 2.
The trypanosomatid-specific N terminus of RPA2 is required for RNA polymerase I assembly, localization, and function.
Infection and innate immune mechanism of goose astrovirus.

Table A2-50, Cluster 49

Cluster 49 focuses on the tumor microenvironment (especially the role of tumor-associated macrophages in promoting immunosuppression), emphasizing the targeting of these tumor-associated macrophages for cancer immunotherapy (435)
The Role of Antigen Presentation in Tumor-Associated Macrophages.
The role of macrophages in the tumor microenvironment and tumor metabolism.
Tumor-associated macrophages in cancer: recent advancements in cancer nanoimmunotherapies.
Tumor-Associated Macrophages as Target for Antitumor Therapy.
Metabolism and polarization regulation of macrophages in the tumor microenvironment.
Exosome-mediated communication between tumor cells and tumor-associated macrophages: implications for tumor microenvironment.
Metabolic reprogramming of tumor-associated macrophages.
The role of tumor-associated macrophages in breast cancer progression (review).

Tumor-Associated Macrophages: Protumoral Macrophages in Inflammatory Tumor Microenvironment.
The cross-talk between tumor-associated macrophages and tumor endothelium: Recent advances in macrophage-based cancer immunotherapy.
Targeting tumor-associated macrophages in the tumor microenvironment.
CAFs Interacting With TAMs in Tumor Microenvironment to Enhance Tumorigenesis and Immune Evasion.
Tumor-associated macrophages: implications in cancer immunotherapy.
Targeting Tumor-Associated Macrophages in Cancer Immunotherapy.
Metabolic Reprogramming Induces Macrophage Polarization in the Tumor Microenvironment.
Role of macrophages in cancer progression and targeted immunotherapies.
Tumor promoting role of anti-tumor macrophages in tumor microenvironment.
Research progress of tumor microenvironment and tumor-associated macrophages.
A role for platelets in metabolic reprogramming of tumor-associated macrophages.
Tumor-associated macrophages employ immunoediting mechanisms in colorectal tumor progression: Current research in Macrophage repolarization immunotherapy.
Immune checkpoint inhibitors as mediators for immunosuppression by cancer-associated fibroblasts: A comprehensive review.
Tumor-associated macrophages as targets for tumor immunotherapy.
Emerging Nanoparticle Strategies for Modulating Tumor-Associated Macrophage Polarization. [Prevention and treatment of lung cancer by regulating tumor-associated macrophages with traditional Chinese medicine].
The Unique Molecular and Cellular Microenvironment of Ovarian Cancer.
Dual Relationship Between Stromal Cells and Immune Cells in the Tumor Microenvironment.
Shedding light on macrophage immunotherapy in lung cancer.
Macrophage responses to hypoxia: implications for tumor progression and anti-cancer therapies.
The impact of hypoxia on tumor-associated macrophages.
Targeting Macrophages for Tumor Therapy.
Tumor-Associated Macrophages in Multiple Myeloma: Key Role in Disease Biology and Potential Therapeutic Implications.
Preclinical and Clinical Therapeutic Strategies Affecting Tumor-Associated Macrophages in Hepatocellular Carcinoma.
Tumor-associated macrophages contribute to tumor progression in ovarian cancer.
Macrophage and Tumor Cell Cross-Talk Is Fundamental for Lung Tumor Progression: We Need to Talk.
Tumor Associated Macrophages in Kidney Cancer.
Tumor-associated macrophages: effectors of angiogenesis and tumor progression.
Tumor Hypoxia: A Key Determinant of Microenvironment Hostility and a Major Checkpoint during the Antitumor Response.
Cancer-Associated Fibroblasts in the Hypoxic Tumor Microenvironment.
Monocyte/macrophage infiltration in tumors: modulators of angiogenesis.
Mechanisms of immune modulation in the tumor microenvironment and implications for targeted therapy.
Targeted Delivery of Zoledronate to Tumor-Associated Macrophages for Cancer Immunotherapy.
The role of tumor-associated macrophages in primary hepatocellular carcinoma and its related targeting therapy.

The Antitumor Cytotoxic Response: If the Killer Cells Play the Music, the Microenvironmental Hypoxia Plays the Tune.
Crosstalk between cancer-associated fibroblasts and immune cells in the tumor microenvironment: new findings and future perspectives.
Hepatocellular Carcinoma Tumor Microenvironment and Its Implications in Terms of Anti-tumor Immunity: Future Perspectives for New Therapeutics.
Perspectives on Epigenetics and Cancer Immunotherapy: A Preface to Special Issue.
Evaluating the Polarization of Tumor-Associated Macrophages Into M1 and M2 Phenotypes in Human Cancer Tissue: Technicalities and Challenges in Routine Clinical Practice.
Reprogramming Hypoxic Tumor-Associated Macrophages by Nanoglycoclusters for Boosted Cancer Immunotherapy.
Cancer-Associated Fibroblasts in Conversation with Tumor Cells in Endometrial Cancers: A Partner in Crime.
Immunosuppressive circuits in tumor microenvironment and their influence on cancer treatment efficacy.
Immune Regulatory Processes of the Tumor Microenvironment under Malignant Conditions.
Current Strategies for Modulating Tumor-Associated Macrophages with Biomaterials in Hepatocellular Carcinoma.
Hypoxia, a key factor in the immune microenvironment.
Differentiation and gene expression profile of tumor-associated macrophages.
The Effect of Hypoxia and Hypoxia-Associated Pathways in the Regulation of Antitumor Response: Friends or Foes?
Prediction of immunotherapy efficacy and immunomodulatory role of hypoxia in colorectal cancer.
Tumor Hypoxia Regulates Immune Escape/Invasion: Influence on Angiogenesis and Potential Impact of Hypoxic Biomarkers on Cancer Therapies.
The regulation of immune checkpoints by the hypoxic tumor microenvironment.
Latest Advances in Targeting the Tumor Microenvironment for Tumor Suppression.
Cancer-Associated Fibroblast Heterogeneity: A Factor That Cannot Be Ignored in Immune Microenvironment Remodeling.
Stromal cells in the tumor microenvironment: accomplices of tumor progression?
Microenvironmental hypoxia orchestrating the cell stroma cross talk, tumor progression and antitumor response.
Nitric oxide, apoptosis and macrophage polarization during tumor progression.
Hypoxic stress: obstacles and opportunities for innovative immunotherapy of cancer.
Development of Immunotherapy Strategies Targeting Tumor Microenvironment Is Fiercely Ongoing.
Macrophages and NF- $\kappa$ B in cancer.
Tumor Microenvironment and Immunotherapy Response in Head and Neck Cancer.
Hijacked Immune Cells in the Tumor Microenvironment: Molecular Mechanisms of Immunosuppression and Cues to Improve T Cell-Based Immunotherapy of Solid Tumors.
PGE2-HIF1 $\alpha$ reciprocal induction regulates migration, phenotypic alteration and immunosuppressive capacity of macrophages in tumor microenvironment.
Biological Effects of Nanoparticles on Macrophage Polarization in the Tumor Microenvironment.
The Dark Side of Fibroblasts: Cancer-Associated Fibroblasts as Mediators of Immunosuppression in the Tumor Microenvironment.
"Re-educating" Tumor Associated Macrophages as a Novel Immunotherapy Strategy for Neuroblastoma.
The interactions of docetaxel with tumor microenvironment.

Understanding Tricky Cellular and Molecular Interactions in Pancreatic Tumor Microenvironment: New Food for Thought.
Targeting the tumor microenvironment by immunotherapy: part 2.
Intratumoral Platelets: Harmful or Incidental Bystanders of the Tumor Microenvironment?
Unusual Association of NF- $\kappa$ B Components in Tumor-Associated Macrophages (TAMs) Promotes HSPG2-Mediated Immune-Escaping Mechanism in Breast Cancer.
Multifunctional biomimetic nanoparticles loading baicalin for polarizing tumor-associated macrophages.
Next frontier in tumor immunotherapy: macrophage-mediated immune evasion.
Macrophages: Key orchestrators of a tumor microenvironment defined by therapeutic resistance.
Activation of phagocytosis by immune checkpoint blockade.
The evolving landscape of N(6)-methyladenosine modification in the tumor microenvironment.
Brain Tumor Microenvironment and Host State: Implications for Immunotherapy.
Crosstalk between IL-15R $\alpha$ (+) tumor-associated macrophages and breast cancer cells reduces CD8(+) T cell recruitment.
The Role of PRMT5 in Immuno-Oncology.
Integrating tumor hypoxic stress in novel and more adaptable strategies for cancer immunotherapy.
Tumor-macrophage crosstalk: how to listen.
Hypoxia and the Tumor Microenvironment.
A Novel Selective Axl/Mer/CSF1R Kinase Inhibitor as a Cancer Immunotherapeutic Agent Targeting Both Immune and Tumor Cells in the Tumor Microenvironment.
Tumor-Derived Retinoic Acid Regulates Intratumoral Monocyte Differentiation to Promote Immune Suppression.
Supramolecular Nanotherapeutics for Macrophage Immunotherapy.
Unraveling the Immune Microenvironment in Classic Hodgkin Lymphoma: Prognostic and Therapeutic Implications.
Immune cells within the tumor microenvironment: Biological functions and roles in cancer immunotherapy.
Targeting interleukin 4 receptor alpha on tumor-associated macrophages reduces the pro-tumor macrophage phenotype.
Aggressiveness Niche: Can It Be the Foster Ground for Cancer Metastasis Precursors?
Reactive oxygen species in the tumor niche triggers altered activation of macrophages and immunosuppression: Role of fluoxetine.
Association between pH regulation of the tumor microenvironment and immunological state.
Specialized Pro-Resolving Mediators Mitigate Cancer-Related Inflammation: Role of Tumor-Associated Macrophages and Therapeutic Opportunities.
M1 Macrophage-Derived Nanovesicles Potentiate the Anticancer Efficacy of Immune Checkpoint Inhibitors.
Imitating Hypoxia and Tumor Microenvironment with Immune Evasion by Employing Three Dimensional In vitro Cellular Models: Impressive Tool in Drug Discovery.
Low doses of gamma irradiation potentially modifies immunosuppressive tumor microenvironment by retuning tumor-associated macrophages: lesson from insulinoma.
Attenuated Listeria monocytogenes reprograms M2-polarized tumor-associated macrophages in ovarian cancer leading to iNOS-mediated tumor cell lysis.
Understanding the Role of Adipocytes and Fibroblasts in Cancer.
Immunomodulatory nanomedicine for colorectal cancer treatment: a landscape to be explored?

The balance between breast cancer and the immune system: Challenges for prognosis and clinical benefit from immunotherapies.
Aspects of the Tumor Microenvironment Involved in Immune Resistance and Drug Resistance.
Hypoxia: a key player in antitumor immune response. A Review in the Theme: Cellular Responses to Hypoxia.
Epigenetic regulation and therapeutic targets in the tumor microenvironment.
Targeting Hypoxia-Driven Metabolic Reprogramming to Constrain Tumor Progression and Metastasis.
Cancer-Associated Fibroblasts: Accomplices in the Tumor Immune Evasion.
Targeting Tumor-Associated Macrophages in the Pediatric Sarcoma Tumor Microenvironment.
IL-6 and leukemia-inhibitory factor are involved in the generation of tumor-associated macrophage: regulation by IFN- $\gamma$ .
Hypoxia-inducible factors: master regulators of hypoxic tumor immune escape.
Dual-targeting tumor cells and tumor associated macrophages with lipid coated calcium zoledronate for enhanced lung cancer chemoimmunotherapy.
The Crosstalk Between Tumor Cells and the Immune Microenvironment in Breast Cancer: Implications for Immunotherapy.
Hypoxia-Regulated Tumor-Derived Exosomes and Tumor Progression: A Focus on Immune Evasion.
CXCL9-expressing tumor-associated macrophages: new players in the fight against cancer.
Membrane Cholesterol Efflux Drives Tumor-Associated Macrophage Reprogramming and Tumor Progression.
Crosstalk among m(6)A RNA methylation, hypoxia and metabolic reprogramming in TME: from immunosuppressive microenvironment to clinical application.
Combinatorial therapy in tumor microenvironment: Where do we stand?
Strategy of targeting the tumor microenvironment via inhibition of fibroblast/fibrosis remodeling new era to cancer chemo-immunotherapy resistance.
Chronic inflammation, cancer development and immunotherapy.
Tumor-associated macrophages expressing galectin-9 identify immunoevasive subtype muscle-invasive bladder cancer with poor prognosis but favorable adjuvant chemotherapeutic response.
CCL2-CCR2 axis recruits tumor associated macrophages to induce immune evasion through PD-1 signaling in esophageal carcinogenesis.
Tumor-associated macrophages: unwitting accomplices in breast cancer malignancy.
Tumor-associated macrophages mediate immunosuppression in the renal cancer microenvironment by activating the 15-lipoxygenase-2 pathway.
TGF- $\beta$ Mediated Immune Evasion in Cancer-Spotlight on Cancer-Associated Fibroblasts.
Nanoparticle Delivery of MnO(2) and Antiangiogenic Therapy to Overcome Hypoxia-Driven Tumor Escape and Suppress Hepatocellular Carcinoma.
Remodeling the Tumor Myeloid Landscape to Enhance Antitumor Antibody Immunotherapies.
The exploitation of enzyme-based cancer immunotherapy.
GADD45 $\beta$ Loss Ablates Innate Immunosuppression in Cancer.
Allies or Enemies-The Multifaceted Role of Myeloid Cells in the Tumor Microenvironment.
The Role of Amino Acid Metabolism of Tumor Associated Macrophages in the Development of Colorectal Cancer.
Liver Tumor Microenvironment.
Turning Cold into Hot: Firing up the Tumor Microenvironment.
Alteration of the Antitumor Immune Response by Cancer-Associated Fibroblasts.

Targeting the Tumor Microenvironment in Acute Myeloid Leukemia: The Future of Immunotherapy and Natural Products.
Cancer-associated fibroblasts: an emerging target of anti-cancer immunotherapy.
MicroRNA regulation in cancer-associated fibroblasts.
Evaluation of Macrophage Polarization in Pancreatic Cancer Microenvironment Under Hypoxia.
The role of matrix stiffness in cancer stromal cell fate and targeting therapeutic strategies.
Targeting extracellular matrix through phytochemicals: a promising approach of multi-step actions on the treatment and prevention of cancer.
Lipid-loaded macrophages as new therapeutic target in cancer.
Targeting cancer glycosylation repolarizes tumor-associated macrophages allowing effective immune checkpoint blockade.
Targeting tumor-infiltrating macrophages to combat cancer.
Hypoxia promotes tumor growth in linking angiogenesis to immune escape.
The impact of hypoxia on tumor-mediated bypassing anti-PD-(L)1 therapy.
ALKBH5 Facilitates Hypoxia-Induced Paraspeckle Assembly and IL8 Secretion to Generate an Immunosuppressive Tumor Microenvironment.
Tumor Cell-derived Extracellular Vesicles in Modulating Phenotypes and Immune Functions of Macrophages: Mechanisms and Therapeutic Applications.
Cancer-associated fibroblasts in non-small cell lung cancer: Recent advances and future perspectives.
Tumor-associated leukemia inhibitory factor and IL-6 skew monocyte differentiation into tumor-associated macrophage-like cells.
Cellular and Molecular Players in the Tumor Microenvironment of Renal Cell Carcinoma.
CD47 Blockade Inhibits Tumor Progression through Promoting Phagocytosis of Tumor Cells by M2 Polarized Macrophages in Endometrial Cancer.
Targeting the Tumor Microenvironment in Brain Metastasis.
Co-opting macrophage traits in cancer progression: a consequence of tumor cell fusion?
miR-210 and hypoxic microvesicles: Two critical components of hypoxia involved in the regulation of killer cells function.
Tumor Extracellular Matrix Stiffness Promptly Modulates the Phenotype and Gene Expression of Infiltrating T Lymphocytes.
The Bladder Tumor Microenvironment Components That Modulate the Tumor and Impact Therapy.
Tumor-Associated Macrophage-Derived Exosomal LINC01232 Induces the Immune Escape in Glioma by Decreasing Surface MHC-I Expression.
Biomimetic Nanoparticles Carrying a Repolarization Agent of Tumor-Associated Macrophages for Remodeling of the Inflammatory Microenvironment Following Photothermal Therapy.
Ferritin light chain promotes the reprogramming of glioma immune microenvironment and facilitates glioma progression.
Hypoxic Stress-Induced Tumor and Immune Plasticity, Suppression, and Impact on Tumor Heterogeneity.
Integrated analysis of the tumor microenvironment using a reconfigurable microfluidic cell culture platform.
Method for the Analysis of the Tumor Microenvironment by Mass Cytometry: Application to Chronic Lymphocytic Leukemia.
Current Advancements of Plant-Derived Agents for Triple-Negative Breast Cancer Therapy through Deregulating Cancer Cell Functions and Reprogramming Tumor Microenvironment.
VentX expression in tumor-associated macrophages promotes phagocytosis and immunity against pancreatic cancers.

Tumor microenvironment (TME)-driven immune suppression in B cell malignancy.
Tumor microenvironment manipulates chemoresistance in ovarian cancer (Review).
Dual role of macrophage in tumor immunity.
Cancer-Associated Fibroblasts Suppress CD8+ T-cell Infiltration and Confer Resistance to Immune-Checkpoint Blockade.
Immunobiology of cancer-associated fibroblasts in the context of radiotherapy.
Arsenic trioxide elicits anti-tumor activity by inhibiting polarization of M2-like tumor-associated macrophages via Notch signaling pathway in lung adenocarcinoma.
RNAi screen in apoptotic cancer cell-stimulated human macrophages reveals co-regulation of IL-6/IL-10 expression.
Role of reactive oxygen species in tumors based on the 'seed and soil' theory: A complex interaction (Review).
Activation of MAT2A-RIP1 signaling axis reprograms monocytes in gastric cancer.
Cancer-associated fibroblasts: tumor defenders in radiation therapy.
Mechanisms of hypoxia-mediated immune escape in cancer.
Dendritic cell-based cancer immunotherapy in the era of immune checkpoint inhibitors: From bench to bedside.
Vascular Microenvironment, Tumor Immunity and Immunotherapy.
A feed-forward loop based on aerobic glycolysis and TGF- $\beta$ between tumor-associated macrophages and bladder cancer cells promoted malignant progression and immune escape.
Perspectives for 3D-Bioprinting in Modeling of Tumor Immune Evasion.
Tumor-associated macrophages promote lung metastasis and induce epithelial-mesenchymal transition in osteosarcoma by activating the COX-2/STAT3 axis.
Nitric Oxide Generated by Tumor-Associated Macrophages Is Responsible for Cancer Resistance to Cisplatin and Correlated With Syntaxin 4 and Acid Sphingomyelinase Inhibition.
What Do We Have to Know about PD-L1 Expression in Prostate Cancer? A Systematic Literature Review. Part 3: PD-L1, Intracellular Signaling Pathways and Tumor Microenvironment.
The role of hypoxia-inducible factor 1 in tumor immune evasion.
Targeting the Tumor Microenvironment of Leukemia and Lymphoma.
Reprogramming the tumor microenvironment by genome editing for precision cancer therapy.
Decoding the Complexity of Immune-Cancer Cell Interactions: Empowering the Future of Cancer Immunotherapy.
Persistent immune response: Twice tumor exfoliation induced by sialic acid-modified vincristine sulfate liposomes.
A novel classifier combining G protein-coupled receptors and the tumor microenvironment is associated with survival status in glioblastoma.
Fatty Acid-Driven Polarization of Suppressive Bone Marrow-Derived Macrophages Including Metabolic and Functional Analysis.
Supraglottic laryngeal tumor microenvironmental factors facilitate STAT3 dependent pro-tumorigenic switch in tumor associated macrophages to render utmost immune evasion.
Resveratrol inhibits lung cancer growth by suppressing M2-like polarization of tumor associated macrophages.
Unveiling the Role of the Tumor Microenvironment in the Treatment of Follicular Lymphoma.
CCL18 in the Progression of Cancer.
Hypoxia and the Tumor Secretome.
Theranostic Radiopharmaceuticals Targeting Cancer-Associated Fibroblasts.

Mapping the tumor microenvironment in clear cell renal carcinoma by single-cell transcriptome analysis.
Cancer-associated fibroblasts and their influence on tumor immunity and immunotherapy.
The Vault Complex Is Significantly Involved in Therapeutic Responsiveness of Endocrine Tumors and Linked to Autophagy under Chemotherapeutic Conditions.
Acute vs. Chronic vs. Cyclic Hypoxia: Their Differential Dynamics, Molecular Mechanisms, and Effects on Tumor Progression.
Targeting Wnt/ $\beta$ -catenin signaling using XAV939 nanoparticles in tumor microenvironment-conditioned macrophages promote immunogenicity.
Poor clinical outcome in metastatic melanoma is associated with a microRNA-modulated immunosuppressive tumor microenvironment.
Role and mechanism of programmed death-ligand 1 in hypoxia-induced liver cancer immune escape.
Pancreatic cancer cell/fibroblast co-culture induces M2 like macrophages that influence therapeutic response in a 3D model.
Immune checkpoints in the tumor microenvironment.
Mutual concessions and compromises between stromal cells and cancer cells: driving tumor development and drug resistance.
Tumors may modulate host immunity partly through hypoxia-induced sympathetic bias.
Tumor microenvironment-related dendritic cell deficiency: a target to enhance tumor immunotherapy.
Assessment of Stabilization and Activity of the HIFs Important for Hypoxia-Induced Signalling in Cancer Cells.
Macrophage polarity in cancer: A review.
Influence of Estrogen on the NSCLC Microenvironment: A Comprehensive Picture and Clinical Implications.
CSF1R inhibition delays cervical and mammary tumor growth in murine models by attenuating the turnover of tumor-associated macrophages and enhancing infiltration by CD8(+) T cells.
MicroRNA-A Tumor Trojan Horse for Tumor-Associated Macrophages.
Stromal Protein-Mediated Immune Regulation in Digestive Cancers.
Hypoxia-driven intratumor heterogeneity and immune evasion.
Local Activation of p53 in the Tumor Microenvironment Overcomes Immune Suppression and Enhances Antitumor Immunity.
Fibroblasts Fuel Immune Escape in the Tumor Microenvironment.
Blood monocytes and tumor-associated macrophages in human cancer: differences in activation levels.
Fibrosis and cancer: A strained relationship.
Peritoneal resident macrophages in tumor metastasis and immunotherapy.
Immuno-oncological Efficacy of RXDX-106, a Novel TAM (TYRO3, AXL, MER) Family Small-Molecule Kinase Inhibitor.
Spi-B Promotes the Recruitment of Tumor-Associated Macrophages via Enhancing CCL4 Expression in Lung Cancer.
Tumor-associated macrophage-specific CD155 contributes to M2-phenotype transition, immunosuppression, and tumor progression in colorectal cancer.
Immunosuppressive Tumor Microenvironment and Immunotherapy of Epstein-Barr Virus-Associated Malignancies.
Role of Hypoxic Stress in Regulating Tumor Immunogenicity, Resistance and Plasticity.



Tumor microenvironment and cellular senescence: Understanding therapeutic resistance and harnessing strategies.
Targeting stromal cell sialylation reverses T cell-mediated immunosuppression in the tumor microenvironment.
Immune microenvironment of cervical cancer and the role of IL-2 in tumor promotion.
Deletion of the RNA regulator HuR in tumor-associated microglia and macrophages stimulates anti-tumor immunity and attenuates glioma growth.
Vy9V62 T-cell immunotherapy in blood cancers: ready for prime time?
IL-15 protects NKT cells from inhibition by tumor-associated macrophages and enhances antimetastatic activity.
Hypoxia-Driven Immune Escape in the Tumor Microenvironment.
Strengthening the AntiTumor NK Cell Function for the Treatment of Ovarian Cancer.
Extracellular Matrix Protein Tenascin C Increases Phagocytosis Mediated by CD47 Loss of Function in Glioblastoma.
Hypoxia Mediates Tumor Malignancy and Therapy Resistance.
Biodegradable NIR-II Pseudo Conjugate Polymeric Nanoparticles Amplify Photodynamic Immunotherapy via Alleviation of Tumor Hypoxia and Tumor-Associated Macrophage Reprogramming.
Targeting tumor microenvironment in ovarian cancer: Premise and promise.
Immune Escape Mechanisms and Future Prospects for Immunotherapy in Neuroblastoma.
Hypoxia-inducible factors in hepatocellular carcinoma (Review).
Hypoxia-Inducible Factors: Master Regulators of Cancer Progression.
Recent progress of autophagy signaling in tumor microenvironment and its targeting for possible cancer therapeutics.
Blockade of DC-SIGN(+) Tumor-Associated Macrophages Reactivates Antitumor Immunity and Improves Immunotherapy in Muscle-Invasive Bladder Cancer.
Bridging angiogenesis and immune evasion in the hypoxic tumor microenvironment.
Mechanisms of Macrophage Plasticity in the Tumor Environment: Manipulating Activation State to Improve Outcomes.
Hypoxia induces escape from innate immunity in cancer cells via increased expression of ADAM10: role of nitric oxide.
The hypoxic tumor microenvironment: driving the tumorigenesis of non-small-cell lung cancer.
Single-cell RNA sequencing reveals intra-tumoral heterogeneity of glioblastoma and a pro-tumor subset of tumor-associated macrophages characterized by EZH2 overexpression.
Tumor-Derived Extracellular Vesicles in Cancer Immunoediting and Their Potential as Oncoimmunotherapeutics.
Targeting UDP- $\alpha$ -d-glucose 6-dehydrogenase alters the CNS tumor immune microenvironment and inhibits glioblastoma growth.
Immune cellular components and signaling pathways in the tumor microenvironment.
Poor clinical outcomes and immunoevasive contexture in SIRP $\alpha$ (+) tumor-associated macrophages enriched muscle-invasive bladder cancer patients.
Clinical relevance of the tumor microenvironment and immune escape of oral squamous cell carcinoma.
Unveiling Tumor Microenvironment Interactions Using Zebrafish Models.
Macrophage-Mediated Tumor Cell Phagocytosis: Opportunity for Nanomedicine Intervention.
The Cancer Cell Dissemination Machinery as an Immunosuppressive Niche: A New Obstacle Towards the Era of Cancer Immunotherapy.

TLR4 expression and functionality are downregulated in glioblastoma cells and in tumor-associated macrophages: A new mechanism of immune evasion?
Glioblastoma Chemoresistance: The Double Play by Microenvironment and Blood-Brain Barrier.
Hypoxia-/HIF-1 $\alpha$ -Driven Factors of the Tumor Microenvironment Impeding Antitumor Immune Responses and Promoting Malignant Progression.
Targeting STAT3 phosphorylation by neem leaf glycoprotein prevents immune evasion exerted by supraglottic laryngeal tumor induced M2 macrophages.
Extracellular vesicle-mediated crosstalk between melanoma and the immune system: Impact on tumor progression and therapy response.
In Situ Vaccination as a Strategy to Modulate the Immune Microenvironment of Hepatocellular Carcinoma.
TAM receptor tyrosine kinases as emerging targets of innate immune checkpoint blockade for cancer therapy.
The tumor microenvironment shapes hallmarks of mature B-cell malignancies.
The Microenvironment in Epstein-Barr Virus-Associated Malignancies.
Role of hypoxia in inhibiting dendritic cells by VEGF signaling in tumor microenvironments: mechanism and application.
Heme catabolism by tumor-associated macrophages controls metastasis formation.
The intriguing roles of Siglec family members in the tumor microenvironment.
Tumor innervation: peripheral nerves take control of the tumor microenvironment.
Autophagy and ncRNAs: Dangerous Liaisons in the Crosstalk between the Tumor and Its Microenvironment.
Reversing insufficient photothermal therapy-induced tumor relapse and metastasis by regulating cancer-associated fibroblasts.
The pathological role of C-X-C chemokine receptor type 4 (CXCR4) in colorectal cancer (CRC) progression; special focus on molecular mechanisms and possible therapeutics.
The Emerging Role of the Single-Cell and Spatial Tumor Microenvironment in High-Grade Serous Ovarian Cancer.
Tumor-associated $\alpha 2$ vacuolar ATPase acts as a key mediator of cancer-related inflammation by inducing pro-tumorigenic properties in monocytes.
HGF/c-Met Signalling in the Tumor Microenvironment.
Activation of the VEGFC/VEGFR3 Pathway Induces Tumor Immune Escape in Colorectal Cancer.
Neuro-immune interactions and immuno-oncology.
Tumor-Infiltrating B Lymphocytes: Promising Immunotherapeutic Targets for Primary Liver Cancer Treatment.
How Cells Communicate with Each Other in the Tumor Microenvironment: Suggestions to Design Novel Therapeutic Strategies in Cancer Disease.
Hypoxia-Driven Oncometabolite L-2HG Maintains Stemness-Differentiation Balance and Facilitates Immune Evasion in Pancreatic Cancer.
Targeting CAMKII to reprogram tumor-associated macrophages and inhibit tumor cells for cancer immunotherapy with an injectable hybrid peptide hydrogel.
Comprehensive Pan-Cancer Analysis of Senescence With Cancer Prognosis and Immunotherapy.
CCL21 Programs Immune Activity in Tumor Microenvironment.
The immune escape mechanism of nasopharyngeal carcinoma.
Recent advances of nanotechnology-based tumor vessel-targeting strategies.
Metabolic reprogramming of tumor-associated macrophages by collagen turnover promotes fibrosis in pancreatic cancer.

Interplay of Viral Infection, Host Cell Factors and Tumor Microenvironment in the Pathogenesis of Nasopharyngeal Carcinoma.
Nanomedicine-Based Immunotherapy for the Treatment of Cancer Metastasis.
Immune evasion in esophageal squamous cell cancer: From the perspective of tumor microenvironment.
Novel PdPtCu Nanozymes for Reprogramming Tumor Microenvironment to Boost Immunotherapy Through Endoplasmic Reticulum Stress and Blocking IDO-Mediated Immune Escape.
Targeting the bicarbonate transporter SLC4A4 overcomes immunosuppression and immunotherapy resistance in pancreatic cancer.
Warburg Effect Is a Cancer Immune Evasion Mechanism Against Macrophage Immunosurveillance.
Wnt Signaling in the Tumor Microenvironment.
The Potential Role of the T2 Ribonucleases in TME-Based Cancer Therapy.
Ligand Activation of TAM Family Receptors-Implications for Tumor Biology and Therapeutic Response.
Targeting macrophage Syk enhances responses to immune checkpoint blockade and radiotherapy in high-risk neuroblastoma.
Microenvironmental Regulation of Tumor Progression and Therapeutic Response in Brain Metastasis.
Hypoxia and Extracellular Acidification as Drivers of Melanoma Progression and Drug Resistance.
New insights into the Immune TME of adult-type diffuse gliomas.
Exploiting E3 ubiquitin ligases to reeducate the tumor microenvironment for cancer therapy.
Functions of Key Enzymes of Glycolytic Metabolism in Tumor Microenvironment.
Addressing activation of WNT beta-catenin pathway in diverse landscape of endometrial carcinogenesis.
Single-Cell Sequencing of Malignant Ascites Reveals Transcriptomic Remodeling of the Tumor Microenvironment during the Progression of Epithelial Ovarian Cancer.
Eph Receptors in the Immunosuppressive Tumor Microenvironment.
Metabolite activation of tumorigenic signaling pathways in the tumor microenvironment.
Tumor-induced reshuffling of lipid composition on the endoplasmic reticulum membrane sustains macrophage survival and pro-tumorigenic activity.
Axl and Mertk Receptors Cooperate to Promote Breast Cancer Progression by Combined Oncogenic Signaling and Evasion of Host Antitumor Immunity.
Tumor derived UBR5 promotes ovarian cancer growth and metastasis through inducing immunosuppressive macrophages.
What Are the Roles of Proprotein Convertases in the Immune Escape of Tumors?
Role of Hypoxia and the Adenosine System in Immune Evasion and Prognosis of Patients with Brain Metastases of Melanoma: A Multiplex Whole Slide Immunofluorescence Study.
Can Radiotherapy Empower the Host Immune System to Counterattack Neoplastic Cells? A Systematic Review on Tumor Microenvironment Radiomodulation.
Tumorous expression of NAC1 restrains antitumor immunity through the LDHA-mediated immune evasion.
Hypoxia induces immunogenic cell death of cancer cells by enhancing the exposure of cell surface calreticulin in an endoplasmic reticulum stress-dependent manner.
Therapeutic Potential of Targeting Stromal Crosstalk-Mediated Immune Suppression in Pancreatic Cancer.
Fibroblasts, an inconspicuous but essential player in colon cancer development and progression.
EHBP1L1 Drives Immune Evasion in Renal Cell Carcinoma through Binding and Stabilizing JAK1.
Pleiotropic effects of the COX-2/PGE2 axis in the glioblastoma tumor microenvironment.
Oral squamous carcinoma cells promote macrophage polarization in an MIF-dependent manner.

Immune escape: A critical hallmark in solid tumors.
Lewis Lung Cancer Cells Promote SIGNR1(CD209b)-Mediated Macrophages Polarization Induced by IL-4 to Facilitate Immune Evasion.
Immune Evasion in Pancreatic Cancer: From Mechanisms to Therapy.
From Whole Blood to Isolated Pro-Metastasis Immune Cells: An Ex Vivo Approach to Isolate and Manipulate Immune Cells Contributing to Tumor Metastasis.
Classification of primary liver cancer with immunosuppression mechanisms and correlation with genomic alterations.
Hyperglycemia Enhances Cancer Immune Evasion by Inducing Alternative Macrophage Polarization through Increased O-GlcNAcylation.
Matricellular proteins in intrahepatic cholangiocarcinoma.
The carcinoma-associated fibroblast expressing fibroblast activation protein and escape from immune surveillance.
MNK2 governs the macrophage antiinflammatory phenotype.
Tumor immunoevasion via acidosis-dependent induction of regulatory tumor-associated macrophages.
Collaborating with the enemy: function of macrophages in the development of neoplastic disease.
Tumor Cell-Intrinsic SETD2 Deficiency Reprograms Neutrophils to Foster Immune Escape in Pancreatic Tumorigenesis.
Blockade of fibroblast activation protein in combination with radiation treatment in murine models of pancreatic adenocarcinoma.
Insights into N6-methyladenosine (m6A) modification of noncoding RNA in tumor microenvironment.
Identification of CD300a as a new hypoxia-inducible gene and a regulator of CCL20 and VEGF production by human monocytes and macrophages.
Systemic Immune Bias Delineates Malignant Astrocytoma Survival Cohorts.
Tumor Acidosis and Hypoxia Differently Modulate the Inflammatory Program: Measurements In Vitro and In Vivo.
Dendritic cell reprogramming by the hypoxic environment.
Myeloid cell-synthesized coagulation factor X dampens antitumor immunity.
Cu(2+)-Pyropheophorbide a-Cystine Conjugate: Synergistic Photodynamic/Chemodynamic Therapy and Glutathione Depletion Improves the Antitumor Efficacy and Downregulates the Hypoxia-Inducing Factor.
Cutting Edge: Targeting Thrombocytes to Rewire Anticancer Immunity in the Tumor Microenvironment and Potentiate Efficacy of PD-1 Blockade.
Luteolin impairs hypoxia adaptation and progression in human breast and colon cancer cells.
Hypoxia-inducible factors and innate immunity in liver cancer.
Induction of systemic immune responses and reversion of immunosuppression in the tumor microenvironment by a therapeutic vaccine for cervical cancer.
Surface Modification of Macrophages with Nucleic Acid Aptamers for Enhancing the Immune Response against Tumor Cells.
An overlooked tumor promoting immunoregulation by non-hematopoietic stromal cells.
Hypoxic Melanoma Cells Deliver microRNAs to Dendritic Cells and Cytotoxic T Lymphocytes through Connexin-43 Channels.
Hybrid cellular membrane nanovesicles amplify macrophage immune responses against cancer recurrence and metastasis.
HIF-1 $\alpha$ signaling: Essential roles in tumorigenesis and implications in targeted therapies.

Cancer Cells Promote Immune Regulatory Function of Macrophages by Upregulating Scavenger Receptor MARCO Expression.
Immune and Inflammatory Cells of the Tumor Microenvironment Represent Novel Therapeutic Targets in Classical Hodgkin Lymphoma.
The Role of Tumor Microenvironment in Multiple Myeloma Development and Progression.
Role of ubiquitin specific proteases in the immune microenvironment of prostate cancer: A new direction.
Transient and Local Expression of Chemokine and Immune Checkpoint Traps To Treat Pancreatic Cancer.
The application of the fibroblast activation protein $\alpha$ -targeted immunotherapy strategy.
Tumor-host colluding through erythroid progenitor cells: Mechanisms and opportunities.
Identification of CD137- and CD137L-Expressing Cells in EL-4 Tumor.
Natural products as potent inhibitors of hypoxia-inducible factor-1 $\alpha$ in cancer therapy.
Lysophosphatidic Acid: Promoter of Cancer Progression and of Tumor Microenvironment Development. A Promising Target for Anticancer Therapies?
Roles of Podoplanin in Malignant Progression of Tumor.
Regulation of Tumor Immunity by Lysophosphatidic Acid.
New extracellular factors in glioblastoma multiforme development: neurotensin, growth differentiation factor-15, sphingosine-1-phosphate and cytomegalovirus infection.
Advances in research on immune escape mechanism of glioma.
Hypoxia accelerates the progression of angiosarcoma through the regulation of angiosarcoma cells and tumor microenvironment.
Pan-cancer transcriptomic analysis of CNS tumor stroma identifies a population of perivascular fibroblasts that predict poor immunotherapy response in glioblastoma patients.
Restoring the Immunity in the Tumor Microenvironment: Insights into Immunogenic Cell Death in Onco-Therapies.
Ferroptosis: A potential opportunity for intervention of pre-metastatic niche.
SRF617 Is a Potent Inhibitor of CD39 with Immunomodulatory and Antitumor Properties.
Neural regulations of the tumor microenvironment.
Biological Insight and Recent Advancement in the Treatment of Neuroblastoma.
DKK1 Promotes Tumor Immune Evasion and Impedes Anti-PD-1 Treatment by Inducing Immunosuppressive Macrophages in Gastric Cancer.
Immune Adaptation of Colorectal Cancer Stem Cells and Their Interaction With the Tumor Microenvironment.
Siglec Signaling in the Tumor Microenvironment.
TIMs, TAMs, and PS- antibody targeting: implications for cancer immunotherapy.
Dual Tumor Microenvironment Remodeling by Glucose-Contained Radical Copolymer for MRI-Guided Photoimmunotherapy.
Neem Leaf Glycoprotein in immunoregulation of cancer.
Hypoxia-inducible miR-210 regulates the susceptibility of tumor cells to lysis by cytotoxic T cells.
Hypoxic Roadmap of Glioblastoma-Learning about Directions and Distances in the Brain Tumor Environment.
Immune microenvironment characteristics in multiple myeloma progression from transcriptome profiling.
Hepatic carcinoma-associated fibroblasts induce IDO-producing regulatory dendritic cells through IL-6-mediated STAT3 activation.
Hypoxia-induced autophagy: a new player in cancer immunotherapy?

Hypoxia-inducible factors: cancer progression and clinical translation.
Combined effects of pericytes in the tumor microenvironment.
Genomic and phenotypic characterization of nurse-like cells that promote drug resistance in chronic lymphocytic leukemia.
DNA methylation repels binding of hypoxia-inducible transcription factors to maintain tumor immunotolerance.
Cutting edge: Hypoxia-induced Nanog favors the intratumoral infiltration of regulatory T cells and macrophages via direct regulation of TGF- $\beta$ 1.
Targeting micro-environmental pathways by PROTACs as a therapeutic strategy.
Targeting focal adhesion kinase renders pancreatic cancers responsive to checkpoint immunotherapy.
A Hypoxia-Sensitive Drug Delivery System Constructed by Nitroimidazole and its Application in the Treatment of Hepatocellular Carcinoma.
The emerging role of Arid5a in cancer: A new target for tumors.
Tryptophan 2,3-dioxygenase 2 controls M2 macrophages polarization to promote esophageal squamous cell carcinoma progression via AKT/GSK3 $\beta$ /IL-8 signaling pathway.
Effect of glucose level on chemical hypoxia- and hydrogen peroxide-induced chemokine expression in human glioblastoma cell lines.
Do Tumor Mechanical Stresses Promote Cancer Immune Escape?
Siglec-15 as an immune suppressor and potential target for normalization cancer immunotherapy.
Genomic testing, tumor microenvironment and targeted therapy of Hedgehog-related human cancers.
Metabolic, Anti-apoptotic and Immune Evasion Strategies of Primary Human Myeloma Cells Indicate Adaptations to Hypoxia.
Role of Immunosuppressive Microenvironment in Acquiring Immunotolerance Post-Photothermal Therapy.
Adenosinergic Signaling as a Key Modulator of the Glioma Microenvironment and Reactive Astrocytes.
Identification of a unique tumor cell subset employing myeloid transcriptional circuits to create an immunomodulatory microenvironment in glioblastoma.
The Interplay Between Innate Lymphoid Cells and the Tumor Microenvironment.
The immunosuppressive microenvironment and immunotherapy in human glioblastoma.
The prognostic and therapeutic implications of distinct patterns of argininosuccinate synthase 1 (ASS1) and arginase-2 (ARG2) expression by cancer cells and tumor stroma in non-small-cell lung cancer.
Mutant RAS and the tumor microenvironment as dual therapeutic targets for advanced colorectal cancer.
$\gamma\delta$ T Lymphocytes as a Double-Edged Sword-State of the Art in Gynecological Diseases.
Melanoma clonal subline analysis uncovers heterogeneity-driven immunotherapy resistance mechanisms.
TGF $\beta$ signaling activation correlates with immune-inflamed tumor microenvironment across human cancers and predicts response to immunotherapy.
The Tumor Microenvironment Impairs Th1 IFN $\gamma$ Secretion through Alternative Splicing Modifications of Irf1 Pre-mRNA.
Tumor-Infiltrating Immune Cell Landscapes in the Lymph Node Metastasis of Papillary Thyroid Cancer.
Activating Immune Recognition in Pancreatic Ductal Adenocarcinoma via Autophagy Inhibition, MEK Blockade, and CD40 Agonism.
Single tumor-initiating cells evade immune clearance by recruiting type II macrophages.

Hypoxia enhances IL-10-producing B cell generation through upregulating high-mobility group B1 on tumor cell-released autophagosomes.
Bioengineering the Oxygen-Deprived Tumor Microenvironment Within a Three-Dimensional Platform for Studying Tumor-Immune Interactions.
Role of hypoxia in cellular senescence.
Siah2 inhibitor and the metabolic antagonist Oxamate retard colon cancer progression and downregulate PD1 expression.
Hypoxia-inducible factor-1 (HIF-1): a potential target for intervention in ocular neovascular diseases.
Targeting CD47 in Anaplastic Thyroid Carcinoma Enhances Tumor Phagocytosis by Macrophages and Is a Promising Therapeutic Strategy.
CCND1 Amplification Contributes to Immunosuppression and Is Associated With a Poor Prognosis to Immune Checkpoint Inhibitors in Solid Tumors.
The Single-Cell Sequencing: A Dazzling Light Shining on the Dark Corner of Cancer.
Assessing the Immune Regulatory Role of Metabolites In Vitro.
The ectonucleotidases CD39 and CD73 on T cells: The new pillar of hematological malignancy.
Fenofibrate inhibits hypoxia-inducible factor-1 alpha and carbonic anhydrase expression through activation of AMP-activated protein kinase/HO-1/Sirt1 pathway in glioblastoma cells.
Phospholipase A(2) Drives Tumorigenesis and Cancer Aggressiveness through Its Interaction with Annexin A1.
Hypoxia-inducible factor-1: a possible link between inhalational anesthetics and tumor progression?
The IKZF1-IRF4/IRF5 Axis Controls Polarization of Myeloma-Associated Macrophages.
Methionine enkephalin (MENK) inhibits human gastric cancer through regulating tumor associated macrophages (TAMs) and PI3K/AKT/mTOR signaling pathway inside cancer cells.
Deoxynivalenol upregulates hypoxia-inducible factor-1 $\alpha$ to promote an "immune evasion" process by activating STAT3 signaling.
Tissue-Resident Macrophages Help Early-Stage Lung Tumors Develop.
Recent Advances in the Role of Arid5a in Immune Diseases and Cancer.
Hypoxia, oxidative stress, and immune evasion: a trinity of the trichothecenes T-2 toxin and deoxynivalenol (DON).
Xenoextracellular matrix-rosiglitazone complex-mediated immune evasion promotes xenogenic bioengineered root regeneration by altering M1/M2 macrophage polarization.
Altered Membrane Expression and Function of CD11b Play a Role in the Immunosuppressive Effects of Morphine on Macrophages at the Nanomolar Level.
HIF-1 $\alpha$ inhibits T-2 toxin-mediated "immune evasion" process by negatively regulating PD-1/PD-L1.
Tumor-Associated Fibroblasts and Microvessels Contribute to the Expression of Immunosuppressive Factor Indoleamine 2, 3-Dioxygenase in Human Esophageal Cancers.
Advances in Immune Microenvironment and Immunotherapy of Isocitrate Dehydrogenase Mutated Glioma.
MUC1-C integrates type II interferon and chromatin remodeling pathways in immunosuppression of prostate cancer.
The neurotoxicity of trichothecenes T-2 toxin and deoxynivalenol (DON): Current status and future perspectives.
Chitinase-3-like 1 protein complexes modulate macrophage-mediated immune suppression in glioblastoma.

Table A2-51, Cluster 50

Cluster 50 focuses on the role of the PD-1 and PD-L1 immune checkpoint in immune escape, emphasizing its cancer immunotherapy targeting to inhibit immune escape (421)
PD-L1 and PD-L2 differ in their molecular mechanisms of interaction with PD-1.
Tumor cell-intrinsic PD-1 receptor is a tumor suppressor and mediates resistance to PD-1 blockade therapy.
PD-1 and PD-L1 in cancer immunotherapy: clinical implications and future considerations.
[PD-1/PD-L1 Signaling Pathway and Its Anti-Tumor Effect in Lymphoma -Review].
Expression of Immunomodulatory Molecules PD-1, PD-L1, and PD-L2, and their Relationship With Clinicopathologic Characteristics in Endometrial Cancer.
[Targeting the PD-1/PD-L1 immune checkpoint signal - a new treatment strategy for cancer].
eEF2K promotes PD-L1 stabilization through inactivating GSK3 $\beta$ in melanoma.
Releasing the Brake on the Immune System: The PD-1 Strategy for Hematologic Malignancies.
Understanding Immune Evasion and Therapeutic Targeting Associated with PD-1/PD-L1 Pathway in Diffuse Large B-cell Lymphoma.
PD-1 and its ligands are important immune checkpoints in cancer.
PD-1/PD-L1 pathway: current researches in cancer.
Characterization of the Anti-PD-1 Antibody REGN2810 and Its Antitumor Activity in Human PD-1 Knock-In Mice.
PD-1/PD-L1 immune checkpoint: Potential target for cancer therapy.
Programmed death 1, ligand 1 and 2 correlated genes and their association with mutation, immune infiltration and clinical outcomes of hepatocellular carcinoma.
Prospects for targeting PD-1 and PD-L1 in various tumor types.
PD-1/PD-L1 blockade in cancer treatment: perspectives and issues.
Blocking tumor escape in hematologic malignancies: the anti-PD-1 strategy.
Intracellular accumulation of PD-1 molecules in circulating T lymphocytes in advanced malignant melanoma: an implication for immune evasion mechanism.
Correlation Between Programmed Death Receptor-1 Expression in Tumor-Infiltrating Lymphocytes and Programmed Death Ligand-1 Expression in Non-Small Cell Lung Carcinoma.
Progressive upregulation of PD-1 in primary and metastatic melanomas associated with blunted TCR signaling in infiltrating T lymphocytes.
Comprehensive insights into the effects and regulatory mechanisms of immune cells expressing programmed death-1/programmed death ligand 1 in solid tumors.
Anti-PD-1 And Anti-PD-L1 Antibodies as Immunotherapy Against Cancer: A Structural Perspective.
Antigen-Presenting Cell-Intrinsic PD-1 Neutralizes PD-L1 in cis to Attenuate PD-1 Signaling in T Cells.
The role of programmed cell death-1 (PD-1) and its ligands in pediatric cancer.
Basis of PD1/PD-L1 Therapies.
PD-1 blockade therapy in renal cell carcinoma: current studies and future promises.
Update on checkpoint blockade therapy for lymphoma.
PD-L1 Expression Correlates With Tumor Infiltrating Lymphocytes And Response To Neoadjuvant Chemotherapy In Cervical Cancer.
Development of a human phage display-derived anti-PD-1 scFv antibody: an attractive tool for immune checkpoint therapy.
Mathematical modeling for the combination treatment of IFN- $\gamma$ and anti-PD-1 in cancer immunotherapy.
Research progress of therapeutic effects and drug resistance of immunotherapy based on PD-1/PD-L1 blockade.



PD-1/PD-L1 Pathway and Its Blockade in Patients with Classic Hodgkin Lymphoma and Non-Hodgkin Large-Cell Lymphomas.
Regulation of PD-1 in T cells for cancer immunotherapy.
Programmed cell death-1 inhibition in lymphoma.
ERK and USP5 govern PD-1 homeostasis via deubiquitination to modulate tumor immunotherapy.
The design of high affinity human PD-1 mutants by using molecular dynamics simulations (MD).
Enhanced programmed death 1 (PD-1) and PD-1 ligand (PD-L1) expression in patients with actinic cheilitis and oral squamous cell carcinoma.
PD1/PD-L1 immune checkpoint as a potential target for preventing brain tumor progression.
Serum-derived exosomes promote CD8+ T cells to overexpress PD-1, affecting the prognosis of hypopharyngeal carcinoma.
Exosomal PD-L1 contributes to immunosuppression and is associated with anti-PD-1 response.
The Role of Anti-PD-1/PD-L1 Agents in Melanoma: Progress to Date.
Activation of the PD-1/PD-L1 immune checkpoint confers tumor cell chemoresistance associated with increased metastasis.
Structure-based small inhibitors search combined with molecular dynamics driven energies for human programmed cell death-1 (PD-1) protein.
Molecular dynamics of the immune checkpoint programmed cell death protein 1, PD-1: conformational changes of the BC-loop upon binding of the ligand PD-L1 and the monoclonal antibody nivolumab.
Current state of anti-PD-L1 and anti-PD-1 agents in cancer therapy.
Immunostaining of PD-1/PD-Ls in liver tissues of patients with hepatitis and hepatocellular carcinoma.
Signaling pathway and dysregulation of PD1 and its ligands in lymphoid malignancies.
Tumor Infiltrating Lymphocyte Expression of PD-1 Predicts Response to Anti-PD-1/PD-L1 Immunotherapy.
KLHL22 maintains PD-1 homeostasis and prevents excessive T cell suppression.
Role of PD-1 in Immunity and Diseases.
Immunohistochemical Analysis of PD-L1 Expression in Canine Malignant Cancers and PD-1 Expression on Lymphocytes in Canine Oral Melanoma.
Pneumonitis in cancer patients receiving anti-PD-1 and radiotherapies: Three case reports.
Tumor Suppression by PD-1/PD-L1 Interaction Blockage in Mice Model.
Targeting the PD-1/PD-L1 Immune Evasion Axis With DNA Aptamers as a Novel Therapeutic Strategy for the Treatment of Disseminated Cancers.
Coinhibitory molecule PD-1 as a potential target for the immunotherapy of multiple myeloma.
Transcriptional and epigenetic regulation of PD-1 expression.
Negative regulation of T-cell function by PD-1.
Mechanisms underlying low-clinical responses to PD-1/PD-L1 blocking antibodies in immunotherapy of cancer: a key role of exosomal PD-L1.
The Extrinsic and Intrinsic Roles of PD-L1 and Its Receptor PD-1: Implications for Immunotherapy Treatment.
The role of exosomal PD-L1 in tumor immunotherapy.
A snapshot of the PD-1/PD-L1 pathway.
Immune Checkpoint Blockade in Cancer Immunotherapy: Mechanisms, Clinical Outcomes, and Safety Profiles of PD-1/PD-L1 Inhibitors.
Computational design of a synthetic PD-1 agonist.
Prospective role of PD-1/PD-L1 immune checkpoint inhibitors in GI cancer.

Programmed Cell Death-1 and Its Ligands as Targets for Therapy of Multiple Myeloma Patients.
Study and analysis of antitumor resistance mechanism of PD1/PD-L1 immune checkpoint blocker.
TGFβ1-Mediated SMAD3 Enhances PD-1 Expression on Antigen-Specific T Cells in Cancer.
New insights into the important roles of tumor cell-intrinsic PD-1.
Co-expression of PD-1 and PD-L1 predicts poor outcome in nasopharyngeal carcinoma.
Safety and efficacy of mutant neoantigen-specific T-cell treatment combined anti-PD-1 therapy in stage IV solid tumors.
Harnessing the PD-1 pathway in renal cell carcinoma: current evidence and future directions.
Basics of PD-1 in self-tolerance, infection, and cancer immunity.
Programmed death-1(+) T cells inhibit effector T cells at the pathological site of miliary tuberculosis.
Boosting anti-PD-1 therapy with metformin-loaded macrophage-derived microparticles.
[Immunotherapy with PD-1 and PD-L1 inhibitors for prostate cancer].
[Eukaryotic expression and functional characterization of PD-1 extracellular domain].
PD-1/PD-L1 in disease.
Overcoming Resistance to Immune Checkpoint Inhibitors in Head and Neck Squamous Cell Carcinomas.
PD-1 and PD-L1 expression in Kaposi sarcoma: A comparative study according to the pathological stage and clinical characteristics.
PARP Targeted Alpha-Particle Therapy Enhances Response to PD-1 Immune-Checkpoint Blockade in a Syngeneic Mouse Model of Glioblastoma.
PD-1-PD-L1 immune-checkpoint blockade in malignant lymphomas.
Anti-PD-1 antibody significantly increases therapeutic efficacy of Listeria monocytogenes (Lm)-LLO immunotherapy.
Evolving landscape of PD-L2: bring new light to checkpoint immunotherapy.
CU06-1004-Induced Vascular Normalization Improves Immunotherapy by Modulating Tumor Microenvironment via Cytotoxic T Cells.
Co-expression of AFAP1-AS1 and PD-1 predicts poor prognosis in nasopharyngeal carcinoma.
Cancer Immunotherapy: An Overview of Small Molecules as Inhibitors of the Immune Checkpoint PD-1/PD-L1 (2015-2021).
Dual pH-sensitive nanodrug blocks PD-1 immune checkpoint and uses T cells to deliver NF-κB inhibitor for antitumor immunotherapy.
Characterization of PD-1 upregulation on tumor-infiltrating lymphocytes in human and murine gliomas and preclinical therapeutic blockade.
Neutralization of PD-L2 is Essential for Overcoming Immune Checkpoint Blockade Resistance in Ovarian Cancer.
Role of programmed death ligands in effective T-cell interactions in extranodal natural killer/T-cell lymphoma.
Cloning and identification of porcine programmed death 1.
PD-1/PD-L1 in Cancer: Pathophysiological, Diagnostic and Therapeutic Aspects.
PD-L1(P146R) is prognostic and a negative predictor of response to immunotherapy in gastric cancer.
Bone marrow-derived mesenchymal stem cells inhibit CD8(+) T cell immune responses via PD-1/PD-L1 pathway in multiple myeloma.
Roles of PD-1/PD-L1 Pathway: Signaling, Cancer, and Beyond.
Programmed cell death 1 ligand 1 and tumor-infiltrating CD8+ T lymphocytes are prognostic factors of human ovarian cancer.
Expression of PD-1/PD-L1 and PD-L2 in peripheral T-cells from non-small cell lung cancer patients.

Immune checkpoint blockade: the role of PD-1-PD-L axis in lymphoid malignancies.
Glucocorticoid and PD-1 Cross-Talk: Does the Immune System Become Confused?
Repeated PD-1/PD-L1 monoclonal antibody administration induces fatal xenogeneic hypersensitivity reactions in a murine model of breast cancer.
BRCA1/2 and TP53 mutation status associates with PD-1 and PD-L1 expression in ovarian cancer.
Advance investigation on synthetic small-molecule inhibitors targeting PD-1/PD-L1 signaling pathway.
Programmed death ligand 2 in cancer-induced immune suppression.
Engineering a High-Affinity PD-1 Peptide for Optimized Immune Cell-Mediated Tumor Therapy.
The role of exosomal PD-L1 in tumor progression and immunotherapy.
The PD-1:PD-L1 immune inhibitory checkpoint in Helicobacter pylori infection and gastric cancer: a comprehensive review and future perspectives.
The current advances and future directions of PD-1/PD-L1 blockade in head and neck squamous cell carcinoma (HNSCC) in the era of immunotherapy.
PD-1/PD-L1 Pathway: A Therapeutic Target in CD30+ Large Cell Lymphomas.
[The Role of PD-1/PD-L1 Signaling Pathway in Antitumor Immune Response].
A tumor-intrinsic PD-L1/NLRP3 inflammasome signaling pathway drives resistance to anti-PD-1 immunotherapy.
PD-1 blockade in rhesus macaques: impact on chronic infection and prophylactic vaccination.
Programmed death ligand-1 over-expression correlates with malignancy and contributes to immune regulation in ovarian cancer.
Programmed cell death 1 expression is associated with inferior survival in patients with primary central nervous system lymphoma.
Myeloid-derived suppressor cells regulate the immunosuppressive functions of PD-1(-)PD-L1(+) Bregs through PD-L1/PI3K/AKT/NF- $\kappa$ B axis in breast cancer.
PD-1-PD-L1 immune-checkpoint blockade in B-cell lymphomas.
Landscape of PD-1/PD-L1 Regulation and Targeted Immunotherapy.
PD-1/PD-L1 Pathway in Breast Cancer.
Subverting the adaptive immune resistance mechanism to improve clinical responses to immune checkpoint blockade therapy.
A soluble form of CD80 enhances antitumor immunity by neutralizing programmed death ligand-1 and simultaneously providing costimulation.
Biological Characteristics and Clinical Significance of Soluble PD-1/PD-L1 and Exosomal PD-L1 in Cancer.
The Microenvironment of Head and Neck Cancers: Papillomavirus Involvement and Potential Impact of Immunomodulatory Treatments.
Interferon Receptor Signaling Pathways Regulating PD-L1 and PD-L2 Expression.
Blockade of bovine PD-1 increases T cell function and inhibits bovine leukemia virus expression in B cells in vitro.
PD-1 Blunts the Function of Ovarian Tumor-Infiltrating Dendritic Cells by Inactivating NF- $\kappa$ B.
Biomarkers that may predict response to immunotherapy in ovarian malignancies.
High Quality Performance of Novel Immunoassays for the Sensitive Quantification of Soluble PD-1, PD-L1 and PD-L2 in Blood.
"Built-in" PD-1 blocker to rescue NK-92 activity from PD-L1-mediated tumor escape mechanisms.
Targeting Protein Kinases to Enhance the Response to anti-PD-1/PD-L1 Immunotherapy.
The crosstalk between H. pylori virulence factors and the PD1:PD-L1 immune checkpoint inhibitors in progression to gastric cancer.

Albendazole induces immunotherapy response by facilitating ubiquitin-mediated PD-L1 degradation.
In vitro assay for the development of small molecule inhibitors targeting PD-1/PD-L1.
Diurnal Expression of PD-1 on Tumor-Associated Macrophages Underlies the Dosing Time-Dependent Antitumor Effects of the PD-1/PD-L1 Inhibitor BMS-1 in B16/BL6 Melanoma-Bearing Mice.
PD-1 blockade augments Th1 and Th17 and suppresses Th2 responses in peripheral blood from patients with prostate and advanced melanoma cancer.
Regulation of PD-1/PD-L1 Pathway in Cancer by Noncoding RNAs.
Blockade of the checkpoint PD-1 by its ligand PD-L1 and the immuno-oncological drugs pembrolizumab and nivolumab.
Clinical significance and therapeutic potential of the programmed death-1 ligand/programmed death-1 pathway in human pancreatic cancer.
Extracellular vesicle-based checkpoint regulation and immune state in cancer.
PD-1 /PD-L1 checkpoint in hematological malignancies.
Combination immunotherapy with interleukin-2 surface-modified tumor cell vaccine and programmed death receptor-1 blockade against renal cell carcinoma.
Expressions of programmed death (PD)-1 and PD-1 ligand (PD-L1) in cervical intraepithelial neoplasia and cervical squamous cell carcinomas are of prognostic value and associated with human papillomavirus status.
[Expression and significance of B7-H1 and its receptor PD-1 in human gastric carcinoma].
Tumor-derived exosomes in the PD-1/PD-L1 axis: Significant regulators as well as promising clinical targets.
Peptide-based and small synthetic molecule inhibitors on PD-1/PD-L1 pathway: A new choice for immunotherapy?
The immune checkpoint molecules PD-1, PD-L1, TIM-3 and LAG-3 in diffuse large B-cell lymphoma.
Development and Fit-for-Purpose Validation of a Soluble Human Programmed Death-1 Protein Assay.
Sequential administration of anti-PD-1 and anti-Tim-3 combined with an SA-GM-CSF-anchored vaccine overcomes adaptive immune resistance to reject established bladder cancer.
Programmed death-1 blockade enhances the antitumor effects of peptide vaccine-induced peptide-specific cytotoxic T lymphocytes.
PD-1-Targeted Discovery of Peptide Inhibitors by Virtual Screening, Molecular Dynamics Simulation, and Surface Plasmon Resonance.
Increased Tim-3 expression on TILs during treatment with the Anchored GM-CSF vaccine and anti-PD-1 antibodies is inversely correlated with response in prostate cancer.
The perivascular microenvironment in Epstein-Barr virus positive primary central nervous system lymphoma: The role of programmed cell death 1 and programmed cell death ligand 1.
Immunophenotype based on inflammatory cells, PD-1/PD-L1 signalling pathway and M2 macrophages predicts survival in gastric cancer.
Correlation of T Cell Immunoglobulin and ITIM Domain (TIGIT) and Programmed Death 1 (PD-1) with Clinicopathological Characteristics of Renal Cell Carcinoma May Indicate Potential Targets for Treatment.
Expression of programmed cell death ligand 1 and programmed cell death 1 in cutaneous warts.
Inhibition of tumor-associated macrophages by trabectedin improves the antitumor adaptive immunity in response to anti-PD-1 therapy.
Immune escape to PD-L1/PD-1 blockade: seven steps to success (or failure).
Lichenoid dermatitis in three patients with metastatic melanoma treated with anti-PD-1 therapy.
PD-L1 and B7-1 Cis-Interaction: New Mechanisms in Immune Checkpoints and Immunotherapies.
Emerging Role of Ubiquitination in the Regulation of PD-1/PD-L1 in Cancer Immunotherapy.

Restoration of p53 activity via intracellular protein delivery sensitizes triple negative breast cancer to anti-PD-1 immunotherapy.
Immune checkpoint expression on peripheral cytotoxic lymphocytes in cervical cancer patients: moving beyond the PD-1/PD-L1 axis.
Emerging treatment options for the management of Hodgkin's lymphoma: clinical utility of nivolumab.
Biomarkers of response to PD-1 pathway blockade.
Pembrolizumab for the treatment of patients with recurrent locally advanced or metastatic gastric or gastroesophageal junction cancer: an evidence-based review of place in therapy.
Prognostic Impact of PD-1 and Tim-3 Expression in Tumor Tissue in Stage I-III Colorectal Cancer.
Atezolizumab-induced type 1 diabetes mellitus in a patient with metastatic renal cell carcinoma.
PD-L1 and IDO1 are potential targets for treatment in patients with primary diffuse large B-cell lymphoma of the CNS.
Blocking programmed death-1 ligand-PD-1 interactions by local gene therapy results in enhancement of antitumor effect of secondary lymphoid tissue chemokine.
Molecular characterization of immunoinhibitory factors PD-1/PD-L1 in chickens infected with Marek's disease virus.
A canine chimeric monoclonal antibody targeting PD-L1 and its clinical efficacy in canine oral malignant melanoma or undifferentiated sarcoma.
Immune checkpoint silencing using RNAi-incorporated nanoparticles enhances antitumor immunity and therapeutic efficacy compared with antibody-based approaches.
Bifunctional PD-1 $\times$ $\alpha$ CD3 $\times$ $\alpha$ CD33 fusion protein reverses adaptive immune escape in acute myeloid leukemia.
Promises and Pitfalls in the Use of PD-1/PD-L1 Inhibitors in Multiple Myeloma.
The Roles of microRNAs in Regulating the Expression of PD-1/PD-L1 Immune Checkpoint.
Abrine, an IDO1 inhibitor, suppresses the immune escape and enhances the immunotherapy of anti-PD-1 antibody in hepatocellular carcinoma.
Immunosuppressive CD10+ALPL+ neutrophils promote resistance to anti-PD-1 therapy in HCC by mediating irreversible exhaustion of T cells.
Targeting the PD-1/ PD-L1 interaction in nasopharyngeal carcinoma.
A single dose of neoadjuvant PD-1 blockade predicts clinical outcomes in resectable melanoma.
Monitoring Immune Checkpoint Regulators as Predictive Biomarkers in Hepatocellular Carcinoma.
Nivolumab for the treatment of cancer.
Molecular Interactions of Antibody Drugs Targeting PD-1, PD-L1, and CTLA-4 in Immuno-Oncology.
Traditional Chinese medicine inhibits PD-1/PD-L1 axis to sensitize cancer immunotherapy: a literature review.
Clinical significance of PD-1/PD-Ls gene amplification and overexpression in patients with hepatocellular carcinoma.
Rs10204525 Polymorphism of the Programmed Death (PD-1) Gene Is Associated with Increased Risk in a Saudi Arabian Population with Colorectal Cancer.
Programmed Cell Death 1 and Programmed Cell Death Ligands in Extranodal Natural Killer/T Cell Lymphoma: Expression Pattern and Potential Prognostic Relevance.
$\beta$ -Catenin Activation Promotes Immune Escape and Resistance to Anti-PD-1 Therapy in Hepatocellular Carcinoma.
Disabling immune tolerance by programmed death-1 blockade with pidilizumab after autologous hematopoietic stem-cell transplantation for diffuse large B-cell lymphoma: results of an international phase II trial.

Exosomal PD-L1: Roles in Tumor Progression and Immunotherapy.
The effect of chemotherapy on programmed cell death 1/programmed cell death 1 ligand axis: some chemotherapeutic drugs may finally work through immune response.
PD-1 Deficiency Promotes Macrophage Activation and T-Helper Cell Type 1/T-Helper Cell Type 17 Response in Pneumocystis Pneumonia.
Robust Preanalytical Performance of Soluble PD-1, PD-L1 and PD-L2 Assessed by Sensitive ELISAs in Blood.
Targeting HIF-1 $\alpha$ abrogates PD-L1-mediated immune evasion in tumor microenvironment but promotes tolerance in normal tissues.
A novel tetravalent bispecific antibody targeting programmed death 1 and tyrosine-protein kinase Met for treatment of gastric cancer.
Signaling pathways and immune evasion mechanisms in classical Hodgkin lymphoma.
Synergistic efficacy of curcumin and anti-programmed cell death-1 in hepatocellular carcinoma.
A review of the PD-1/PD-L1 checkpoint in bladder cancer: From mediator of immune escape to target for treatment.
Microenvironment Remodeling and Subsequent Clinical Implications in Diffuse Large B-Cell Histologic Variant of Richter Syndrome.
Overexpressed miR-195 attenuated immune escape of diffuse large B-cell lymphoma by targeting PD-L1.
Several immune escape patterns in non-Hodgkin's lymphomas.
Abrogation of HnRNP L enhances anti-PD-1 therapy efficacy via diminishing PD-L1 and promoting CD8(+) T cell-mediated ferroptosis in castration-resistant prostate cancer.
Coexisting Alterations of MHC Class I Antigen Presentation and IFN $\gamma$ Signaling Mediate Acquired Resistance of Melanoma to Post-PD-1 Immunotherapy.
Association of CDKN2A/B mutations, PD-1, and PD-L1 with the risk of acute lymphoblastic leukemia in children.
PD-1/PD-L1 interactions inhibit antitumor immune responses in a murine acute myeloid leukemia model.
Immune evasion phenotype is common in Richter transformation diffuse large B-cell lymphoma variant.
Elevated nuclear PIGL disrupts the cMyc/BRD4 axis and improves PD-1 blockade therapy by dampening tumor immune evasion.
Dynamic single-cell RNA sequencing identifies immunotherapy persister cells following PD-1 blockade.
Treatment of refractory/relapsed extranodal NK/T cell lymphoma with decitabine plus anti-PD-1: A case report.
Ex Vivo Expanded and Activated Natural Killer Cells Prolong the Overall Survival of Mice with Glioblastoma-like Cell-Derived Tumors.
Therapy Resistance in Neoadjuvantly Treated Gastric Cancer and Cancer of the Gastroesophageal Junction is Associated with an Increased Expression of Immune Checkpoint Inhibitors-Comparison Against a Therapy Naïve Cohort.
Mouse-Derived Gastric Organoid and Immune Cell Co-culture for the Study of the Tumor Microenvironment.
Midazolam exhibits antitumour and enhances the efficiency of Anti-PD-1 immunotherapy in hepatocellular carcinoma.
CHI3L1 regulates PD-L1 and anti-CHI3L1-PD-1 antibody elicits synergistic antitumor responses.
C. tropicalis promotes CRC by down-regulating tumor cell-intrinsic PD-1 receptor via autophagy.

Increasing the expression of programmed death ligand 2 (PD-L2) but not 4-1BB ligand in colorectal cancer cells.
Programmed cell death ligands expression in pheochromocytomas and paragangliomas: Relationship with the hypoxic response, immune evasion and malignant behavior.
Combination PD-1 and PD-L1 Blockade Promotes Durable Neoantigen-Specific T Cell-Mediated Immunity in Pancreatic Ductal Adenocarcinoma.
Comprehensive immunohistochemical analysis of immune checkpoint molecules in adult T cell leukemia/lymphoma.
Development and characterization of mouse anti-canine PD-L1 monoclonal antibodies and their expression in canine tumors by immunohistochemistry in vitro.
PD-1/PD-L1 pathway: A double-edged sword in periodontitis.
An unexpected role for PD-L1 in front-rear polarization and directional migration.
Programmed Cell Death-Ligand 1 in Head and Neck Squamous Cell Carcinoma: Molecular Insights, Preclinical and Clinical Data, and Therapies.
PD-L1 is highly expressed in Enzalutamide resistant prostate cancer.
Microneedles loaded with anti-PD-1-cisplatin nanoparticles for synergistic cancer immuno-chemotherapy.
PD-1 Coexpression Gene Analysis and the Regulatory Network in Endometrial Cancer Based on Bioinformatics Analysis.
Recent advance of peptide-based molecules and nonpeptidic small-molecules modulating PD-1/PD-L1 protein-protein interaction or targeting PD-L1 protein degradation.
Norcholic Acid Promotes Tumor Progression and Immune Escape by Regulating Farnesoid X Receptor in Hepatocellular Carcinoma.
Biological Function of PD-L2 and Correlation With Overall Survival in Type II Endometrial Cancer.
CircSOD2 Contributes to Tumor Progression, Immune Evasion and Anti-PD-1 Resistance in Hepatocellular Carcinoma by Targeting miR-497-5p/ANXA11 Axis.
Virtual Screening and In Vitro Evaluation of PD-1 Dimer Stabilizers for Uncoupling PD-1/PD-L1 Interaction from Natural Products.
Immune evasion mechanisms and immune checkpoint inhibition in advanced merkel cell carcinoma.
Diagnostic Predictors of Immunotherapy Response in Head and Neck Squamous Cell Carcinoma.
Inducible expression of B7-H1 (PD-L1) and its selective role in tumor site immune modulation.
Engineered DBCO+PD-1 Nanovesicles Carrying 1-MT for Cancer-Targeted Immunotherapy.
Lack of an association of programmed cell death-1 PD1.3 polymorphism with risk of hepatocellular carcinoma susceptibility in Turkish population: a case-control study.
Soluble Programmed Death Ligand-1 (sPD-L1): A Pool of Circulating Proteins Implicated in Health and Diseases.
Sulfisoxazole Elicits Robust Antitumour Immune Response Along with Immune Checkpoint Therapy by Inhibiting Exosomal PD-L1.
The expression of PD-1 ligands and IDO1 by macrophage/microglia in primary central nervous system lymphoma.
Synergistic efficacy of simultaneous anti-TGF- $\beta$ /VEGF bispecific antibody and PD-1 blockade in cancer therapy.
Increased PD-1 Level in Severe Cervical Injury Is Associated With the Rare Programmed Cell Death 1 (PDCD1) rs36084323 A Allele in a Dominant Model.
Ubiquilin-4 induces immune escape in gastric cancer by activating the notch signaling pathway.
The expression and clinical significance of programmed cell death receptor 1 and its ligand in tumor tissues of patients with extranodal nasal NK/T cell lymphoma.

The Stone Guest: How Does pH Affect Binding Properties of PD-1/PD-L1 Inhibitors?
Evaluation of circulating PD-1 and PD-L1 as diagnostic biomarkers in dogs with tumors.
An IL-27/Stat3 axis induces expression of programmed cell death 1 ligands (PD-L1/2) on infiltrating macrophages in lymphoma.
STAT3 Induces Immunosuppression by Upregulating PD-1/PD-L1 in HNSCC.
Small molecule inhibitors targeting the PD-1/PD-L1 signaling pathway.
PD-1 blockade inhibits hematogenous spread of poorly immunogenic tumor cells by enhanced recruitment of effector T cells.
B2M and JAK1/2-mutated MSI-H Colorectal Carcinomas Can Benefit From Anti-PD-1 Therapy.
Soluble PD-1 but Not PD-L1 Levels Predict Poor Outcome in Patients with High-Risk Diffuse Large B-Cell Lymphoma.
Mutations Associated with Acquired Resistance to PD-1 Blockade in Melanoma.
The Circular RNA circFGFR4 Facilitates Resistance to Anti-PD-1 of Triple-Negative Breast Cancer by Targeting the miR-185-5p/CXCR4 Axis.
Bispecific antibody targeting TGF- $\beta$ and PD-L1 for synergistic cancer immunotherapy.
[Development of programmed death receptor-1 and programmed death receptor-1 ligand in oral squamous cell carcinoma].
Expression regulation and function of PD-1 and PD-L1 in T lymphoma cells.
ATR-mediated CD47 and PD-L1 up-regulation restricts radiotherapy-induced immune priming and abscopal responses in colorectal cancer.
Combining PD-1/PD-L1 blockade with type I interferon in cancer therapy.
Tumor immune response and immunotherapy in gastric cancer.
Dosimetry Prediction for Clinical Translation of (64)Cu-Pembrolizumab ImmunoPET Targeting Human PD-1 Expression.
Melatonin inhibits EMT and PD-L1 expression through the ERK1/2/FOSL1 pathway and regulates anti-tumor immunity in HNSCC.
PD-1/PD-L1 Checkpoint Inhibitors Are Active in the Chicken Embryo Model and Show Antitumor Efficacy In Ovo.
Expression and clinical significance of PD-1 in hepatocellular carcinoma tissues detected by a novel mouse anti-human PD-1 monoclonal antibody.
Upregulation of PD-1 on CD4 <sup>+</sup> CD25 <sup>+</sup> T cells is associated with immunosuppression in liver of mice infected with Echinococcus multilocularis.
Prognostic influence of tumor microenvironment after hypofractionated radiation and surgery for mesothelioma.
Overexpression of PD-L1 significantly associates with tumor aggressiveness and postoperative recurrence in human hepatocellular carcinoma.
The PD-1 expressing immune phenotype of T cell exhaustion is prominent in the 'immunoreactive' microenvironment of colorectal carcinoma.
Programmed death ligand 2 regulates arginase induction and modifies Trypanosoma cruzi survival in macrophages during murine experimental infection.
PD-1 blockade with nivolumab in relapsed or refractory Hodgkin's lymphoma.
Intracellular mechanisms of tumor cells' immunoresistance.
The effect of Wnt/ $\beta$ -catenin signaling on PD-1/PDL-1 axis in HPV-related cervical cancer.
STINGing the Tumor's immune evasion mechanism.
Discrepant antitumor efficacies of three CpG oligodeoxynucleotide classes in monotherapy and co-therapy with PD-1 blockade.



Soluble programmed cell death protein 1 (sPD-1) and the soluble programmed cell death ligands 1 and 2 (sPD-L1 and sPD-L2) in lymphoid malignancies.
From Tumor Immunology to Immunotherapy in Gastric and Esophageal Cancer.
Dihydroartemisinin broke the tumor immunosuppressive microenvironment by inhibiting YAP1 expression to enhance anti-PD-1 efficacy.
Merkel Cell Carcinoma in the Age of Immunotherapy: Facts and Hopes.
Radiosynthesis and preclinical PET evaluation of (89)Zr-nivolumab (BMS-936558) in healthy non-human primates.
Beyond PD-L1-Identification of Further Potential Therapeutic Targets in Oral Cancer.
Genomic analyses of PMBL reveal new drivers and mechanisms of sensitivity to PD-1 blockade.
Cisplatin-induced programmed cell death ligand-2 expression is associated with metastasis ability in oral squamous cell carcinoma.
Inhibition of histone lysine-specific demethylase 1 elicits breast tumor immunity and enhances antitumor efficacy of immune checkpoint blockade.
LncRNA SNHG14/miR-5590-3p/ZEB1 positive feedback loop promoted diffuse large B cell lymphoma progression and immune evasion through regulating PD-1/PD-L1 checkpoint.
Immune-checkpoint molecules on regulatory T-cells as a potential therapeutic target in head and neck squamous cell cancers.
Relationship between expression of PD-L1 and PD-L2 on esophageal squamous cell carcinoma and the antitumor effects of CD8 <sup>+</sup> T cells.
Emerging role of immunotherapy in urothelial carcinoma-Immunobiology/biomarkers.
PD-L1 Is Not Constitutively Expressed on Tasmanian Devil Facial Tumor Cells but Is Strongly Upregulated in Response to IFN- $\gamma$ and Can Be Expressed in the Tumor Microenvironment.
Circulating PD-1 (+) cells may participate in immune evasion in peripheral T-cell lymphoma and chidamide enhance antitumor activity of PD-1 (+) cells.
SMYD3 represses tumor-intrinsic interferon response in HPV-negative squamous cell carcinoma of the head and neck.
Intratumoral Immune Response to Gastric Cancer Varies by Molecular and Histologic Subtype.
Molecular dynamics identifies semi-rigid domains in the PD-1 checkpoint receptor bound to its natural ligand PD-L1.
Dynamic host immune response in virus-associated cancers.
Prognostic significance of circulating PD-1, PD-L1, pan-BTN3As, BTN3A1 and BTLA in patients with pancreatic adenocarcinoma.
MAGE-C3 promotes cancer metastasis by inducing epithelial-mesenchymal transition and immunosuppression in esophageal squamous cell carcinoma.
Selection of new immunotherapy targets for NK/T cell lymphoma.
Tumor PD-L1 engages myeloid PD-1 to suppress type I interferon to impair cytotoxic T lymphocyte recruitment.
A Preclinical Human-Derived Autologous Gastric Cancer Organoid/Immune Cell Co-Culture Model to Predict the Efficacy of Targeted Therapies.
Met inhibition revokes IFN $\gamma$ -induction of PD-1 ligands in MET-amplified tumours.
Association of Ipilimumab With Safety and Antitumor Activity in Women With Metastatic or Recurrent Human Papillomavirus-Related Cervical Carcinoma.
Proteogenomic analysis of NCC-S1M, a gastric cancer stem cell-like cell line that responds to anti-PD-1.
Development of Novel ImmunoPET Tracers to Image Human PD-1 Checkpoint Expression on Tumor-Infiltrating Lymphocytes in a Humanized Mouse Model.

Programmed cell death-ligand 1 (PD-L1)(+) tumour cells and low-reacting programmed cell death 1 (PD1)(+) tumour-infiltrating lymphocytes predict poor prognosis in Epstein-Barr virus(+) diffuse large B-cell lymphoma.
Coexpression of PD-L1/PD-1 with CXCR3/CD36 and IL-19 Increase in Extranodal Lymphoma.
Sphingosine kinase 1 promotes tumor immune evasion by regulating the MTA3-PD-L1 axis.
Immune Checkpoint Glycoproteins Have Polymorphism: Are Monoclonal Antibodies Too Specific? [Microenvironment in classical Hodgkin lymphoma].
The Expression Pattern and Clinical Significance of the Immune Checkpoint Regulator VISTA in Human Breast Cancer.
Pembrolizumab for the treatment of disease relapse after allogeneic hematopoietic stem cell transplantation.
T-cell coregulatory molecule expression in urothelial cell carcinoma: clinicopathologic correlations and association with survival.
Phase IA/IB study of single-agent tislelizumab, an investigational anti-PD-1 antibody, in solid tumors.
The implication of the PD-1/PD-L1 checkpoint in chronic periodontitis suggests novel therapeutic opportunities with natural products.
Evaluation of the Association between Programmed Cell Death-1 Gene Polymorphisms and Hepatocellular Carcinoma Susceptibility in Turkish Subjects. A Pilot Study.
ERBB2D16 Expression in HER2 Positive Gastric Cancer Is Associated With Resistance to Trastuzumab.
Checkpoint inhibition of PD-L1 and CTLA-4 in a child with refractory acute leukemia.
PD-1 blockade combined with ICE regimen in relapsed/refractory diffuse large B-cell lymphoma.
Program death 1 (PD1) haplotyping in patients with breast carcinoma.
Increased level of myeloid-derived suppressor cells, programmed death receptor ligand 1/programmed death receptor 1, and soluble CD25 in Sokal high risk chronic myeloid leukemia.
PD-1 Haplotype Combinations and Susceptibility of Patients to Squamous Cell Carcinomas of Head and Neck.
New antibody approaches to lymphoma therapy.
The Prognostic Role of CD8(+) T Lymphocytes in Childhood Adrenocortical Carcinomas Compared to Ki-67, PD-1, PD-L1, and the Weiss Score.
PD-L2 glycosylation promotes immune evasion and predicts anti-EGFR efficacy.
Calcium Channel Blocker Nifedipine Suppresses Colorectal Cancer Progression and Immune Escape by Preventing NFAT2 Nuclear Translocation.
Blockade of IL-6 inhibits tumor immune evasion and improves anti-PD-1 immunotherapy.
Multifunctional Immunoliposomes Combining Catalase and PD-L1 Antibodies Overcome Tumor Hypoxia and Enhance Immunotherapeutic Effects Against Melanoma.
MTDH antisense oligonucleotides reshape the immunosuppressive tumor microenvironment to sensitize Hepatocellular Carcinoma to immune checkpoint blockade therapy.
Codelivery of Anti-PD-1 Antibody and Paclitaxel with Matrix Metalloproteinase and pH Dual-Sensitive Micelles for Enhanced Tumor Chemoimmunotherapy.
B7-CD28 gene family expression is associated with prognostic and immunological characteristics of diffuse large B-cell lymphoma.
Large cell morphology, CMYC+ tumour cells, and PD-1+ tumour cell/intense PD-L1+ cell reactions are important prognostic factors in nodal peripheral T-cell lymphomas with T follicular helper markers.
A Systematic Review and Meta-Analysis on the Significance of TIGIT in Solid Cancers: Dual TIGIT/PD-1 Blockade to Overcome Immune-Resistance in Solid Cancers.
Glutaminyl cyclase is an enzymatic modifier of the CD47- SIRPα axis and a target for cancer immunotherapy.

Clinical Role of Programmed Cell Death-1 Expression in Patients with Non-muscle-invasive Bladder Cancer Recurring After Initial Bacillus Calmette-Guérin Therapy.
A novel bispecific c-MET/PD-1 antibody with therapeutic potential in solid cancer.
Monocyte programmed death ligand-1 expression is an early marker for predicting infectious complications in acute pancreatitis.
Immune-checkpoint expression in Epstein-Barr virus positive and negative plasmablastic lymphoma: a clinical and pathological study in 82 patients.
Pembrolizumab, pomalidomide, and low-dose dexamethasone for relapsed/refractory multiple myeloma.
The novel negative checkpoint regulator VISTA is expressed in gastric carcinoma and associated with PD-L1/PD-1: A future perspective for a combined gastric cancer therapy?
Combination immunotherapy with $\alpha$ -CTLA-4 and $\alpha$ -PD-L1 antibody blockade prevents immune escape and leads to complete control of metastatic osteosarcoma.
Down Regulation of c-FLIP(L) Enhance PD-1 Blockade Efficacy in B16 Melanoma.
VISTA expression on tumor-infiltrating inflammatory cells in primary cutaneous melanoma correlates with poor disease-specific survival.
Increased Expression of Programmed Death Ligand 1 in Hepatocellular Carcinoma of Patients with Hepatitis B Virus Pre-S2 Mutant.
Precursor cells and implications of a T-cell inflamed immune response in the pre-malignant setting in Hodgkin lymphoma.
The prognostic significance of VISTA and CD33-positive myeloid cells in cutaneous melanoma and their relationship with PD-1 expression.
The effect of polymorphisms in PD-1 gene on the risk of epithelial ovarian cancer and patients' outcomes.
Effects of Tussilago farfara L. Polysaccharides on the Expression of PD-1 (CD279) and PD-L1 (CD274) in Peripheral Blood and Tumor Tissue Lymphocytes in Mice with Lewis Lung Carcinoma.
Study of Serum Soluble Programmed Death Ligand 1 as a Prognostic Factor in Hepatocellular Carcinoma in Egyptian Patients.
The B7-H1 (PD-L1) T lymphocyte-inhibitory molecule is expressed in breast cancer patients with infiltrating ductal carcinoma: correlation with important high-risk prognostic factors.
Molecular pathogenic pathways in extranodal NK/T cell lymphoma.
Fatal immune-related hepatitis with intrahepatic cholestasis and pneumonia associated with camrelizumab: A case report and literature review.
The Concordant Disruption of B7/CD28 Immune Regulators Predicts the Prognosis of Oral Carcinomas.
Oral Squamous Carcinoma Cells Express B7-H1 and B7-DC Receptors in Vivo.
miR-146a Controls Immune Response in the Melanoma Microenvironment.
Cisplatin remodels the tumor immune microenvironment via the transcription factor EB in ovarian cancer.
Therapeutic Effect of Catgut Implantation at Acupoint in a Mouse Model of Hepatocellular Carcinoma by Suppressing Immune Escape.
Nanotechnology-mediated immunochemotherapy combined with docetaxel and PD-L1 antibody increase therapeutic effects and decrease systemic toxicity.
Insights into anti-tumor immunity via the polyomavirus shared across human Merkel cell carcinomas.
Macrophage-mediated anti-tumor immunity against high-risk neuroblastoma.
B7-H1 overexpression regulates epithelial-mesenchymal transition and accelerates carcinogenesis in skin.

The potential of soluble programmed death-ligand 1 (sPD-L1) as a diagnosis marker for colorectal cancer.
Pyroptosis, a new bridge to tumor immunity.
Investigation of protein-protein interactions and hot spot region between PD-1 and PD-L1 by fragment molecular orbital method.
Recent development of selective inhibitors targeting the HDAC6 as anti-cancer drugs: Structure, function and design.
Targeting fibrinogen-like protein 1 enhances immunotherapy in hepatocellular carcinoma.
Molecular mechanism of SHP2 activation by PD-1 stimulation.
Cerebrospinal fluid soluble programmed death-ligand 1 is a useful prognostic biomarker in primary central nervous system lymphoma.
Nanobody against PDL1.
Effect of TLR4 and B7-H1 on immune escape of urothelial bladder cancer and its clinical significance.
SNP-SNP Interaction in Genes Encoding PD-1/PD-L1 Axis as a Potential Risk Factor for Clear Cell Renal Cell Carcinoma.
Immunoprofiling in Neuroendocrine Neoplasms Unveil Immunosuppressive Microenvironment.
B7-H1 expression model for immune evasion in human papillomavirus-related oropharyngeal squamous cell carcinoma.
Interferon-Gamma at the Crossroads of Tumor Immune Surveillance or Evasion.
Expression of PD-1 and PD-Ls in Kaposi's sarcoma and regulation by oncogenic herpesvirus lytic reactivation.
Immuno-Contexture and Immune Checkpoint Molecule Expression in Mismatch Repair Proficient Colorectal Carcinoma.
Tumor B7-H1 and B7-H3 expression in squamous cell carcinoma of the lung.
The COX2 Effector Microsomal PGE2 Synthase 1 is a Regulator of Immunosuppression in Cutaneous Melanoma.
LncRNA MIAT correlates with immune infiltrates and drug reactions in hepatocellular carcinoma.
The CXCL11-CXCR3A axis influences the infiltration of CD274 and IDO1 in oral squamous cell carcinoma.
The PD-1/PD-L1 axis contributes to immune metabolic dysfunctions of monocytes in chronic lymphocytic leukemia.
Characterization of the Clinical Significance of PD-1/PD-Ls Expression and Methylation in Patients With Low-Grade Glioma.
Impact of serum soluble programed death ligand 1 on end of treatment metabolic response of diffuse large B cell lymphoma patients.
Lysosome activable polymeric vorinostat encapsulating PD-L1KD for a combination of HDACi and immunotherapy.
Automated Analysis of PD1 and PDL1 Expression in Lymph Nodes and the Microenvironment of Transmissible Tumors in Tasmanian Devils.
Recall of reactive cutaneous capillary endothelial proliferation (RCCEP) induced by stereotactic body radiation therapy (SBRT) in a patient with lung squamous cell carcinoma.
Expression of Immune Checkpoint Regulators IDO, VISTA, LAG3, and TIM3 in Resected Pancreatic Ductal Adenocarcinoma.
Inverse association between programmed death ligand 1 and genes in the VEGF pathway in primary clear cell renal cell carcinoma.
Inter- and inpatient heterogeneity of indoleamine 2,3-dioxygenase expression in primary and metastatic melanoma cells and the tumour microenvironment.

Outlooks on Epstein-Barr virus associated gastric cancer.
Label-free and specific detection of soluble programmed death ligand-1 using a localized surface plasmon resonance biosensor based on excessively tilted fiber gratings.
S100a9 deficiency accelerates MDS-associated tumor escape via PD-1/PD-L1 overexpression.
HDAC is indispensable for IFN- $\gamma$ -induced B7-H1 expression in gastric cancer.
The clinical application of camrelizumab on advanced hepatocellular carcinoma.
Shuyu pills inhibit immune escape and enhance chemosensitization in hepatocellular carcinoma.
Expression of immune checkpoint molecules in endometrial carcinoma.
In situ expression and significance of B7 costimulatory molecules within tissues of human gastric carcinoma.
Liver fibrosis promotes immunity escape but limits the size of liver tumor in a rat orthotopic transplantation model.
Tumor-infiltrating lymphocytes associate with outcome in nonendemic nasopharyngeal carcinoma: a multicenter study.
PKM2 Drives Hepatocellular Carcinoma Progression by Inducing Immunosuppressive Microenvironment.
Predictive and prognostic value of PDL1 protein expression in breast cancer patients in neoadjuvant setting.
Loss of tumor suppressor PTEN function increases B7-H1 expression and immunoresistance in glioma.
Combined induction with anti-PD-1 and anti-CTLA-4 antibodies provides synergistic antitumor effects in DC-CIK cells in renal carcinoma cell lines.
HDAC3 increases HMGB3 expression to facilitate the immune escape of breast cancer cells via down-regulating microRNA-130a-3p.
Sustained low-level expression of interferon-gamma promotes tumor development: potential insights in tumor prevention and tumor immunotherapy.
The Upregulation of Molecules Related to Tumor Immune Escape in Human Pituitary Adenomas.
Recurrent PDL1 expression and PDL1 (CD274) copy number alterations in breast implant-associated anaplastic large cell lymphomas.
Increased BTLA and HVEM in gastric cancer are associated with progression and poor prognosis.
B7-H1 and B7-DC receptors of oral squamous carcinoma cells are upregulated by Porphyromonas gingivalis.
Serum soluble immune checkpoint levels predict cervical lymph node metastasis of differentiated thyroid carcinoma patients.
Protein kinase D3 regulates the expression of the immunosuppressive protein, PD-L1, through STAT1/STAT3 signaling.
Interferon regulatory factor-1 is prerequisite to the constitutive expression and IFN-gamma-induced upregulation of B7-H1 (CD274).
Rationale for immune-based therapies in Merkel polyomavirus-positive and -negative Merkel cell carcinomas.
Analysis and therapeutic targeting of the EP300 and CREBBP acetyltransferases in anaplastic large cell lymphoma and Hodgkin lymphoma.
KCNQ10T1 contributes to sorafenib resistance and programmed death-ligand-1-mediated immune escape via sponging miR-506 in hepatocellular carcinoma cells.
Thirty years of Epstein-Barr virus-associated gastric carcinoma.
B7-H4 expression in ovarian serous carcinoma: a study of 306 cases.
[Effects of lncRNA-UCA1 targeting miR-204-5p on the proliferation, migration, apoptosis and immune escape of endometrial carcinoma cells].

MACC1 regulates PDL1 expression and tumor immunity through the c-Met/AKT/mTOR pathway in gastric cancer cells.
Interleukin 23/interleukin 17 axis activated by Mycobacterium avium complex (MAC) is attenuated in patients with MAC-lung disease.
Paclitaxel induced B7-H1 expression in cancer cells via the MAPK pathway.
B7 family molecules are favorably positioned at the human maternal-fetal interface.
[The relationship between CD44 and immune escape in gastric carcinoma].
Evaluation of myrrh (Mirazid) therapy in fascioliasis and intestinal schistosomiasis in children: immunological and parasitological study.
Expression Depression of CD300LG-γ in Human Pulmonary Carcinoma.
[Relationship between MAP3K5 and Epstein-Barr virus-encoded miR-BART22 expression in nasopharyngeal carcinoma].
The Immune Checkpoint Molecule CD200 Is Associated with Tumor Grading and Metastasis in Bladder Cancer.
Maintained CD40 and loss of polarised CD40 ligand expression in oral squamous cell carcinoma.

Table A2-52, Cluster 51

Cluster 51 focuses on parasites (especially Trypanosoma cruzi, Leishmania, and Toxoplasma gondii), emphasizing their immune system evasion strategies (495)
Molecular basis of Trypanosoma cruzi and Leishmania interaction with their host(s): exploitation of immune and defense mechanisms by the parasite leading to persistence and chronicity, features reminiscent of immune system evasion strategies in cancer diseases.
Living off a fish: a trade-off between parasites and the immune system.
Intracellular parasitism: cell biological adaptations of parasitic protozoa to a life inside cells.
Purinergic signaling and infection by Leishmania: A new approach to evasion of the immune response.
The oligopeptidase B of Leishmania regulates parasite enolase and immune evasion.
The pathogenicity and virulence of Toxoplasma gondii.
How to master the host immune system? Leishmania parasites have the solutions!
Malaria Parasites: The Great Escape.
Leishmania Hijacks Myeloid Cells for Immune Escape.
The role of host autophagy machinery in controlling Toxoplasma infection.
Nematoda: genes, genomes and the evolution of parasitism.
Mechanisms of cellular invasion by intracellular parasites.
Effects of a natural parasitical infection (Lernanthropus kroyeri) on the immune system of European sea bass, Dicentrarchus labrax L.
Mechanisms of Human Innate Immune Evasion by Toxoplasma gondii.
The pathogenicity and virulence of Leishmania - interplay of virulence factors with host defenses.
Apoptotic-like Leishmania exploit the host's autophagy machinery to reduce T-cell-mediated parasite elimination.
Determination of Toxoplasma gondii Replication in Naïve and Activated Macrophages.
Deception and manipulation: the arms of leishmania, a successful parasite.
Immune defence, parasite evasion strategies and their relevance for 'macroscopic phenomena' such as virulence.
Host-parasite interactions: an intimate epigenetic relationship.
The enemy within: lipid asymmetry in intracellular parasite-host interactions.

Subversion of innate and adaptive immune responses by <i>Toxoplasma gondii</i> .
Modulation of Host-Pathogen Communication by Extracellular Vesicles (EVs) of the Protozoan Parasite <i>Leishmania</i> .
Use and abuse of dendritic cells by <i>Toxoplasma gondii</i> .
DNA vaccines against tropical parasitic diseases.
[Malaria--biological aspects of an infectious disease of importance to humans].
Modelling parasite dissemination: host cell subversion and immune evasion by <i>Toxoplasma gondii</i> .
Repeat-enriched proteins are related to host cell invasion and immune evasion in parasitic protozoa.
<i>Leishmania donovani</i> Secretory Mevalonate Kinase Regulates Host Immune Response and Facilitates Phagocytosis.
<i>Leishmania</i> and the macrophage: a multifaceted interaction.
<i>Toxoplasma gondii</i> virulence factor ROP1 reduces parasite susceptibility to murine and human innate immune restriction.
Mechanisms of pathogenesis and the evolution of parasite virulence.
Effects of <i>Toxoplasma gondii</i> infection on the brain.
Defence against the immune barrage: helminth survival strategies.
Are neutrophils important host cells for <i>Leishmania</i> parasites?
[Modulation of inflammatory cells in helminth infections].
Enlisting the mRNA Vaccine Platform to Combat Parasitic Infections.
Different regulatory mechanisms in protozoan parasitic infections.
Neuroimmunoendocrine modulation in the host by helminth parasites: a novel form of host-parasite coevolution?
Complement Evasion: An Effective Strategy That Parasites Utilize to Survive in the Host.
Integrative Approaches to Understand the Mastery in Manipulation of Host Cytokine Networks by Protozoan Parasites with Emphasis on <i>Plasmodium</i> and <i>Leishmania</i> Species.
Communication between <i>Toxoplasma gondii</i> and its host: impact on parasite growth, development, immune evasion, and virulence.
Protozoa: Pathogenesis and Defenses.
How much epigenetics and quantitative trait loci (QTL) mapping tell us about parasitism maintenance and resistance/susceptibility to hosts.
How protozoan parasites evade the immune response.
Toward an understanding of the interaction between filarial parasites and host antigen-presenting cells.
The Rhoptry Pseudokinase ROP54 Modulates <i>Toxoplasma gondii</i> Virulence and Host GBP2 Loading.
<i>Toxoplasma gondii</i> and subversion of the immune system.
The <i>Leishmania donovani</i> Ortholog of the Glycosylphosphatidylinositol Anchor Biosynthesis Cofactor PBN1 Is Essential for Host Infection.
<i>Toxoplasma</i> on the brain: understanding host-pathogen interactions in chronic CNS infection.
Induction of suppressor of cytokine signaling-1 by <i>Toxoplasma gondii</i> contributes to immune evasion in macrophages by blocking IFN-gamma signaling.
A Protective and Pathogenic Role for Complement During Acute <i>Toxoplasma gondii</i> Infection.
Effects of parasites coinfection with other pathogens on animal host: A literature review.
The Influence of Parasite Infections on Host Immunity to Co-infection With Other Pathogens.
<i>Leishmania</i> Spp-Host Interaction: There Is Always an Onset, but Is There an End?
Unravelling the cellular and molecular pathogenesis of bovine babesiosis: is the sky the limit?
Host- <i>Toxoplasma gondii</i> Coadaptation Leads to Fine Tuning of the Immune Response.

Strategies Developed by <i>Toxoplasma gondii</i> to Survive in the Host.
Histones and histone modifications in protozoan parasites.
Immune evasion, immunopathology and the regulation of the immune system.
Revisiting the Mechanisms of Immune Evasion Employed by Human Parasites.
<i>Toxoplasma gondii</i> manipulates host cell signaling pathways via its secreted effector molecules.
Mammalian apoptotic signalling pathways: multiple targets of protozoan parasites to activate or deactivate host cell death.
Cell biology of <i>Leishmania</i> .
<i>Toxoplasma gondii</i> Dissemination in the Brain Is Facilitated by Infiltrating Peripheral Immune Cells.
<i>Eimeria falciformis</i> infection of the mouse caecum identifies opposing roles of IFN $\gamma$ -regulated host pathways for the parasite development.
Protein quality control machinery in intracellular protozoan parasites: hopes and challenges for therapeutic targeting.
Modulation of innate immunity by <i>Toxoplasma gondii</i> virulence effectors.
Interactions between <i>Leishmania</i> parasites and host cells.
<i>Leishmania</i> apolipoprotein A-I expression: a possible strategy used by the parasite to evade the host's immune response.
Released Parasite-Derived Kinases as Novel Targets for Antiparasitic Therapies.
Antigenic variation as an exploitable weakness of babesial parasites.
<i>Leishmania</i> major infection in humanized mice induces systemic infection and provokes a nonprotective human immune response.
Immunomodulation in trichinellosis: does <i>Trichinella</i> really escape the host immune system?
Host Lipid Bodies as Platforms for Intracellular Survival of Protozoan Parasites.
The <i>Toxoplasma</i> effector GRA28 promotes parasite dissemination by inducing dendritic cell-like migratory properties in infected macrophages.
Parasite Cystatin: Immunomodulatory Molecule with Therapeutic Activity against Immune Mediated Disorders.
Role of Cytokines in the Pathogenesis of Visceral Leishmaniasis.
A review on the interactions between dendritic cells, filarial parasite and parasite-derived molecules in regulating the host immune responses.
In Vivo Biotinylation of the <i>Toxoplasma</i> Parasitophorous Vacuole Reveals Novel Dense Granule Proteins Important for Parasite Growth and Pathogenesis.
Protein glycosylation in <i>Leishmania</i> spp.
<i>Toxoplasma gondii</i> Extends the Life Span of Infected Human Neutrophils by Inducing Cytosolic PCNA and Blocking Activation of Apoptotic Caspases.
<i>Leishmania</i> kinetoplast DNA contributes to parasite burden in infected macrophages: Critical role of the cGAS-STING-TBK1 signaling pathway in macrophage parasitemia.
Computational and Experimental Approaches to Predict Host-Parasite Protein-Protein Interactions.
Cytokine saga in visceral leishmaniasis.
Parasite immune escape: new views into host-parasite interactions.
The role of macrophages in protective and pathological responses to <i>Toxoplasma gondii</i> .
Parasite immune evasion and exploitation: reflections and projections.
Host-parasite interactions in some fish species.
Sink or swim: lipid rafts in parasite pathogenesis.
Controlling and coordinating development in vector-transmitted parasites.
Regulation of gene expression in protozoa parasites.



Leishmania infantum and Leishmania braziliensis: Differences and Similarities to Evade the Innate Immune System.
Src- and Abl-family kinases activate spleen tyrosine kinase to maximize phagocytosis and Leishmania infection.
Proteins involved in the biosynthesis of lipophosphoglycan in Leishmania: a comparative genomic and evolutionary analysis.
A lesson in survival, by Giardia lamblia.
Pathogenicity and virulence of Entamoeba histolytica, the agent of amoebiasis.
New approaches in vaccine development for parasitic infections.
Antibodies to variable surface antigens induce antigenic variation in the intestinal parasite Giardia lamblia.
Integrative omics analysis highlights the immunomodulatory effects of the parasitic dinoflagellate hhematodinium on crustacean hemocytes.
Fancy a gene? A surprisingly complex evolutionary history of peroxiredoxins.
Determinants of Innate Immunity in Visceral Leishmaniasis and Their Implication in Vaccine Development.
Role of calreticulin from parasites in its interaction with vertebrate hosts.
Mechanisms of immune evasion in leishmaniasis.
Nitric oxide hinders antibody clearance from the surface of Trypanoplasma borreli and increases susceptibility to complement-mediated lysis.
Diverse mechanisms employed by Toxoplasma gondii to inhibit IFN-gamma-induced major histocompatibility complex class II gene expression.
Biological roles of proteases in parasitic protozoa.
Parasitism as the main factor shaping peptide vocabularies in current organisms.
Genes and susceptibility to leishmaniasis.
Host-parasite interactions and the evolution of ploidy.
Selective modification and immune evasion: a hypothesis.
Paradoxical immune response in leishmaniasis: The role of toll-like receptors in disease progression.
Biliary parasites.
Parasite-Produced MIF Cytokine: Role in Immune Evasion, Invasion, and Pathogenesis.
Long-term live imaging reveals cytosolic immune responses of host hepatocytes against Plasmodium infection and parasite escape mechanisms.
B-Cell Response during Protozoan Parasite Infections.
How to succeed in parasitic life without sex? Asking Leishmania.
Proteases in parasitic diseases.
Recombinant Mouse Prolactin Confers Partial Protection Against Toxoplasma gondii Infection in a Pre-treated Experimental Murine Model.
Protein O-fucosyltransferase 2-mediated O-glycosylation of the adhesin MIC2 is dispensable for Toxoplasma gondii tachyzoite infection.
Parasite-Derived Proteins for the Treatment of Allergies and Autoimmune Diseases.
Epigenetic Control of IFN- $\gamma$ Host Responses During Infection With Toxoplasma gondii.
Evasion of Human Neutrophil-Mediated Host Defense during Toxoplasma gondii Infection.
Phosphorylation of mouse immunity-related GTPase (IRG) resistance proteins is an evasion strategy for virulent Toxoplasma gondii.
Toxoplasma gondii inhibits MHC class II expression in neural antigen-presenting cells by down-regulating the class II transactivator CIITA.

On being a parasite in an invertebrate host: a short survival course.
Understanding the role of antibodies in murine infections with <i>Heligmosomoides (polygyrus) bakeri</i> : 35 years ago, now and 35 years ahead.
Identification and Characterization of miRNAs in Response to <i>Leishmania donovani</i> Infection: Delineation of Their Roles in Macrophage Dysfunction.
Host parasite communications-Messages from helminths for the immune system: Parasite communication and cell-cell interactions.
Why does immunity to parasites take so long to develop?
Apoptosis as a host defense mechanism in <i>Crassostrea virginica</i> and its modulation by <i>Perkinsus marinus</i> .
Cell biology and immunology of <i>Leishmania</i> .
Down-regulation of MHC class II molecules and inability to up-regulate class I molecules in murine macrophages after infection with <i>Toxoplasma gondii</i> .
Touching the Surface: Diverse Roles for the Flagellar Membrane in Kinetoplastid Parasites.
Phosphatidylserine exposure on the surface of <i>Leishmania amazonensis</i> amastigotes modulates in vivo infection and dendritic cell function.
Virus-like Particle Display of the $\alpha$ -Gal Carbohydrate for Vaccination against <i>Leishmania</i> Infection.
Delayed IL-12 production by macrophages during <i>Toxoplasma gondii</i> infection is regulated by miR-187.
Immune evasive mechanisms contributing to persistent <i>Leishmania donovani</i> infection.
Understanding host-parasite relationship: the immune central nervous system microenvironment and its effect on brain infections.
A nucleotide sugar transporter involved in glycosylation of the <i>Toxoplasma</i> tissue cyst wall is required for efficient persistence of bradyzoites.
<i>Toxoplasma</i> exports dense granule proteins beyond the vacuole to the host cell nucleus and rewires the host genome expression.
Parasite epigenetics and immune evasion: lessons from budding yeast.
Cysteine proteases in protozoan parasites.
<i>Toxoplasma gondii</i> infection and schizophrenia: an inter-kingdom communication perspective.
Role of prostaglandin F <sub>2</sub> $\alpha$ production in lipid bodies from <i>Leishmania infantum</i> chagasi: insights on virulence.
Sphingolipid degradation by <i>Leishmania major</i> is required for its resistance to acidic pH in the mammalian host.
Parasite regulation by host hormones: an old mechanism of host exploitation?
Host cell signalling and leishmania mechanisms of evasion.
Immunity to gastrointestinal microparasites of fish.
<i>Leishmania major</i> parasite stage-dependent host cell invasion and immune evasion.
The genetics of <i>Leishmania</i> virulence.
Quantification of Nitric Oxide and Reactive Oxygen Species in <i>Leishmania</i> -infected J774A.1 Macrophages as a Response to the in vitro treatment with a Natural Product Compound.
Histone deacetylase inhibitor MS-275 augments expression of a subset of IFN- $\gamma$ -regulated genes in <i>Toxoplasma gondii</i> -infected macrophages but does not improve parasite control.
Genetic mapping identifies novel highly protective antigens for an apicomplexan parasite.
In silico characterization of multiple genes encoding the GP63 virulence protein from <i>Leishmania braziliensis</i> : identification of sources of variation and putative roles in immune evasion.
The kidney form of <i>Trypanosoma musculi</i> : A distinct stage in the life cycle?

Ecto-nucleotidase activities of promastigotes from <i>Leishmania (Viannia) braziliensis</i> relates to parasite infectivity and disease clinical outcome.
Exposure of phosphatidylserine on <i>Leishmania amazonensis</i> isolates is associated with diffuse cutaneous leishmaniasis and parasite infectivity.
Effects of <i>Pseudoloma neurophilia</i> infection on the brain transcriptome in zebrafish ( <i>Danio rerio</i> ).
Protective immune response against cutaneous leishmaniasis by prime/booster immunization regimens with vaccinia virus recombinants expressing <i>Leishmania infantum</i> p36/LACK and IL-12 in combination with purified p36.
Why are male malaria parasites in such a rush?: Sex-specific evolution and host-parasite interactions.
Proximity biotinylation reveals novel secreted dense granule proteins of <i>Toxoplasma gondii</i> bradyzoites.
Crosstalk of PD-1 signaling with the SIRT1/FOXO-1 axis during the progression of visceral leishmaniasis.
[Immune evasion molecules of <i>Toxoplasma gondii</i> ].
Immune Response of Amebiasis and Immune Evasion by <i>Entamoeba histolytica</i> .
The role of extracellular vesicles in <i>Plasmodium</i> and other protozoan parasites.
LCCL proteins of apicomplexan parasites.
<i>Trichomonas vaginalis</i> : Pathogenesis, Symbiont Interactions, and Host Cell Immune Responses.
Genome-wide identification of molecular mimicry candidates in parasites.
<i>Leishmania donovani</i> infection suppresses Allograft Inflammatory Factor-1 in monocytes and macrophages to inhibit inflammatory responses.
Parasitism and chromosome dynamics in protozoan parasites: is there a connection?
IL-12p70 production by <i>Leishmania</i> major-harboring human dendritic cells is a CD40/CD40 ligand-dependent process.
<i>Leishmania</i> survival in the macrophage: where the ends justify the means.
<i>Toxoplasma gondii</i> Recruits Factor H and C4b-Binding Protein to Mediate Resistance to Serum Killing and Promote Parasite Persistence in vivo.
[Functional roles of macrophage migration inhibitory factor in anti-parasitic diseases].
Rhoptry and Dense Granule Secreted Effectors Regulate CD8(+) T Cell Recognition of <i>Toxoplasma gondii</i> Infected Host Cells.
The roles of intramembrane proteases in protozoan parasites.
Parasite-mediated selection drives an immunogenetic trade-off in plains zebras ( <i>Equus quagga</i> ).
Parasite immune evasion: a momentous molecular war.
Transcriptional Profiling Suggests T Cells Cluster around Neurons Injected with <i>Toxoplasma gondii</i> Proteins.
Host persistence: exploitation of anti-inflammatory pathways by <i>Toxoplasma gondii</i> .
Interplay Between <i>Toxoplasma gondii</i> , Autophagy, and Autophagy Proteins.
Surface antigens and potential virulence factors from parasites detected by comparative genomics of perfect amino acid repeats.
Evolution and diversity of the EMA families of the divergent equid parasites, <i>Theileria equi</i> and <i>T. haneyi</i> .
Immune evasion and the evolution of molecular mimicry in parasites.
Integration of DNA Repair, Antigenic Variation, Cytoadhesion, and Chance in <i>Babesia</i> Survival: A Perspective.
Fc receptors and immunity to parasites.
Innate immunity against <i>Leishmania</i> infections.

Evolution and genetic diversity of Theileria.
Sequence variation and structural conservation allows development of novel function and immune evasion in parasite surface protein families.
Novel roles of dense granule protein 12 (GRA12) in <i>Toxoplasma gondii</i> infection.
The kinetics of cellular and humoral immune responses of common carp to presporogonic development of the myxozoan <i>Sphaerospora molnari</i> .
How to get away with murder: The multiple strategies employed by pathogenic protozoa to avoid complement killing.
Secreted virulence factors and immune evasion in visceral leishmaniasis.
A novel high throughput invasion screen identifies host actin regulators required for efficient cell entry by <i>Toxoplasma gondii</i> .
Host-parasite interactions: resist or tolerate but never stop running.
Inflammasomes and Leishmania: in good times or bad, in sickness or in health.
Surface carbohydrate composition of a tapeworm in its consecutive intermediate hosts: individual variation and fitness consequences.
Neutrophil interactions with the sexually transmitted parasite <i>Trichomonas vaginalis</i> : implications for immunity and pathogenesis.
[Immunopathogenesis of Leishmania infections].
Prior infections or defence priming: what determines the risk of trematode infections in amphipod hosts?
The intermediate host immune response in cystic echinococcosis.
Polarisation of human macrophages towards an M1 subtype triggered by an atypical Brazilian strain of <i>Toxoplasma gondii</i> results in a reduction in parasite burden.
<i>Trypanosoma carassii</i> calreticulin binds host complement component C1q and inhibits classical complement pathway-mediated lysis.
Prostaglandin E2/leukotriene B4 balance induced by <i>Lutzomyia longipalpis</i> saliva favors <i>Leishmania infantum</i> infection.
<i>Toxoplasma gondii</i> -induced immune suppression by human peripheral blood monocytes: role of gamma interferon.
Regulation of intrinsic apoptosis in cycloheximide-treated macrophages by the Sichuan human strain of Chinese <i>Leishmania</i> isolates.
Coevolution of virulence and immunosuppression in multiple infections.
Advances in the Development of Anti- <i>Toxoplasma gondii</i> Vaccines: Challenges, Opportunities, and Perspectives.
Can myxosporean parasites compromise fish and amphibian reproduction?
<i>Toxoplasma</i> Effectors Targeting Host Signaling and Transcription.
Differential phenoloxidase activity between native and invasive gammarids infected by local acanthocephalans: differential immunosuppression?
Pathology of CNS parasitic infections.
Modulation of early $\beta$ -defensin-2 production as a mechanism developed by type I <i>Toxoplasma gondii</i> to evade human intestinal immunity.
<i>Leishmania</i> -induced inactivation of the macrophage transcription factor AP-1 is mediated by the parasite metalloprotease GP63.
Influence of parasitic life style on the patterns of codon usage and base frequencies of <i>Ancylostoma</i> and <i>Necator</i> species.
Impact of <i>Leishmania</i> metalloprotease GP63 on macrophage signaling.

The evolution of parasite recognition genes in the innate immune system: purifying selection on <i>Drosophila melanogaster</i> peptidoglycan recognition proteins.
The immunomodulatory potential of the arylmethylaminosteroid sc1o.
Mechanisms of adaptation in the intestinal parasite <i>Giardia lamblia</i> . [ <i>Giardia</i> and giardiasis].
Effect of malaria parasite shape on its alignment at erythrocyte membrane.
Evasion of innate immunity by parasitic protozoa.
<i>Leishmania infantum</i> modulates host macrophage mitochondrial metabolism by hijacking the SIRT1-AMPK axis.
[Progress of researches on the involvement of indoleamine 2, 3-dioxygenase in regulation of parasite-host immune interactions].
Immune Response to the Enteric Parasite <i>Entamoeba histolytica</i> .
Metabolomic Reprogramming of C57BL/6-Macrophages during Early Infection with <i>L. amazonensis</i> .
Lipoxins as an immune-escape mechanism.
Cytokines in the generation and function of regulatory T cell subsets in leishmaniasis.
Activation of phosphatidylinositol 3-kinase/Akt and impairment of nuclear factor-kappaB: molecular mechanisms behind the arrested maturation/activation state of <i>Leishmania infantum</i> -infected dendritic cells.
Adhesion molecules and other secreted host-interaction determinants in Apicomplexa: insights from comparative genomics.
Parasite-mediated steps in immune response failure during primary <i>Theileria annulata</i> infection.
A <i>Leishmania</i> ortholog of macrophage migration inhibitory factor modulates host macrophage responses.
<i>Ichthyophthirius multifiliis</i> Fouquet and Ichthyophthiriosis in Freshwater Teleosts.
Parasite protein phosphatases: biological function, virulence, and host immune evasion.
The role of SIRT1 in the process of <i>Toxoplasma gondii</i> infection of RAW 264.7 macrophages.
<i>Leishmania</i> lipophosphoglycan activates the transcription factor activating protein 1 in J774A.1 macrophages through the extracellular signal-related kinase (ERK) and p38 mitogen-activated protein kinase.
<i>Toxoplasma gondii</i> Parasitophorous Vacuole Membrane-Associated Dense Granule Proteins Orchestrate Chronic Infection and GRA12 Underpins Resistance to Host Gamma Interferon.
Macrophages behaving badly: infected cells and subversion of immune responses to <i>Theileria annulata</i> .
Host-Parasite Interactions: Regulation of <i>Leishmania</i> Infection in Sand Fly.
Brazilian strains of <i>Toxoplasma gondii</i> are controlled by azithromycin and modulate cytokine production in human placental explants.
Effect of different serine protease inhibitors in validating the 115 kDa <i>Leishmania donovani</i> secretory serine protease as chemotherapeutic target.
Variation in <i>Leishmania</i> chemokine suppression driven by diversification of the GP63 virulence factor.
<i>Babesia microti</i> Immunoreactive Rhoptry-Associated Protein-1 Paralogs Are Ancestral Members of the Piroplasmid-Confined RAP-1 Family.
Influence of apoptosis on the cutaneous and peripheral lymph node inflammatory response in dogs with visceral leishmaniasis.
A comparative study between excretory/secretory and autoclaved vaccines against RH strain of <i>Toxoplasma gondii</i> in murine models.
Propulsive cell entry diverts pathogens from immune degradation by remodeling the phagocytic synapse.

Immunization With a Live-Attenuated RH:ΔNPT1 Strain of <i>Toxoplasma gondii</i> Induces Strong Protective Immunity Against Toxoplasmosis in Mice.
Regulation of the immune system in metazoan parasite infections.
Mast cells at the host-pathogen interface: host-protection versus immune evasion in leishmaniasis.
Analysis of mixed parasite populations of <i>Theileria sergenti</i> using cDNA probes encoding a major piroplasm surface protein.
The oxygen reduction pathway and heat shock stress response are both required for <i>Entamoeba histolytica</i> pathogenicity.
Aberrant receptor-mediated endocytosis of <i>Schistosoma mansoni</i> glycoproteins on host lipoproteins.
Alice in microbes' land: adaptations and counter-adaptations of vector-borne parasitic protozoa and their hosts.
In vitro leukocyte response of three-spined sticklebacks ( <i>Gasterosteus aculeatus</i> ) to helminth parasite antigens.
Directing differentiation in <i>Theileria annulata</i> : old methods and new possibilities for control of apicomplexan parasites.
<i>Leishmania</i> virulence factors: focus on the metalloprotease GP63.
Zoite migration during infection: parasite adaptation to host defences.
Plastic and micro-evolutionary responses of a nematode to the host immune environment.
Interplay between Attenuation- and Virulence-Factors of <i>Babesia bovis</i> and Their Contribution to the Establishment of Persistent Infections in Cattle.
Contribution of the TIM-3/Gal-9 immune checkpoint to tropical parasitic diseases.
Biological selection of variant-specific surface proteins in <i>Giardia lamblia</i> .
Host cell invasion by the apicomplexans: the significance of microneme protein proteolysis.
Disruption of antigenic variation is crucial for effective parasite vaccine.
Host-directed antileishmanial interventions: Harvesting unripe fruits to reach fruition.
Natural insect host-parasite systems show immune priming and specificity: puzzles to be solved.
To React or Not to React: The Dilemma of Fish Immune Systems Facing Myxozoan Infections.
<i>Toxoplasma gondii</i> Cathepsin C1 inhibits NF-κB signalling through the positive regulation of the HIF-1α/EPO axis.
<i>Leishmania donovani</i> infection of a susceptible host results in apoptosis of Th1-like cells: rescue of anti-leishmanial CMI by providing Th1-specific bystander costimulation.
Immune Evasion Mechanisms of <i>Entamoeba histolytica</i> : Progression to Disease.
Advances in understanding red blood cell modifications by <i>Babesia</i> .
Epigenetics in the early divergent eukaryotic <i>Giardia duodenalis</i> : An update.
<i>Leishmania donovani</i> evades Caspase 1 dependent host defense mechanism during infection.
RNA-seq analysis of early enteromyxosis in turbot ( <i>Scophthalmus maximus</i> ): new insights into parasite invasion and immune evasion strategies.
TLR9/MyD88/TRIF signaling activates host immune inhibitory CD200 in <i>Leishmania</i> infection.
<i>Toxoplasma</i> Effector Recruits the Mi-2/NuRD Complex to Repress STAT1 Transcription and Block IFN-γ-Dependent Gene Expression.
Structural characterization of the bradyzoite surface antigen (BSR4) from <i>Toxoplasma gondii</i> , a unique addition to the surface antigen glycoprotein 1-related superfamily.
Interaction between parasite-encoded JAB1/CSN5 and macrophage migration inhibitory factor proteins attenuates its proinflammatory function.
<i>Leishmania donovani</i> : impairment of the cellular immune response against recombinant ornithine decarboxylase protein as a possible evasion strategy of <i>Leishmania</i> in visceral leishmaniasis.

Reciprocal CD40 signals through p38MAPK and ERK-1/2 induce counteracting immune responses.
The soluble fraction of <i>Neospora caninum</i> treated with PI-PLC is dominated by NcSRS29B and NcSRS29C.
<i>Toxoplasma gondii</i> inhibits granzyme B-mediated apoptosis by the inhibition of granzyme B function in host cells.
Starch Branching Enzyme 1 Is Important for Amylopectin Synthesis and Cyst Reactivation in <i>Toxoplasma gondii</i> .
CRISPR screen to determine the in vivo fitness of <i>Toxoplasma</i> genes.
Congenital Transmission of Apicomplexan Parasites: A Review.
<i>Leishmania donovani</i> infection differentially regulates small G-proteins.
Ecological and histopathological aspects of <i>Didymodictylus</i> sp. (Trematoda: Didymozoidae) parasite of the dusky grouper, <i>Epinephelus marginatus</i> (Osteichthyes: Serranidae), from the western Mediterranean Sea.
Genomic Analysis of Colombian <i>Leishmania panamensis</i> strains with different level of virulence.
Molecular characterization of NCLIV_011700 of <i>Neospora caninum</i> , a low sequence identity rhostry protein.
Metallic Nanoparticles: A New Frontier in the Fight Against Leishmaniasis.
Inhibition of dendritic cell apoptosis by <i>Leishmania mexicana</i> amastigotes.
Sand fly saliva enhances <i>Leishmania amazonensis</i> infection by modulating interleukin-10 production.
Homologous Recombination in Protozoan Parasites and Recombinase Inhibitors.
Cyclooxygenase (COX)-2 modulates <i>Toxoplasma gondii</i> infection, immune response and lipid droplets formation in human trophoblast cells and villous explants.
[Characteristics of the immune response in protozoan infections].
Phylogenetic evidence for recombination in SAG5 locus in <i>Toxoplasma gondii</i> .
Prevalence and histopathology of the parasitic barnacle, <i>Sacculina carcini</i> in shore crabs, <i>Carcinus maenas</i> .
<i>Leishmania donovani</i> promastigotes evade the activation of mitogen-activated protein kinases p38, c-Jun N-terminal kinase, and extracellular signal-regulated kinase-1/2 during infection of naive macrophages.
<i>Leishmania</i> Uses Mincle to Target an Inhibitory ITAM Signaling Pathway in Dendritic Cells that Dampens Adaptive Immunity to Infection.
<i>Leishmania</i> evades host immunity by inhibiting antigen cross-presentation through direct cleavage of the SNARE VAMP8.
Visiting Molecular Mimicry Once More: Pathogenicity, Virulence, and Autoimmunity.
Functional Analysis of the Role of <i>Toxoplasma gondii</i> Nucleoside Triphosphate Hydrolases I and II in Acute Mouse Virulence and Immune Suppression.
Molecular communication between the monogenea and fish immune system.
African trypanosomosis: from immune escape and immunopathology to immune intervention.
Proteophosphoglycan, a major secreted product of intracellular <i>Leishmania mexicana</i> amastigotes, is a poor B-cell antigen and does not elicit a specific conventional CD4+ T-cell response.
Molecular mechanisms of hookworm disease: stealth, virulence, and vaccines.
Immunomodulatory effects of helminths and protozoa in multiple sclerosis and experimental autoimmune encephalomyelitis.
PI3K signaling in <i>Leishmania</i> infections.
Flagellum assembly and function during the <i>Leishmania</i> life cycle.
Potential biomarkers of immune protection in human leishmaniasis.
Major Surface Antigens in Zoonotic Babesia.

Crosstalk between purinergic receptors and lipid mediators in leishmaniasis.
Setaria digitata secreted filarial lipids modulate IL-12 signaling through JAK-STAT pathway leading to the development of Th1 response.
Ligand-binding renders the 160 kDa Trypanosoma cruzi complement regulatory protein susceptible to proteolytic cleavage.
New cell motility model observed in parasitic cnidarian Sphaerospora molnari (Myxozoa:Myxosporea) blood stages in fish.
Peroxisome proliferator-activated receptor (PPAR): balance for survival in parasitic infections.
The dynamics of neutrophils in zebrafish (Danio rerio) during infection with the parasite Ichthyophthirius multifiliis.
[Reaction of bovine endothelial cells in vitro to coccidia (Eimeria bovis, Toxoplasma gondii, Neospora caninum) infections as the expression of a non-adaptive immune response].
A genome-wide analysis of coatamer protein (COP) subunits of apicomplexan parasites and their evolutionary relationships.
Leishmania amazonensis impairs DC function by inhibiting CD40 expression via A2B adenosine receptor activation.
Shared epigenetic mechanisms control virulence factors in protozoan parasites.
Genome-Wide Identification and Evolutionary Analysis of Sarcocystis neurona Protein Kinases.
Thioredoxin reductase from Toxoplasma gondii: an essential virulence effector with antioxidant function.
Myxozoan Adhesion and Virulence: Ceratonyx shasta on the Move.
Representational difference analysis identifies specific genes in the interaction of Giardia duodenalis with the murine intestinal epithelial cell line, IEC-6.
Adaptive amino acid composition in collagens of parasitic nematodes.
Principles of ecological immunology.
Transcriptomic immune response of Tenebrio molitor pupae to parasitization by Scleroderma guani.
Regulation of antigenic variation in Giardia lamblia.
Characterization of the B-cell immune response elicited in BALB/c mice challenged with Neospora caninum tachyzoites.
A short note on predicted motifs in the Thaparocleidus wallagonius genome.
Gene Expression Profiling of Neospora caninum in Bovine Macrophages Reveals Differences Between Isolates Associated With Key Parasite Functions.
Leishmania amazonensis downregulates macrophage iNOS expression via Histone Deacetylase 1 (HDAC1): a novel parasite evasion mechanism.
Immune responses in kala-azar.
The role of myeloid-derived suppressor cells in chronic infectious diseases and the current methodology available for their study.
Leishmania amazonensis-Induced cAMP Triggered by Adenosine A(2B) Receptor Is Important to Inhibit Dendritic Cell Activation and Evade Immune Response in Infected Mice.
African Trypanosomiasis-Associated Anemia: The Contribution of the Interplay between Parasites and the Mononuclear Phagocyte System.
Surface Glycans: A Therapeutic Opportunity for Kinetoplastid Diseases.
Hemoglobin uptake and utilization by human protozoan parasites: a review.
Dynamic development of parasitophorous vacuole of Eimeria tenella transfected with the yellow fluorescent protein gene fused to different signal sequences from apicomplexan parasites.
The Toxoplasma gondii cyst wall protein CST1 is critical for cyst wall integrity and promotes bradyzoite persistence.



Regulation of host cell function by glycosylphosphatidylinositols of the parasitic protozoa.
Population, genetic, and antigenic diversity of the apicomplexan <i>Eimeria tenella</i> and their relevance to vaccine development.
Impact of lipoxin-mediated regulation on immune response to infectious disease.
What Do We Know about Surface Proteins of Chicken Parasites <i>Eimeria</i> ?
Cell-to-cell transfer of <i>Leishmania amazonensis</i> amastigotes is mediated by immunomodulatory LAMP-rich parasitophorous extrusions.
Reprint of: The non-mammalian MIF superfamily.
Subversion of the B-cell compartment during parasitic, bacterial, and viral infections.
Genome-wide analysis of gene expression and protein secretion of <i>Babesia canis</i> during virulent infection identifies potential pathogenicity factors.
Molecular variation and diversity in candidate vaccine antigens from <i>Babesia</i> .
Harnessing Immune Evasion Strategy of Lymphatic Filariae: A Therapeutic Approach against Inflammatory and Infective Pathology.
[Research progress on the mechanisms of antigenic variation in <i>Giardia lamblia</i> ].
Genome-Wide Bimolecular Fluorescence Complementation-Based Proteomic Analysis of <i>Toxoplasma gondii</i> ROP18's Human Interactome Shows Its Key Role in Regulation of Cell Immunity and Apoptosis.
Comparative genomic analysis and phylogenetic position of <i>Theileria equi</i> .
Immune modulation by fish kinetoplastid parasites: a role for nitric oxide.
<i>Toxoplasma gondii</i> Type-I ROP18 Targeting Human E3 Ligase TRIM21 for Immune Escape.
CD4(+) T Cell-Dependent Macrophage Activation Modulates Sustained PS Exposure on Intracellular Amastigotes of <i>Leishmania amazonensis</i> .
Immune evasion by Trypanosomatidae: normal aggregated immunoglobulin protects against lysis by the alternative complement pathway.
Reprint of "fish immunity to scuticociliate parasites".
Method for Isolation of Myxozoan Proliferative Stages from Fish at High Yield and Purity: An Essential Prerequisite for In Vitro, In Vivo and Genomics-Based Research Developments.
Evidence of gene conversion in genes encoding the Gal/GalNac lectin complex of <i>Entamoeba</i> .
Update on Lymphatic Filarial Infections.
Ichthyophthiriasis in carp <i>Cyprinus carpio</i> : infectivity of trophonts prematurely exiting both the immune and non-immune host.
Detection in human sera of IgG, IgM and IgA to excreted/secreted antigens from <i>Toxoplasma gondii</i> by use of dot-ELISA and immunoblot assay.
Genomic resources for a unique, low-virulence <i>Babesia</i> taxon from China.
SHP-1 Regulates Antigen Cross-Presentation and Is Exploited by <i>Leishmania</i> to Evade Immunity.
<i>Leishmania donovani</i> affects antigen presentation of macrophage by disrupting lipid rafts.
Interdependencies between Toll-like receptors in <i>Leishmania</i> infection.
RAP-1a is the main rhoptry-associated-protein-1 (RAP-1) recognized during infection with <i>Babesia</i> sp. BQ1 (Lintan) ( <i>B. motasi</i> -like phylogenetic group), a pathogen of sheep in China.
<i>Leishmania donovani</i> inhibits ferroportin translation by modulating FBXL5-IRP2 axis for its growth within host macrophages.
Vaccination against babesiosis using recombinant GPI-anchored proteins.
Trichomoniasis immunity and the involvement of the purinergic signaling.
The Mechanism of Facultative Intracellular Parasitism of <i>Brucella</i> .
<i>Leishmania braziliensis</i> : Strain-Specific Modulation of Phagosome Maturation.
Antigenic variation in <i>Giardia lamblia</i> is regulated by RNA interference.

Differences in cap formation between invasive <i>Entamoeba histolytica</i> and non-invasive <i>Entamoeba dispar</i> .
Clinical pleiomorphism in human leishmaniasis, with special mention of asymptomatic infection.
Molecular technology and antigenic variation among intraerythrocytic hemoparasites: do we see reality?
Human babesiosis: Indication of a molecular mimicry between thrombospondin domains from a novel <i>Babesia microti</i> BmP53 protein and host platelets molecules.
Harbouring in the brain: A focus on immune evasion mechanisms and their deleterious effects in malaria and human African trypanosomiasis.
The NF-kappaB signaling pathway: immune evasion and immunoregulation during toxoplasmosis.
(1)H, (13)C and (15)N backbone and side-chain resonance assignments of $\Delta\Delta$ BmSA1, the surface antigen of <i>Babesia microti</i> .
<i>Leishmania infantum</i> Exoproducts Inhibit Human Invariant NKT Cell Expansion and Activation.
Babesiosis: persistence in the face of adversity.
<i>Lepeophtheirus salmonis</i> : characterization of prostaglandin E(2) in secretory products of the salmon louse by RP-HPLC and mass spectrometry.
The complex fibronectin-- <i>Trichomonas vaginalis</i> interactions and Trichomonosis.
Surface antigenic variation in <i>Giardia lamblia</i> .
Regulation of host epithelial responses to <i>Cryptosporidium</i> infection by microRNAs.
Early stages of infection of three-spined stickleback ( <i>Gasterosteus aculeatus</i> ) with the cestode <i>Schistocephalus solidus</i> .
G protein signaling in the parasite <i>Entamoeba histolytica</i> .
Mechanisms for a novel immune evasion strategy in the scabies mite <i>Sarcoptes scabiei</i> : a multigene family of inactivated serine proteases.
Evidence of high-efficiency cross fertilization in <i>Eimeria acervulina</i> revealed using two lines of transgenic parasites.
Sex differences in the protection of host immune systems by a polyembryonic parasitoid.
Apoptosis and apoptotic mimicry: the <i>Leishmania</i> connection.
Proteomic analysis in <i>Giardia duodenalis</i> yields insights into strain virulence and antigenic variation.
A <i>Leishmania infantum</i> cytosolic trypanothionein activates B cells to secrete interleukin-10 and specific immunoglobulin.
<i>Leishmania donovani</i> : dynamics of <i>L. donovani</i> evasion of innate immune cell attack due to malnutrition in visceral leishmaniasis.
Microvesicles and exosomes as vehicles between protozoan and host cell communication.
Emerging perspectives in the research of bovine babesiosis and anaplasmosis.
Modulation of Host Immune Response during <i>Leishmania infantum</i> Natural Infection: A Whole-Transcriptome Analysis of the Popliteal Lymph Nodes in Dogs.
Clinical and microbiological aspects of <i>Trichomonas vaginalis</i> .
A stochastic microscopic model for the dynamics of antigenic variation.
<i>Trichomonas vaginalis</i> Lipophosphoglycan Exploits Binding to Galectin-1 and -3 to Modulate Epithelial Immunity.
Identification and characterization of a cyclooxygenase-like enzyme from <i>Entamoeba histolytica</i> .
Cryptosporidial Infection Suppresses Intestinal Epithelial Cell MAPK Signaling Impairing Host Anti-Parasitic Defense.
Identification of variant-specific surface proteins in <i>Giardia muris</i> trophozoites.
Genetic Diversity of Serine Protease Inhibitors in Myxozoan (Cnidaria, Myxozoa) Fish Parasites.

Reciprocal regulation of protein kinase C isoforms results in differential cellular responsiveness.
Functional Characterization of an Interferon Gamma Receptor-Like Protein on <i>Entamoeba histolytica</i> .
Modulation of systemic and mucosal immune responses of <i>Catla catla</i> (Hamilton, 1822) experimentally challenged with gill monogeneans.
RepSeq--a database of amino acid repeats present in lower eukaryotic pathogens.
Recent Development in Identification of Potential Novel Therapeutic Targets Against Trypanosomatids.
Extracellular vesicles in infectious diseases caused by protozoan parasites in buffaloes.
Genetic basis for GPI-anchor merozoite surface antigen polymorphism of <i>Babesia</i> and resulting antigenic diversity.
Polynucleotidyl transfer reactions in site-specific DNA recombination.
<i>Babesia microti</i> : Pathogen Genomics, Genetic Variability, Immunodominant Antigens, and Pathogenesis.
Transition-state-guided drug design for treatment of parasitic neglected tropical diseases.
CD40-modulated dual-specificity phosphatases MAPK phosphatase (MKP)-1 and MKP-3 reciprocally regulate <i>Leishmania major</i> infection.
Eukaryote-conserved histone post-translational modification landscape in <i>Giardia duodenalis</i> revealed by mass spectrometry.
An <i>Entamoeba histolytica</i> rhomboid protease with atypical specificity cleaves a surface lectin involved in phagocytosis and immune evasion.
A Devil of a Transmissible Cancer.
Immunoproteomic analysis of the protein repertoire of unsporulated <i>Eimeria tenella</i> oocysts.
Relationships between leukocytes and Hepatozoon spp. In green frogs, <i>Rana clamitans</i> .
Inoculating <i>Anopheles gambiae</i> Mosquitoes with Beads to Induce and Measure the Melanization Immune Response.
Selection of diversity at putative glycosylation sites in the immunodominant merozoite/piroplasm surface antigen of <i>Theileria</i> parasites.
Adenosine and immune imbalance in visceral leishmaniasis: the possible role of ectonucleotidases.
Reproductive failures associated with <i>Trypanosoma</i> ( <i>Duttonella</i> ) <i>vivax</i> .
Immunology of anaphylaxis: lessons from murine models.
Immunization with <i>Neospora caninum</i> profilin induces limited protection and a regulatory T-cell response in mice.
Suppression of host PTEN gene expression for <i>Leishmania donovani</i> survival in Indian visceral leishmaniasis.
Recent developments in the molecular, biochemical and functional characterization of GPI8 and the GPI-anchoring mechanism [review].
Sequestration of host-CD59 as potential immune evasion strategy of <i>Trichomonas vaginalis</i> .
Role of scuticociliate proteinases in infection success in turbot, <i>Psetta maxima</i> (L.).
Immunoglobulin subclass responses of wild brown rats to <i>Sarcocystis singaporensis</i> .
Modulation of the host Th1 immune response in pigeon protozoal encephalitis caused by <i>Sarcocystis calchasi</i> .
The rhoptry proteins ROP18 and ROP5 mediate <i>Toxoplasma gondii</i> evasion of the murine, but not the human, interferon-gamma response.
A perspective on clonal phenotypic (antigenic) variation in protozoan parasites.
<i>Leishmania infantum-chagasi</i> activates SHP-1 and reduces NFAT5/TonEBP activity in the mouse kidney inner medulla.
Lateral Gene Transfer in the Adaptation of the Anaerobic Parasite <i>Blastocystis</i> to the Gut.

Apoptotic killing and phagocytosis of host cells by the parasite <i>Entamoeba histolytica</i> .
Alveolar echinococcosis: what triggers emergence in North America, Central Europe and Asia?
<i>Theileria</i> hijacks JNK2 into a complex with the macroschizont GPI (GlycosylPhosphatidyInositol)-anchored surface protein p104.
Interactions between the Parasite <i>Philasterides dicentrarchi</i> and the Immune System of the Turbot <i>Scophthalmus maximus</i> . A Transcriptomic Analysis.
<i>Neospora caninum</i> Evades Immunity via Inducing Host Cell Mitophagy to Inhibit Production of Proinflammatory Cytokines in a ROS-Dependent Manner.
O-deGlcNAcylation is required for <i>Entamoeba histolytica</i> -induced HepG2 cell death.
Cutaneous myxidiosis in European eel, <i>Anguilla anguilla</i> (Linnaeus, 1758): histopathology, histochemistry and laminin immunohistochemistry.
Identification and expression of macrophage migration inhibitory factor in <i>Sarcoptes scabiei</i> .
Sneaking Out for Happy Hour: Yeast-Based Approaches to Explore and Modulate Immune Response and Immune Evasion.
Making the cut: central roles of intramembrane proteolysis in pathogenic microorganisms.
Discovering a vaccine against neosporosis using computers: is it feasible?
<i>Entamoeba histolytica</i> induces intestinal cathelicidins but is resistant to cathelicidin-mediated killing.
Comprehensive characterization of Cysteine-rich protein-coding genes of <i>Giardia lamblia</i> and their role during antigenic variation.
<i>Entamoeba histolytica</i> and <i>E. dispar</i> trophozoites in the liver of hamsters: in vivo binding of antibodies and complement.
Antimony-Resistant <i>Leishmania donovani</i> Exploits miR-466i To Deactivate Host MyD88 for Regulating IL-10/IL-12 Levels during Early Hours of Infection.
Secretion of multilamellar whorls by <i>Eimeria tenella</i> zoites.
Evolutionary dynamics of invasion and escape.
Serum concentration of sialic acids in naturally occurring ovine babesiosis.
Modulation of Cholesterol Pathways in Human Macrophages Infected by Clinical Isolates of <i>Leishmania infantum</i> .
Emergence of new types of <i>Theileria orientalis</i> in Australian cattle and possible cause of theileriosis outbreaks.
The Price equation framework to study disease within-host evolution.
Dinoflagellate symbionts escape vomocytosis by host cell immune suppression.
Inhibition of CD40-induced N-Ras activation reduces leishmania major infection.
Fulminant amebic colitis: An unusual postoperative complication of intraabdominal malignancy.
Cholesterol depletion associated with <i>Leishmania major</i> infection alters macrophage CD40 signalosome composition and effector function.
Immunoproteomic analysis of the sporozoite antigens of <i>Eimeria necatrix</i> .
Proteomic analysis of <i>Eimeria acervulina</i> sporozoite proteins interaction with duodenal epithelial cells by shotgun LC-MS/MS.
Macrophage migration inhibitory factor of <i>Thelazia callipaeda</i> induces M2-like macrophage polarization through TLR4-mediated activation of the PI3K-Akt pathway.
<i>Tritrichomonas foetus</i> displays classical detergent-resistant membrane microdomains on its cell surface.
<i>Babesia microti</i> -induced fulminant sepsis in an immunocompromised host: A case report and the case-specific literature review.
Evaluation of cellular and humoral systemic immune response against <i>Giardia duodenalis</i> infection in cattle.

Oxidative stress response in <i>Paracoccidioides brasiliensis</i> .
Coexistence of sense and anti-sense mRNAs of variant surface protein in <i>Giardia lamblia</i> trophozoites.
On the age of leprosy.
The immunology of susceptibility and resistance to scabies.
<i>Eimeria bovis</i> modulates adhesion molecule gene transcription in and PMN adhesion to infected bovine endothelial cells.
In silico and phylogenetic analyses of partial BbRAP-1, BbCP2, BbSBP-4 and Bb $\beta$ TUB gene sequences of <i>Babesia bovis</i> isolates from cattle in South Africa.
The basic isoform of profilin in pathogenic <i>Entamoeba histolytica</i> . cDNA cloning, heterologous expression, and actin-binding properties.
T-cell receptor beta-chain binding to enterotoxin superantigens.
Apoptosis of macrophages induced by <i>Trichomonas vaginalis</i> through the phosphorylation of p38 mitogen-activated protein kinase that locates at downstream of mitochondria-dependent caspase activation.
Recombinant immunoglobulin A: powerful tools for fundamental and applied research.
Calpain-dependent cleavage of SHP-1 and SHP-2 is involved in the dephosphorylation of Jurkat T cells induced by <i>Entamoeba histolytica</i> .
Immunology of scabies and translational outcomes: identifying the missing links.
<i>Demodex canis</i> targets TLRs to evade host immunity and induce canine demodicosis.
Effects of the histiophagous ciliate <i>Philasterides dicentrarchi</i> on turbot phagocyte responses.
Global haplotype distribution of <i>Babesia ovis</i> inferred by 18S rRNA sequences; a phylogeographical systematic review.
An African perspective on mucosal immunity and HIV-1.
An overview on kinetoplastid paraflagellar rod.
Microparticles and infectious diseases.
Intrinsically disordered proteins (IDPs) in trypanosomatids.
<i>Sarcocystis neurona</i> major surface antigen gene 1 (SAG1) shows evidence of having evolved under positive selection pressure.
[Expression of CD47 and its ligands in pregnant mice infected with <i>Toxoplasma gondii</i> during pregnancy].
Quantitative PCR-based genome size estimation of the astigmatid mites <i>Sarcoptes scabiei</i> , <i>Psoroptes ovis</i> and <i>Dermatophagoides pteronyssinus</i> .
Guest list or black list: heritable small RNAs as immunogenic memories.
Influence of experimental <i>Anaplasma marginale</i> infection and splenectomy on NTPDase and 5' nucleotidase activities in platelets of cattle.
Expression and localization of grass carp <i>pkc-<math>\theta</math></i> (protein kinase C theta) gene after its activation.

Table A2-53, Cluster 52

Cluster 52 focuses on <i>Streptococcus pneumoniae</i> , emphasizing pathogenicity, virulence, and immune system evasion (366)
Oligopeptide Transporters of Nonencapsulated <i>Streptococcus pneumoniae</i> Regulate CbpAC and PspA Expression and Reduce Complement-Mediated Clearance.
Diverse Mechanisms of Protective Anti-Pneumococcal Antibodies.
Sugar-Coated Killer: Serotype 3 Pneumococcal Disease.
Molecular Evolution of Clinical Pathogenic Streptococci.

Streptococci and the complement system: interplay during infection, inflammation and autoimmunity.
Natural transformation and genome evolution in <i>Streptococcus pneumoniae</i> .
PspK of <i>Streptococcus pneumoniae</i> increases adherence to epithelial cells and enhances nasopharyngeal colonization.
<i>Streptococcus pneumoniae</i> Evades Host Cell Phagocytosis and Limits Host Mortality Through Its Cell Wall Anchoring Protein PfbA.
Pneumococcal Surface Protein A: A Promising Candidate for the Next Generation of Pneumococcal Vaccines.
Heterologous expression of Ralp3 in <i>Streptococcus pyogenes</i> M2 and M6 strains affects the virulence characteristics.
Recombination in <i>Streptococcus pneumoniae</i> Lineages Increase with Carriage Duration and Size of the Polysaccharide Capsule.
Clinical Relevance and Molecular Pathogenesis of the Emerging Serotypes 22F and 33F of <i>Streptococcus pneumoniae</i> in Spain.
Combat pneumococcal infections: adhesins as candidates for protein-based vaccine development.
The antimicrobial resistance profile of <i>Streptococcus pneumoniae</i> .
Non-typeable <i>Streptococcus pneumoniae</i> carriage isolates genetically similar to invasive and carriage isolates expressing capsular type 14 in Brazilian infants.
In Vivo Relationship between the Nano-Biomechanical Properties of Streptococcal Polysaccharide Capsules and Virulence Phenotype.
Two cases of serotypeable and non-serotypeable variants of <i>Streptococcus pneumoniae</i> detected simultaneously during invasive disease.
Immunity to <i>Streptococcus pneumoniae</i> : Factors affecting production and efficacy.
JMM Profile: <i>Streptococcus pneumoniae</i> : sugar-coated captain of the men of death.
Immunization with Pneumococcal Surface Protein K of Nonencapsulated <i>Streptococcus pneumoniae</i> Provides Protection in a Mouse Model of Colonization.
Responses of innate immune cells to group A <i>Streptococcus</i> .
Fisher scientific award lecture - the capsular polysaccharides of Group B <i>Streptococcus</i> and <i>Streptococcus suis</i> differently modulate bacterial interactions with dendritic cells.
Polyamine transporter in <i>Streptococcus pneumoniae</i> is essential for evading early innate immune responses in pneumococcal pneumonia.
Role of <i>Streptococcus pneumoniae</i> extracellular glycosidases in immune evasion.
<i>Streptococcus pyogenes</i> evades adaptive immunity through specific IgG glycan hydrolysis.
Variants of <i>Streptococcus pneumoniae</i> Serotype 14 from Papua New Guinea with the Potential to Be Mistyped and Escape Vaccine-Induced Protection.
The choline-binding proteins PspA, PspC, and LytA of <i>Streptococcus pneumoniae</i> and their interaction with human endothelial and red blood cells.
Pathogenicity and virulence of <i>Streptococcus pneumoniae</i> : Cutting to the chase on proteases.
Gene elements that regulate <i>Streptococcus pneumoniae</i> virulence and immunity evasion.
Surface-associated lipoprotein PpmA of <i>Streptococcus pneumoniae</i> is involved in colonization in a strain-specific manner.
Role for streptococcal collagen-like protein 1 in M1T1 group A <i>Streptococcus</i> resistance to neutrophil extracellular traps.
Parameters governing invasive disease propensity of non-M1 serotype group A streptococci.
High multiple carriage and emergence of <i>Streptococcus pneumoniae</i> vaccine serotype variants in Malawian children.

A Jack of All Trades: The Role of Pneumococcal Surface Protein A in the Pathogenesis of <i>Streptococcus pneumoniae</i> .
Targeted control of pneumolysin production by a mobile genetic element in <i>Streptococcus pneumoniae</i> .
Molecular pathogenesis of <i>Klebsiella pneumoniae</i> .
<i>Streptococcus pyogenes</i> nuclease A (SpnA) mediated virulence does not exclusively depend on nuclease activity.
T4 Pili Promote Colonization and Immune Evasion Phenotypes of Nonencapsulated M4 <i>Streptococcus pyogenes</i> .
Surface charge of <i>Streptococcus pneumoniae</i> predicts serotype distribution.
The Heme Transporter HtsABC of Group A <i>Streptococcus</i> Contributes to Virulence and Innate Immune Evasion in Murine Skin Infections.
Neutrophil responsiveness to IL-10 impairs clearance of <i>Streptococcus pneumoniae</i> from the lungs.
Changed Expression of Cytoskeleton Proteins During Lung Injury in a Mouse Model of <i>Streptococcus pneumoniae</i> Infection.
The Group A <i>Streptococcus</i> serotype M2 pilus plays a role in host cell adhesion and immune evasion.
Serotype-independent pneumococcal experimental vaccines that induce cellular as well as humoral immunity.
Genetic diversity of capsular polysaccharide biosynthesis in <i>Klebsiella pneumoniae</i> clinical isolates.
A chemokine-degrading extracellular protease made by group A <i>Streptococcus</i> alters pathogenesis by enhancing evasion of the innate immune response.
Stand-alone response regulators controlling global virulence networks in <i>streptococcus pyogenes</i> .
Transcription of the <i>Streptococcus pyogenes</i> hyaluronic acid capsule biosynthesis operon is regulated by previously unknown upstream elements.
Genetics, Structure, and Function of Group A Streptococcal Pili.
Drug treatment of pneumococcal pneumonia in the elderly.
Upregulation of capsule enables <i>Streptococcus pyogenes</i> to evade immune recognition by antigen-specific antibodies directed to the G-related alpha2-macroglobulin-binding protein GRAB located on the bacterial surface.
<i>Streptococcus pneumoniae</i> , le transformiste.
Nasal immunization of mice with <i>Lactobacillus casei</i> expressing the Pneumococcal Surface Protein A: induction of antibodies, complement deposition and partial protection against <i>Streptococcus pneumoniae</i> challenge.
A brief review on Group A <i>Streptococcus</i> pathogenesis and vaccine development.
Evidence for soft selective sweeps in the evolution of pneumococcal multidrug resistance and vaccine escape.
Structural determination of <i>Streptococcus pyogenes</i> M1 protein interactions with human immunoglobulin G using integrative structural biology.
Molecular epidemiology and serogroup 6 capsular gene evolution of pneumococcal carriage in a Japanese birth cohort study.
Competence in <i>Streptococcus pneumoniae</i> and Close Commensal Relatives: Mechanisms and Implications.
<i>Streptococcus suis</i> serotype 2 strains can induce the formation of neutrophil extracellular traps and evade trapping.
Current views of haemolytic streptococcal pathogenesis.
Structural characterization of the virulence factor Sda1 nuclease from <i>Streptococcus pyogenes</i> .

Addiction of Hypertransformable Pneumococcal Isolates to Natural Transformation for In Vivo Fitness and Virulence.
IgG protease Mac/IdeS is not essential for phagocyte resistance or mouse virulence of M1T1 group A Streptococcus.
The Pneumococcal Serotype 15C Capsule Is Partially O-Acetylated and Allows for Limited Evasion of 23-Valent Pneumococcal Polysaccharide Vaccine-Elicited Anti-Serotype 15B Antibodies.
Development of a high-throughput screen and its use in the discovery of Streptococcus pneumoniae immunoglobulin A1 protease inhibitors.
Streptococcus suis vaccines: candidate antigens and progress.
Carriage burden, multiple colonization and antibiotic pressure promote emergence of resistant vaccine escape pneumococci.
Streptococcus pneumoniae galU gene mutation has a direct effect on biofilm growth, adherence and phagocytosis in vitro and pathogenicity in vivo.
Streptococcus suis: a re-emerging pathogen associated with occupational exposure to pigs or pork products. Part II - Pathogenesis.
Group A Streptococcal S Protein Utilizes Red Blood Cells as Immune Camouflage and Is a Critical Determinant for Immune Evasion.
Role of teichoic acid choline moieties in the virulence of Streptococcus pneumoniae.
Emergence of Amoxicillin-Resistant Variants of Spain9V-ST156 Pneumococci Expressing Serotype 11A Correlates with Their Ability to Evade the Host Immune Response.
EndoS and SpeB from Streptococcus pyogenes inhibit immunoglobulin-mediated opsonophagocytosis.
Inflammasomes in Pneumococcal Infection: Innate Immune Sensing and Bacterial Evasion Strategies.
Global emergence and population dynamics of divergent serotype 3 CC180 pneumococci.
Pathogenesis of Streptococcus pneumoniae serotype 3 during natural colonization and infections among children and its IgG correlate of protection in a mouse model.
A type IV pilus mediates DNA binding during natural transformation in Streptococcus pneumoniae.
Immunomodulatory Effects of Pneumococcal Extracellular Vesicles on Cellular and Humoral Host Defenses.
RNA thermosensors facilitate Streptococcus pneumoniae and Haemophilus influenzae immune evasion.
Vaccine escape recombinants emerge after pneumococcal vaccination in the United States.
Extracellular deoxyribonuclease made by group A Streptococcus assists pathogenesis by enhancing evasion of the innate immune response.
Position of O-Acetylation within the Capsular Repeat Unit Impacts the Biological Properties of Pneumococcal Serotypes 33A and 33F.
Serotype 1 and 8 Pneumococci Evade Sensing by Inflammasomes in Human Lung Tissue.
Deletion of the Zinc Transporter Lipoprotein AdcAll Causes Hyperencapsulation of Streptococcus pneumoniae Associated with Distinct Alleles of the Type I Restriction-Modification System.
Nonencapsulated Streptococcus pneumoniae resists extracellular human neutrophil elastase- and cathepsin G-mediated killing.
Unique genomic arrangements in an invasive serotype M23 strain of Streptococcus pyogenes identify genes that induce hypervirulence.
Pneumococcal VncR Strain-Specifically Regulates Capsule Polysaccharide Synthesis.
Immunometabolic control by Klebsiella pneumoniae.
Clinical Implications of Genomic Adaptation and Evolution of Carbapenem-Resistant Klebsiella pneumoniae.
Surface-exposed adherence molecules of Streptococcus pneumoniae.



Insertion of an immunodominant T helper cell epitope within the Group A Streptococcus M protein promotes an IFN- $\gamma$ -dependent shift from a non-protective to a protective immune response.
Hypercapsule is the cornerstone of <i>Klebsiella pneumoniae</i> in inducing pyogenic liver abscess.
Pneumococcal lipoproteins involved in bacterial fitness, virulence, and immune evasion.
Pneumolysin suppresses the initial macrophage pro-inflammatory response to <i>Streptococcus pneumoniae</i> .
The Role of Pneumococcal Virulence Factors in Ocular Infectious Diseases.
Streptococcal inhibitor of complement promotes innate immune resistance phenotypes of invasive M1T1 group A <i>Streptococcus</i> .
The group A <i>Streptococcus</i> interleukin-8 protease SpyCEP promotes bacterial intracellular survival by evasion of autophagy.
<i>Streptococcus pyogenes</i> escapes from autophagy.
Cloning, expression, purification, crystallization and preliminary X-ray diffraction analysis of SpyCEP, a candidate antigen for a vaccine against <i>Streptococcus pyogenes</i> .
Increasing incidence of <i>Streptococcus pneumoniae</i> serotype 19A and emergence of two vaccine escape recombinant ST695 strains in Liguria, Italy, 7 years after implementation of the 7-valent conjugated vaccine.
<i>Streptococcus pneumoniae</i> resists intracellular killing by olfactory ensheathing cells but not by microglia.
The Transcriptional Regulator CpsY Is Important for Innate Immune Evasion in <i>Streptococcus pyogenes</i> .
Cytocidal effect of <i>Streptococcus pyogenes</i> on mouse neutrophils in vivo and the critical role of streptolysin S.
DNase Sda1 allows invasive M1T1 Group A <i>Streptococcus</i> to prevent TLR9-dependent recognition.
Characterization of M-Type-Specific Pilus Expression in Group A <i>Streptococcus</i> .
The deficient cleavage of M protein-bound IgG by IdeS: insight into the escape of <i>Streptococcus pyogenes</i> from antibody-mediated immunity.
Nasopharyngeal colonization and invasive disease are enhanced by the cell wall hydrolases LytB and LytC of <i>Streptococcus pneumoniae</i> .
Fhb, a novel factor H-binding surface protein, contributes to the antiphagocytic ability and virulence of <i>Streptococcus suis</i> .
Capsular polysaccharide production in <i>Enterococcus faecalis</i> and contribution of CpsF to capsule serospecificity.
Cytokine mRNA expression in pneumococcal carriage, pneumonia, and sepsis in young mice.
Synergistic effects of streptolysin S and streptococcal pyrogenic exotoxin B on the mouse model of group A streptococcal infection.
Pathogenesis and pathophysiology of pneumococcal meningitis.
Human IgG Increases Virulence of <i>Streptococcus pyogenes</i> through Complement Evasion.
Role of extracellular GAPDH in <i>Streptococcus pyogenes</i> virulence.
<i>Klebsiella pneumoniae</i> Lipopolysaccharides Serotype O2afg Induce Poor Inflammatory Immune Responses Ex Vivo.
Penicillin-binding proteins in <i>Streptococcus agalactiae</i> : a novel mechanism for evasion of immune clearance.
Expanding strain coverage of a group A <i>Streptococcus</i> pilus-expressing <i>Lactococcus lactis</i> mucosal vaccine.
A novel extracellular vesicle-associated endodeoxyribonuclease helps <i>Streptococcus pneumoniae</i> evade neutrophil extracellular traps and is required for full virulence.

Group A Streptococcus cation diffusion facilitator proteins contribute to immune evasion by regulating intracellular metal concentrations.
Binding of Plasminogen to Streptococcus suis Protein Endopeptidase O Facilitates Evasion of Innate Immunity in Streptococcus suis.
Rapid pneumococcal evolution in response to clinical interventions.
Pharmacological Evaluation of Synthetic Dominant-Negative Peptides Derived from the Competence-Stimulating Peptide of Streptococcus pneumoniae.
Analysis of Global Collection of Group A Streptococcus Genomes Reveals that the Majority Encode a Trio of M and M-Like Proteins.
Bacterial genome-wide association study of hyper-virulent pneumococcal serotype 1 identifies genetic variation associated with neurotropism.
amrA encodes a putative membrane protein necessary for maximal exponential phase expression of the Mga virulence regulon in Streptococcus pyogenes.
Exposure to Acyl Homoserine Lactone Enhances Survival of Streptococcus pyogenes in Murine Macrophages.
Functional Genomic Screen Identifies Klebsiella pneumoniae Factors Implicated in Blocking Nuclear Factor $\kappa$ B (NF- $\kappa$ B) Signaling.
Characterization of protective mucosal and systemic immune responses elicited by pneumococcal surface protein PspA and PspC nasal vaccines against a respiratory pneumococcal challenge in mice.
Streptococcal 5'-Nucleotidase A (S5nA), a Novel Streptococcus pyogenes Virulence Factor That Facilitates Immune Evasion.
An hemolytic pneumolysin of Streptococcus pneumoniae manipulates human innate and CD4 <sup>+</sup> T-cell responses and reduces resistance to colonization in mice in a serotype-independent manner.
Streptolysin O promotes group A Streptococcus immune evasion by accelerated macrophage apoptosis.
The M1 protein of Streptococcus pyogenes triggers an innate uptake mechanism into polarized human endothelial cells.
The globally disseminated M1T1 clone of group A Streptococcus evades autophagy for intracellular replication.
Mpn491, a secreted nuclease of Mycoplasma pneumoniae, plays a critical role in evading killing by neutrophil extracellular traps.
Mycoplasma pneumoniae Infections: Pathogenesis and Vaccine Development.
Use of Proteins Identified through a Functional Genomic Screen To Develop a Protein Subunit Vaccine That Provides Significant Protection against Virulent Streptococcus suis in Pigs.
DNA methylation from a Type I restriction modification system influences gene expression and virulence in Streptococcus pyogenes.
Critical role for Streptococcus suis cell wall modifications and suilysin in resistance to complement-dependent killing by dendritic cells.
Clonal Changes in the Pneumococcal Population Carried by Portuguese Children during Six Years of Private Use of the 13-Valent Pneumococcal Conjugate Vaccine: the Relative Contribution of Clonal Expansion, Clonal Emergence, and Capsular Switch Events.
Evasion of human innate and acquired immunity by a bacterial homolog of CD11b that inhibits opsonophagocytosis.
Penicillin-binding protein 1a promotes resistance of group B streptococcus to antimicrobial peptides.
Streptococcus suis 2 Transcriptional Regulator TstS Stimulates Cytokine Production and Bacteremia to Promote Streptococcal Toxic Shock-Like Syndrome.
Streptococcus pyogenes Hijacks Host Glutathione for Growth and Innate Immune Evasion.

Klebsiella pneumoniae targets an EGF receptor-dependent pathway to subvert inflammation.
The streptococcal protease IdeS modulates bacterial IgG Fc binding and generates 1/2Fc fragments with the ability to prime polymorphonuclear leucocytes.
Pneumococcal immune evasion: ZmpC inhibits neutrophil influx.
Cationic antimicrobial peptide resistance mechanisms of streptococcal pathogens.
Specific C-terminal cleavage and inactivation of interleukin-8 by invasive disease isolates of Streptococcus pyogenes.
Characterization of an Immunoglobulin Binding Protein (IbpM) From Mycoplasma pneumoniae.
Nuclease A (Gbs0661), an extracellular nuclease of Streptococcus agalactiae, attacks the neutrophil extracellular traps and is needed for full virulence.
Antigenic Variation in Streptococcus pneumoniae PspC Promotes Immune Escape in the Presence of Variant-Specific Immunity.
Pneumococcal genome sequencing tracks a vaccine escape variant formed through a multi-fragment recombination event.
Structural basis of host cell recognition by the pilus adhesin from Streptococcus pneumoniae.
Streptococcus iniae cpsG alters capsular carbohydrate composition and is a cause of serotype switching in vaccinated fish.
Multicomponent Vaccines against Group A Streptococcus Can Effectively Target Broad Disease Presentations.
Novel pneumococcal capsule type 33E results from the inactivation of glycosyltransferase WciE in vaccine type 33F.
A streptococcal penicillin-binding protein is critical for resisting innate airway defenses in the neonatal lung.
Group A Streptococcal M1 Protein Provides Resistance against the Antimicrobial Activity of Histones.
Analysis of group A Streptococcus gene expression in humans with pharyngitis using a microarray.
Within-host microevolution of Streptococcus pneumoniae is rapid and adaptive during natural colonisation.
Structural characterization of the virulence factor nuclease A from Streptococcus agalactiae.
The cysteine protease ApdS from Streptococcus suis promotes evasion of innate immune defenses by cleaving the antimicrobial peptide cathelicidin LL-37.
How Streptococcus suis serotype 2 attempts to avoid attack by host immune defenses.
Streptococcus pyogenes recruits collagen via surface-bound fibronectin: a novel colonization and immune evasion mechanism.
Group A Streptococcal M1 Protein Sequesters Cathelicidin to Evade Innate Immune Killing.
Group B Streptococcus suppression of phagocyte functions by protein-mediated engagement of human Siglec-5.
Crystal structures of Wzb of Escherichia coli and CpsB of Streptococcus pneumoniae, representatives of two families of tyrosine phosphatases that regulate capsule assembly.
Identification of conditionally essential genes for Streptococcus suis infection in pigs.
The VraSR regulatory system contributes to virulence in Streptococcus suis via resistance to innate immune defenses.
Unusual Hypermucoviscous Clinical Isolate of Klebsiella pneumoniae with No Known Determinants of Hypermucoviscosity.
Streptococcus suis adenosine synthase functions as an effector in evasion of PMN-mediated innate immunity.
Klebsiella pneumoniae outer membrane protein A is required to prevent the activation of airway epithelial cells.

Inclusion bodies and pH lowering: as an effect of gold nanoparticles in <i>Streptococcus pneumoniae</i> .
<i>Kingella kingae</i> Surface Polysaccharides Promote Resistance to Neutrophil Phagocytosis and Killing.
Recent advances in understanding the molecular basis of group B <i>Streptococcus</i> virulence.
Short communication: comparison of virulence factors in <i>Klebsiella pneumoniae</i> strains associated with multiple or single cases of mastitis.
Risk of red queen dynamics in pneumococcal vaccine strategy.
Neutrophil extracellular traps in the central nervous system hinder bacterial clearance during pneumococcal meningitis.
Autophagy, cell death, and cytokines in <i>K. pneumoniae</i> infection: therapeutic perspectives.
Bacterial exploitation of phosphorylcholine mimicry suppresses inflammation to promote airway infection.
The evolutionary trade-offs in phage-resistant <i>Klebsiella pneumoniae</i> entail cross-phage sensitization and loss of multidrug resistance.
Diminished Capsule Exacerbates Virulence, Blood-Brain Barrier Penetration, Intracellular Persistence, and Antibiotic Evasion of Hyperhemolytic Group B <i>Streptococci</i> .
Neutrophils select hypervirulent CovRS mutants of M1T1 group A <i>Streptococcus</i> during subcutaneous infection of mice.
Deciphering Key Residues Involved in the Virulence-promoting Interactions between <i>Streptococcus pneumoniae</i> and Human Plasminogen.
The two-component response regulator LiaR regulates cell wall stress responses, pili expression and virulence in group B <i>Streptococcus</i> .
PspA facilitates evasion of pneumococci from bactericidal activity of neutrophil extracellular traps (NETs).
Modular Architecture and Unique Teichoic Acid Recognition Features of Choline-Binding Protein L (CbpL) Contributing to Pneumococcal Pathogenesis.
A high-resolution view of genome-wide pneumococcal transformation.
Comparative genome analysis of two <i>Streptococcus phocae</i> subspecies provides novel insights into pathogenicity.
A novel streptococcal leucine zipper protein (Lzp) binds to human immunoglobulins.
Conformation and Cross-Protection in Group B <i>Streptococcus</i> Serotype III and <i>Streptococcus pneumoniae</i> Serotype 14: A Molecular Modeling Study.
One More Disguise in the Stealth Behavior of <i>Streptococcus pyogenes</i> .
Parallel Evolution of Group B <i>Streptococcus</i> Hypervirulent Clonal Complex 17 Unveils New Pathoadaptive Mutations.
<i>Klebsiella pneumoniae</i> Reduces SUMOylation To Limit Host Defense Responses.
Immune-mediated phagocytosis and killing of <i>Streptococcus pneumoniae</i> are associated with direct and bystander macrophage apoptosis.
The Mga Regulon but Not Deoxyribonuclease Sda1 of Invasive M1T1 Group A <i>Streptococcus</i> Contributes to In Vivo Selection of CovRS Mutations and Resistance to Innate Immune Killing Mechanisms.
A bacterial pathogen co-opts host plasmin to resist killing by cathelicidin antimicrobial peptides.
Plasmin(ogen) acquisition by group A <i>Streptococcus</i> protects against C3b-mediated neutrophil killing.
The hypermucoviscosity of hypervirulent <i>K. pneumoniae</i> confers the ability to evade neutrophil-mediated phagocytosis.
Secretion of a pneumococcal type II secretion system pilus correlates with DNA uptake during transformation.

Emerging Fatal Ib/CC12 Hypervirulent Multiresistant <i>Streptococcus agalactiae</i> in Young Infants With Bloodstream Infection in China.
Group A <i>Streptococcus</i> Induces LAPosomes via SLO/ $\beta$ 1 Integrin/NOX2/ROS Pathway in Endothelial Cells That Are Ineffective in Bacterial Killing and Suppress Xenophagy.
A Neutralizing Monoclonal IgG1 Antibody of Platelet-Activating Factor Acetylhydrolase SsE Protects Mice against Lethal Subcutaneous Group A <i>Streptococcus</i> Infection.
Variation of pneumococcal Pilus-1 expression results in vaccine escape during Experimental Otitis Media [EOM].
Pleiotropic virulence factor - <i>Streptococcus pyogenes</i> fibronectin-binding proteins.
Insight into the Pathogenic Mechanism of <i>Mycoplasma pneumoniae</i> .
The Mga virulence regulon: infection where the grass is greener.
Impact of pneumococcal microbial surface components recognizing adhesive matrix molecules on colonization.
Directed vaccination against pneumococcal disease.
Group A streptococci induce stronger M protein-fibronectin interaction when specific human antibodies are bound.
Two multi-fragment recombination events resulted in the $\beta$ -lactam-resistant serotype 11A-ST6521 related to Spain9V-ST156 pneumococcal clone spreading in south-western Europe, 2008 to 2016.
Mga is sufficient to activate transcription in vitro of sof-sfbX and other Mga-regulated virulence genes in the group A <i>Streptococcus</i> .
NAD-Glycohydrolase Depletes Intracellular NAD(+) and Inhibits Acidification of Autophagosomes to Enhance Multiplication of Group A <i>Streptococcus</i> in Endothelial Cells.
<i>Streptococcus pyogenes</i> Escapes Killing from Extracellular Histones through Plasminogen Binding and Activation by Streptokinase.
Intrinsic Maturation Neonatal Immune Deficiencies and Susceptibility to Group B <i>Streptococcus</i> Infection.
Molecular analyses identifies new domains and structural differences among <i>Streptococcus pneumoniae</i> immune evasion proteins PspC and Hic.
Screening for phagocytosis resistance-related genes via a transposon mutant library of <i>Streptococcus suis</i> serotype 2.
Structure of ScpC, a virulence protease from <i>Streptococcus pyogenes</i> , reveals the functional domains and maturation mechanism.
Antibiotic Resistance and Virulence Profiles of <i>Klebsiella pneumoniae</i> Strains Isolated From Different Clinical Sources.
Antibody-Mediated Killing of Carbapenem-Resistant ST258 <i>Klebsiella pneumoniae</i> by Human Neutrophils.
Monitoring the long-term molecular epidemiology of the pneumococcus and detection of potential 'vaccine escape' strains.
<i>Klebsiella pneumoniae</i> subverts the activation of inflammatory responses in a NOD1-dependent manner.
Contribution of Secreted NADase and Streptolysin O to the Pathogenesis of Epidemic Serotype M1 <i>Streptococcus pyogenes</i> Infections.
<i>Streptococcus suis</i> Serotype 2 Infection Causes Host Immunomodulation through Induction of Thymic Atrophy.
Proteomics identification of novel fibrinogen-binding proteins of <i>Streptococcus suis</i> contributing to antiphagocytosis.
Antibody blocks acquisition of bacterial colonization through agglutination.

Contribution of fucose-containing capsules in <i>Klebsiella pneumoniae</i> to bacterial virulence in mice.
Domains required for transcriptional activation show conservation in the <i>mga</i> family of virulence gene regulators.
Group A streptococcal cysteine protease degrades C3 (C3b) and contributes to evasion of innate immunity.
<i>Streptococcus suis</i> Serotype 2 Infection Impairs Interleukin-12 Production and the MHC-II-Restricted Antigen Presentation Capacity of Dendritic Cells.
<i>Streptococcus pneumoniae</i> endopeptidase O (PepO) is a multifunctional plasminogen- and fibronectin-binding protein, facilitating evasion of innate immunity and invasion of host cells.
The Road to Infection: Host-Microbe Interactions Defining the Pathogenicity of <i>Streptococcus bovis</i> / <i>Streptococcus equinus</i> Complex Members.
Bacterial transformation: ComFA is a DNA-dependent ATPase that forms complexes with ComFC and DprA.
<i>Streptococcus agalactiae</i> <i>cadD</i> alleviates metal stress and promotes intracellular survival in macrophages and ascending infection during pregnancy.
Requirement and Synergistic Contribution of Platelet-Activating Factor Acetylhydrolase Sse and Streptolysin S to Inhibition of Neutrophil Recruitment and Systemic Infection by Hypervirulent <i>emm3</i> Group A <i>Streptococcus</i> in Subcutaneous Infection of Mice.
Identification of novel pig and human immunoglobulin G-binding proteins and characterization of the binding regions of enolase from <i>Streptococcus suis</i> serotype 2.
Immunization With a Secreted Esterase Protects Mice Against Multiple Serotypes (M1, M3, and M28) of Group A <i>Streptococcus</i> .
Unbiased homeologous recombination during pneumococcal transformation allows for multiple chromosomal integration events.
The serotype of type Ia and III group B streptococci is determined by the polymerase gene within the polycistronic capsule operon.
Capsular Polysaccharide Is Essential for the Virulence of the Antimicrobial-Resistant Pathogen <i>Enterobacter hormaechei</i> .
Generic determinants of <i>Streptococcus</i> colonization and infection.
Vaccine escape of pilated <i>Streptococcus pneumoniae</i> strains.
Streptolysin O Deficiency in <i>Streptococcus pyogenes</i> M1T1 <i>covR/S</i> Mutant Strain Attenuates Virulence in In Vitro and In Vivo Infection Models.
Greedy de novo motif discovery to construct motif repositories for bacterial proteomes.
Variation in a surface-exposed region of the <i>Mycoplasma pneumoniae</i> P40 protein as a consequence of homologous DNA recombination between RepMP5 elements.
Immunobiology of the Classical Lancefield Group A Streptococcal Carbohydrate Antigen.
A role for LHC1 in higher order structure and complement binding of the <i>Cryptococcus neoformans</i> capsule.
Neutrophil membrane-coated nanoparticles exhibit increased antimicrobial activities in an anti-microbial resistant <i>K. pneumoniae</i> infection model.
<i>Enterococcus faecalis</i> capsular polysaccharide serotypes C and D and their contributions to host innate immune evasion.
Group A <i>Streptococcus</i> secreted esterase hydrolyzes platelet-activating factor to impede neutrophil recruitment and facilitate innate immune evasion.
Differential bacterial gene expression during experimental pneumococcal endophthalmitis.
Separating Bacteria by Capsule Amount Using a Discontinuous Density Gradient.
Envelope Structures of Gram-Positive Bacteria.

Regulation of inhibition of neutrophil infiltration by the two-component regulatory system CovRS in subcutaneous murine infection with group A streptococcus.
Role of colanic acid polysaccharide in serum resistance in vivo and in adherence.
Streptococcus pyogenes CAMP factor attenuates phagocytic activity of RAW 264.7 cells.
[Correlation between Type IV secretion system component VirD4 and virulence for Streptococcus suis 2].
The streptococcal cysteine protease SpeB is not a natural immunoglobulin-cleaving enzyme.
Refining the Pneumococcal Competence Regulon by RNA Sequencing.
Structure, Function, and Regulation of the Essential Virulence Factor Capsular Polysaccharide of Vibrio vulnificus.
Hypervirulent Group A Streptococcus of Genotype emm3 Invades the Vascular System in Pulmonary Infection of Mice.
Nasopharyngeal microbiome composition associated with Streptococcus pneumoniae colonization suggests a protective role of Corynebacterium in young children.
Identification of residues responsible for the defective virulence gene regulator Mga produced by a natural mutant of Streptococcus pyogenes.
A multiomics analysis of direct interkingdom dynamics between influenza A virus and Streptococcus pneumoniae uncovers host-independent changes to bacterial virulence fitness.
Complement C3aR/C5aR-binding protein Suilysin of Streptococcus suis contributes to monocyte chemotaxis.
Defining the Mga regulon: Comparative transcriptome analysis reveals both direct and indirect regulation by Mga in the group A streptococcus.
[Mechanism behind streptococcus toxic shock-like syndrome onset--immune evasion and bacterial properties].
Molecular mimicry of host sialylated glycans allows a bacterial pathogen to engage neutrophil Siglec-9 and dampen the innate immune response.
The crystal structure of the streptococcal collagen-like protein 2 globular domain from invasive M3-type group A Streptococcus shows significant similarity to immunomodulatory HIV protein gp41.
Antiphagocytic function of an IgG glycosyl hydrolase from Streptococcus equi subsp. equi and its use as a vaccine component.
A MyD88-JAK1-STAT1 complex directly induces SOCS-1 expression in macrophages infected with Group A Streptococcus.
The DNases of pathogenic Lancefield streptococci.
Cell wall-anchored nuclease of Streptococcus sanguinis contributes to escape from neutrophil extracellular trap-mediated bacteriocidal activity.
Fragments of bacterial endoglycosidase s and immunoglobulin g reveal subdomains of each that contribute to deglycosylation.
Crystal structure of Streptococcus pyogenes EndoS, an immunomodulatory endoglycosidase specific for human IgG antibodies.
Molecular pathogenesis of the hyaluronic acid capsule of Pasteurella multocida.
Distinct effects on diversifying selection by two mechanisms of immunity against Streptococcus pneumoniae.
A pivotal role for the Streptococcus iniae extracellular polysaccharide in triggering proinflammatory cytokines transcription and inducing death in rainbow trout.
Flying Under the Radar: Immune Evasion by Group B Streptococcus.
Similar genomic patterns of clinical infective endocarditis and oral isolates of Streptococcus sanguinis and Streptococcus gordonii.

A common theme in interaction of bacterial immunoglobulin-binding proteins with immunoglobulins illustrated in the equine system.
Chemokine degradation by the Group A streptococcal serine proteinase ScpC can be reconstituted in vitro and requires two separate domains.
Discovery and expression of 3 siglecs-like in <i>Oreochromis niloticus</i> neutrophil, and their interaction with group B streptococcal sialylated capsular polysaccharides.
Functional Analysis of Two Novel <i>Streptococcus iniae</i> Virulence Factors Using a Zebrafish Infection Model.
A case of neck abscess caused by rare hypervirulent <i>Klebsiella pneumoniae</i> , capsular type K20 and sequence type 420.
The key surface components of <i>Pasteurella multocida</i> : capsule and lipopolysaccharide.
Hemolytic Membrane Vesicles of Group B <i>Streptococcus</i> Promote Infection.
Adherence molecules of pathogenic pneumococci.
Serotype 3 pneumococci sequester platelet-derived human thrombospondin-1 via the adhesin and immune evasion protein Hic.
Toward Clinical use of the IgG Specific Enzymes IdeS and EndoS against Antibody-Mediated Diseases.
Structural and functional insights into peptidoglycan access for the lytic amidase LytA of <i>Streptococcus pneumoniae</i> .
Biosynthesis and regulation mechanisms of the <i>Pasteurella multocida</i> capsule.
Minimal Peptidoglycan (PG) Turnover in Wild-Type and PG Hydrolase and Cell Division Mutants of <i>Streptococcus pneumoniae</i> D39 Growing Planktonically and in Host-Relevant Biofilms.
A novel bacterial resistance mechanism against human group IIA-secreted phospholipase A2: role of <i>Streptococcus pyogenes</i> sortase A.
Comparative transcriptomic analysis reveal genes involved in the pathogenicity increase of <i>Streptococcus suis</i> epidemic strains.
The Hypervariable region of <i>Streptococcus pyogenes</i> M protein escapes antibody attack by antigenic variation and weak immunogenicity.
Mechanism of antibody-specific deglycosylation and immune evasion by Streptococcal IgG-specific endoglycosidases.
Resistance of <i>Klebsiella pneumoniae</i> to the innate immune system of African green monkeys.
Pneumolysin, PspA, and PspC contribute to pneumococcal evasion of early innate immune responses during bacteremia in mice.
The Retropepsin-Type Protease APRc as a Novel Ig-Binding Protein and Moonlighting Immune Evasion Factor of <i>Rickettsia</i> .
Sequence variability is correlated with weak immunogenicity in <i>Streptococcus pyogenes</i> M protein.
Enzyme Characterization of Pro-virulent SntA, a Cell Wall-Anchored Protein of <i>Streptococcus suis</i> , With Phosphodiesterase Activity on cyclic-di-AMP at a Level Suited to Limit the Innate Immune System.
Liquid-liquid diffusion crystallization improves the X-ray diffraction of EndoS, an endo- $\beta$ -N-acetylglucosaminidase from <i>Streptococcus pyogenes</i> with activity on human IgG.
Mechanistic basis of choline import involved in teichoic acids and lipopolysaccharide modification.
Identification and functional characterization of the putative polysaccharide biosynthesis protein (CapD) of <i>Enterococcus faecium</i> U0317.
The pneumococcal $\sigma(X)$ activator, ComW, is a DNA-binding protein critical for natural transformation.
Immunoglobulins and their receptors, and subversion of their protective roles by bacterial pathogens.
Role of glycan synthesis in colonization of the mammalian gut by the bacterial symbiont <i>Bacteroides fragilis</i> .



Genetic Structure, Function, and Evolution of Capsule Biosynthesis Loci in <i>Vibrio parahaemolyticus</i> .
Mannheimia haemolytica: bacterial-host interactions in bovine pneumonia.
Genomic Stability of Aggregatibacter actinomycetemcomitans during Persistent Oral Infection in Human.
Fc-mediated nonspecific binding between fibronectin-binding protein I of Streptococcus pyogenes and human immunoglobulins.
Genetic diversity of the Pneumococcal CbpA: Implications for next-generation vaccine development.
Near-Infrared Light-Sensitive Nano Neuro-Immune Blocker Capsule Relieves Pain and Enhances the Innate Immune Response for Necrotizing Infection.
The Role of Streptococcal Cell-Envelope Proteases in Bacterial Evasion of the Innate Immune System.
PepO is a target of the two-component systems VicRK and CovR required for systemic virulence of Streptococcus mutans.
Preterm infants harbour diverse Klebsiella populations, including atypical species that encode and produce an array of antimicrobial resistance- and virulence-associated factors.
Two capsular polysaccharides enable Bacillus cereus G9241 to cause anthrax-like disease.
The capsule of Bacillus anthracis protects it from the bactericidal activity of human defensins and other cationic antimicrobial peptides.
Wza the translocon for E. coli capsular polysaccharides defines a new class of membrane protein.
Comprehensive genomic analysis and characterization of a new ST 174 type Klebsiella variicola strain isolated from chicken embryos.
Shifting genetic structure of invasive serotype 19A pneumococci in the United States.
Characterization of streptococcal platelet-activating factor acetylhydrolase variants that are involved in innate immune evasion.
Insight of host immune evasion mediated by two variants of group A Streptococcus Mac protein.
The Wzi outer membrane protein mediates assembly of a tight capsular polysaccharide layer on the Acinetobacter baumannii cell surface.
Product Specificity of C4-Reductases in the Biosynthesis of GDP-6-Deoxy-Heptoses during Capsular Polysaccharide Formation in Campylobacter jejuni.
Cleavage of IgGs by proteases associated with invasive diseases: an evasion tactic against host immunity?
Peroxide responsive regulator PerR of group A Streptococcus is required for the expression of phage-associated DNase Sda1 under oxidative stress.
Functional characterization of the gonococcal polyphosphate pseudo-capsule.
The crystal structure of the major pneumococcal autolysin LytA in complex with a large peptidoglycan fragment reveals the pivotal role of glycans for lytic activity.
Hyaluronidase Impairs Neutrophil Function and Promotes Group B Streptococcus Invasion and Preterm Labor in Nonhuman Primates.
Functional improvement of porcine neonatal pancreatic cell clusters via conformational encapsulation using an air-driven encapsulator.
IgG-binding proteins of bacteria.
Homophilic protein interactions facilitate bacterial aggregation and IgG-dependent complex formation by the Streptococcus canis M protein SCM.
Comparative Genomic Analysis of Virulent Vibrio (Listonella) anguillarum Serotypes Revealed Genetic Diversity and Genomic Signatures in the O-Antigen Biosynthesis Gene Cluster.
[Neonatal group B streptococcus infection in the Children's Hospital of Gansu Province through PCR array].

Involvement of lipoprotein PpiA of <i>Streptococcus gordonii</i> in evasion of phagocytosis by macrophages.
Identification and Co-complex Structure of a New <i>S. pyogenes</i> SpeB Small Molecule Inhibitor.
Group B <i>Streptococcus</i> (GBS) disrupts by calpain activation the actin and microtubule cytoskeleton of macrophages.
Extracellular Nucleases of <i>Streptococcus equi</i> subsp. <i>zooepidemicus</i> Degrade Neutrophil Extracellular Traps and Impair Macrophage Activity of the Host.
Characterization of novel nuclease and protease activities among <i>Leptospiral</i> immunoglobulin-like proteins.
Molecular Basis of Broad Spectrum N-Glycan Specificity and Processing of Therapeutic IgG Monoclonal Antibodies by Endoglycosidase S2.
SIGN-R1, a C-type lectin, binds to Bip/GRP78 and this interaction mediates the regurgitation of T-cell-independent type 2 antigen dextran through the endoplasmic reticulum.
<i>Streptococcal</i> protein FOG, a novel matrix adhesin interacting with collagen I in vivo.
Crystal structure of group A <i>streptococcus</i> Mac-1: insight into dimer-mediated specificity for recognition of human IgG.
Expression of the type 1 pneumococcal pilus is bistable and negatively regulated by the structural component RrgA.
Insights into substrate recognition and specificity for IgG by Endoglycosidase S2.
<i>Streptococcus equi</i> : a pathogen restricted to one host.
Group B <i>Streptococcus</i> induces macrophage apoptosis by calpain activation.
C3- and C3/C5-Epimerases Required for the Biosynthesis of the Capsular Polysaccharides from <i>Campylobacter jejuni</i> .
Immunoglobulin-binding domains: Protein L from <i>Peptostreptococcus magnus</i> .
Heterogeneous expression of Pil3 pilus is critical for <i>Streptococcus gallolyticus</i> translocation across polarized colonic epithelial monolayers.
A Requirement for Neutrophil Glycosaminoglycans in Chemokine:Receptor Interactions Is Revealed by the <i>Streptococcal</i> Protease SpyCEP.
Lipoteichoic acid of <i>Streptococcus gordonii</i> as a negative regulator of human dendritic cell activation.
<i>Streptococcal</i> M1 Strikes by Neutralizing Cathelicidins.
CP40 from <i>Corynebacterium pseudotuberculosis</i> is an endo- $\beta$ -N-acetylglucosaminidase.
Identification of NAD <sup>+</sup> synthetase from <i>Streptococcus sobrinus</i> as a B-cell-stimulatory protein.
Investigating the molecular basis for the virulence of <i>Escherichia coli</i> K5 by nuclear magnetic resonance analysis of the capsule polysaccharide.
Fab fragment glycosylated IgG may play a central role in placental immune evasion.
Mechanism of cooperative N-glycan processing by the multi-modular endoglycosidase EndoE.
Direct Staudinger-Phosphonite Reaction Provides Methylphosphonamidates as Inhibitors of CE4 De-N-acetylases.
A case of structure determination using pseudosymmetry.
Biosynthesis of d-glycero-l-gluco-Heptose in the Capsular Polysaccharides of <i>Campylobacter jejuni</i> .
Engineering an Fc-Fusion of a Capsule Degrading Enzyme for the Treatment of Anthrax.
A Broad-Spectrum Chemokine-Binding Protein of Bovine Papular Stomatitis Virus Inhibits Neutrophil and Monocyte Infiltration in Inflammatory and Wound Models of Mouse Skin.
Ivermectin-dependent attachment of neutrophils and peripheral blood mononuclear cells to <i>Dirofilaria immitis</i> microfilariae in vitro.

Table A2-54, Cluster 53

Cluster 53 focuses on <i>Salmonella typhimurium</i> and other bacteria, emphasizing their pathogenicity, virulence, and the molecular mechanisms they use to evade the immune system (375)
<i>Salmonella enterica</i> serovar typhimurium exploits Toll-like receptor signaling during the host-pathogen interaction.
The SseL protein inhibits the intracellular NF- $\kappa$ B pathway to enhance the virulence of <i>Salmonella Pullorum</i> in a chicken model.
Innate immune response to <i>Salmonella typhimurium</i> , a model enteric pathogen.
Molecular Mechanisms Used by <i>Salmonella</i> to Evade the Immune System.
<i>Salmonella</i> rapidly kill dendritic cells via a caspase-1-dependent mechanism.
<i>Salmonella</i> : a model for bacterial pathogenesis.
Host Cell Death Responses to Non-typhoidal <i>Salmonella</i> Infection.
<i>Salmonella</i> Virulence and Immune Escape.
Methods to Illuminate the Role of <i>Salmonella</i> Effector Proteins during Infection: A Review.
Cellular aspects of immunity to intracellular <i>Salmonella enterica</i> .
Bacterial adhesion and host cell factors leading to effector protein injection by type III secretion system.
Evidence and speculation: the response of <i>Salmonella</i> confronted by autophagy in macrophages.
Cooperative Immune Suppression by <i>Escherichia coli</i> and <i>Shigella</i> Effector Proteins.
<i>Salmonella enterica</i> serovars Typhimurium and Typhi as model organisms: revealing paradigm of host-pathogen interactions.
GH18 family glycoside hydrolase Chitinase A of <i>Salmonella</i> enhances virulence by facilitating invasion and modulating host immune responses.
<i>Salmonella</i> 's long-term relationship with its host.
Identification of Novel Host Interactors of Effectors Secreted by <i>Salmonella</i> and <i>Citrobacter</i> .
<i>Salmonella</i> persistence and transmission strategies.
Typhoidal <i>Salmonella</i> : Distinctive virulence factors and pathogenesis.
<i>Salmonella</i> AvrA Coordinates Suppression of Host Immune and Apoptotic Defenses via JNK Pathway Blockade.
<i>Salmonella</i> Suppresses the TRIF-Dependent Type I Interferon Response in Macrophages.
Visiting the cell biology of <i>Salmonella</i> infection.
The N terminus of type III secretion needle protein YscF from <i>Yersinia pestis</i> functions to modulate innate immune responses.
The <i>Salmonella</i> pathogenicity island-2 subverts human NLRP3 and NLRC4 inflammasome responses.
The flagellar regulator TviA reduces pyroptosis by <i>Salmonella enterica</i> serovar Typhi.
The impact of lipid A modification on biofilm and related pathophysiological phenotypes, endotoxicity, immunogenicity, and protection of <i>Salmonella Typhimurium</i> .
Horizontally acquired glycosyltransferase operons drive salmonellae lipopolysaccharide diversity.
Interactions of <i>Salmonella enterica</i> with dendritic cells.
Innate immune detection of flagellin positively and negatively regulates salmonella infection.
<i>Salmonella</i> produce microRNA-like RNA fragment Sal-1 in the infected cells to facilitate intracellular survival.
A type 6 secretion system (T6SS) encoded gene within <i>Salmonella enterica</i> serovar Enteritidis contributes to virulence.
<i>Salmonella</i> Persistence and Host Immunity Are Dictated by the Anatomical Microenvironment.
<i>Shigella</i> IpaH Family Effectors as a Versatile Model for Studying Pathogenic Bacteria.

Molecular Mechanisms of Salmonella Effector Proteins: A Comprehensive Review.
Salmonella downregulates Nod-like receptor family CARD domain containing protein 4 expression to promote its survival in B cells by preventing inflammasome activation and cell death.
Attaching and effacing pathogens: the effector ABC of immune subversion.
A Yersinia effector protein promotes virulence by preventing inflammasome recognition of the type III secretion system.
Bringing down the host: enteropathogenic and enterohaemorrhagic Escherichia coli effector-mediated subversion of host innate immune pathways.
Mechanisms used by virulent Salmonella to impair dendritic cell function and evade adaptive immunity.
Modulation of innate immune responses by Yersinia type III secretion system translocators and effectors.
[Outer membrane remodeling of Salmonella typhimurium and host innate immunity].
Salmonella-regulated lipopolysaccharide modifications.
Exploiting host immunity: the Salmonella paradigm.
Formate oxidation in the intestinal mucus layer enhances fitness of Salmonella enterica serovar Typhimurium.
Modulation of host innate immune response in the bladder by uropathogenic Escherichia coli.
Single-cell RNA-seq ties macrophage polarization to growth rate of intracellular Salmonella.
Bacterial growth rate and host factors as determinants of intracellular bacterial distributions in systemic Salmonella enterica infections.
Virulence factors of Yersinia pestis are overcome by a strong lipopolysaccharide response.
BopC is a type III secreted effector protein of Burkholderia pseudomallei.
Yersinia pestis Ail: multiple roles of a single protein.
Suppression of dendritic cell and T-cell activation by the pRST <sup>98</sup> Salmonella plasmid.
Virulence factors of uropathogenic E. coli and their interaction with the host.
Differential timing of antibody-mediated phagocytosis and cell-free killing of invasive African Salmonella allows immune evasion.
Identification of HilD-regulated genes in Salmonella enterica serovar Typhimurium.
Neutrophils are resistant to Yersinia YopJ/P-induced apoptosis and are protected from ROS-mediated cell death by the type III secretion system.
Yersinia V-antigen exploits toll-like receptor 2 and CD14 for interleukin 10-mediated immunosuppression.
Switching off Bacterial Flagellar Biogenesis by YdiU-Mediated UMPylation of FlhDC.
Hijacking of the pleiotropic cytokine interferon- $\gamma$ by the type III secretion system of Yersinia pestis.
Yersinia adhesins: An arsenal for infection.
Structural Modifications of Bacterial Lipopolysaccharide that Facilitate Gram-Negative Bacteria Evasion of Host Innate Immunity.
Immunity to intestinal pathogens: lessons learned from Salmonella.
Quinolone/fluoroquinolone susceptibility in Escherichia coli correlates with human polymicrobial bacteriuria and with in vitro interleukine-8 suppression.
Loss of very-long O-antigen chains optimizes capsule-mediated immune evasion by Salmonella enterica serovar Typhi.
Immunology of Yersinia pestis Infection.
Interaction of Yersinia with the gut: mechanisms of pathogenesis and immune evasion.
The Biology of the Escherichia coli Extracellular Matrix.

Lipopolysaccharides belonging to different Salmonella serovars are differentially capable of activating Toll-like receptor 4.
Modulation of host signaling in the inflammatory response by enteropathogenic Escherichia coli virulence proteins.
Acquisition of ompT reveals cryptic virulence function of autotransporter YapE in Yersinia pestis.
Phase variation of the lpf operon is a mechanism to evade cross-immunity between Salmonella serotypes.
Evasion of human innate immunity without antagonizing TLR4 by mutant Salmonella enterica serovar Typhimurium having penta-acylated lipid A.
Yersinia interactions with regulated cell death pathways.
Impact of plasmids, including those encoding VirB4/D4 type IV secretion systems, on Salmonella enterica serovar Heidelberg virulence in macrophages and epithelial cells.
Activation and Evasion of Inflammasomes by Yersinia.
Typhoid fever: "you can't hit what you can't see".
The role of the galU gene of uropathogenic Escherichia coli in modulating macrophage TNF- $\alpha$ response.
Oxidative metabolism enables Salmonella evasion of the NLRP3 inflammasome.
The Yersinia Virulence Factor YopM Hijacks Host Kinases to Inhibit Type III Effector-Triggered Activation of the Pyrin Inflammasome.
Type I interferon induces necroptosis in macrophages during infection with Salmonella enterica serovar Typhimurium.
Subversion of Host Innate Immunity by Uropathogenic Escherichia coli.
The role of the QseC quorum-sensing sensor kinase in colonization and norepinephrine-enhanced motility of Salmonella enterica serovar Typhimurium.
Host acid signal controls Salmonella flagella biogenesis through CadC-YdiV axis.
Lipopolysaccharide modification in Gram-negative bacteria during chronic infection.
O-antigen delays lipopolysaccharide recognition and impairs antibacterial host defense in murine intestinal epithelial cells.
RIPK3-Dependent Recruitment of Low-Inflammatory Myeloid Cells Does Not Protect from Systemic Salmonella Infection.
Suppression of T and B lymphocyte activation by a Yersinia pseudotuberculosis virulence factor, yopH.
Thrombin-activatable fibrinolysis inhibitor is degraded by Salmonella enterica and Yersinia pestis.
Deceiving the big eaters: Salmonella Typhimurium SopB subverts host cell xenophagy in macrophages via dual mechanisms.
(p)ppGpp-Dependent Regulation of the Nucleotide Hydrolase PpnN Confers Complement Resistance in Salmonella enterica Serovar Typhimurium.
Modulation of the Inflammasome Signaling Pathway by Enteropathogenic and Enterohemorrhagic Escherichia coli.
Yersinia pestis subverts the dermal neutrophil response in a mouse model of bubonic plague.
Yersinia outer protein E, YopE. A versatile type III effector molecule for cytosolic targeting of heterologous antigens by attenuated Salmonella.
Arg-GlcNAcylation on TRADD by NleB and SseK1 Is Crucial for Bacterial Pathogenesis.
Structural and Functional Variation in Outer Membrane Polysaccharide Export (OPX) Proteins from the Two Major Capsule Assembly Pathways Present in Escherichia coli.
Salmonella escapes antigen presentation through K63 ubiquitination mediated endosomal proteolysis of MHC II via modulation of endosomal acidification in dendritic cells.
Recognition of lipid A variants by the TLR4-MD-2 receptor complex.

Salmonella enterica Serovar Typhi Lipopolysaccharide O-Antigen Modification Impact on Serum Resistance and Antibody Recognition.
Functional characterizations of effector protein BipC, a type III secretion system protein, in Burkholderia pseudomallei pathogenesis.
The Multivalent Adhesion Molecule SSO1327 plays a key role in Shigella sonnei pathogenesis.
Attenuation of human neutrophil migration and function by uropathogenic bacteria.
Escherichia coli heme oxygenase modulates host innate immune responses.
Salmonella enterica infection stimulates macrophages to hemophagocytose.
The roles of the virulence factor IpaB in Shigella spp. in the escape from immune cells and invasion of epithelial cells.
New Insights on the Early Interaction Between Typhoid and Non-typhoid Salmonella Serovars and the Host Cells.
Regulation of ydiV-induced biological characteristics permits Escherichia coli evasion of the host STING inflammatory response.
Host-pathogen interactions in urinary tract infection.
Targeting bacterial virulence: inhibitors of type III secretion in Yersinia.
EK3D: an E. coli K antigen 3-dimensional structure database.
Bacterial type III translocation: a unique mechanism for cytosolic display of heterologous antigens by attenuated Salmonella.
Active suppression of early immune response in tobacco by the human pathogen Salmonella Typhimurium.
The role of lipopolysaccharide moieties in macrophage response to Escherichia coli.
EsiB, a novel pathogenic Escherichia coli secretory immunoglobulin A-binding protein impairing neutrophil activation.
New insights into the crosstalk between Shigella and T lymphocytes.
Salmonella evades D-amino acid oxidase to promote infection in neutrophils.
The genetic basis of Escherichia coli pathoadaptation to macrophages.
Role of immune response in Yersinia pestis infection.
Early evolutionary loss of the lipid A modifying enzyme PagP resulting in innate immune evasion in Yersinia pestis.
Investigations of Salmonella enterica serovar newport infections of oysters by using immunohistochemistry and knockout mutagenesis.
Yersinia pestis two-component gene regulatory systems promote survival in human neutrophils.
NleC, a type III secretion protease, compromises NF- $\kappa$ B activation by targeting p65/RelA.
Enteropathogenic E. coli non-LEE encoded effectors NleH1 and NleH2 attenuate NF- $\kappa$ B activation.
Yersinia adhesin A (YadA)--beauty & beast.
The Salmonella Effector SteD Mediates MARCH8-Dependent Ubiquitination of MHC II Molecules and Inhibits T Cell Activation.
Helicobacter and salmonella persistent infection strategies.
Enteropathogens: Tuning Their Gene Expression for Hassle-Free Survival.
Genome-based characterization of Escherichia coli causing bloodstream infection through next-generation sequencing.
Uropathogenic Escherichia coli-Associated Exotoxins.
YopN Is Required for Efficient Effector Translocation and Virulence in Yersinia pseudotuberculosis.
Molecular basis of uropathogenic Escherichia coli evasion of the innate immune response in the bladder.

Yersinia pestis Targets the Host Endosome Recycling Pathway during the Biogenesis of the Yersinia-Containing Vacuole To Avoid Killing by Macrophages.
Non-typhoidal Salmonella Typhimurium ST313 isolates that cause bacteremia in humans stimulate less inflammasome activation than ST19 isolates associated with gastroenteritis.
Immune evasion by Salmonella: exploiting the VPAC1/VIP axis in human monocytes.
Yersinia pestis: mechanisms of entry into and resistance to the host cell.
Broad-Spectrum Regulation of Nonreceptor Tyrosine Kinases by the Bacterial ADP-Ribosyltransferase EspJ.
Intracellular Shigella remodels its LPS to dampen the innate immune recognition and evade inflammasome activation.
Galectin-1-Driven Tolerogenic Programs Aggravate Yersinia enterocolitica Infection by Repressing Antibacterial Immunity.
Yersinia enterocolitica outer protein T (YopT).
DamX Controls Reversible Cell Morphology Switching in Uropathogenic Escherichia coli.
A rationally designed oral vaccine induces immunoglobulin A in the murine gut that directs the evolution of attenuated Salmonella variants.
Genomic Insights and Its Comparative Analysis with Yersinia enterocolitica Reveals the Potential Virulence Determinants and Further Pathogenicity for Foodborne Outbreaks.
Clinical pathogenesis of typhoid fever.
The pathogen-associated iroA gene cluster mediates bacterial evasion of lipocalin 2.
Insight into bacterial virulence mechanisms against host immune response via the Yersinia pestis-human protein-protein interaction network.
Yersinia signals macrophages to undergo apoptosis and YopJ is necessary for this cell death.
Identification and Characterization of the Nuclease Activity of the Extracellular Proteins from Salmonella enterica Serovar Typhimurium.
Burkholderia pseudomallei type III secreted protein BipC: role in actin modulation and translocation activities required for the bacterial intracellular lifecycle.
c-KIT signaling is targeted by pathogenic Yersinia to suppress the host immune response.
Lipid A 3'-O-deacylation by Salmonella outer membrane enzyme LpxR modulates the ability of lipid A to stimulate Toll-like receptor 4.
Yersinia versus host immunity: how a pathogen evades or triggers a protective response.
Seeing is understanding: Salmonella's way to penetrate the intestinal epithelium.
Deletion of Yersinia pestis ail Causes Temperature-Sensitive Pleiotropic Effects, Including Cell Lysis, That Are Suppressed by Carbon Source, Cations, or Loss of Phospholipase A Activity.
Yersinia pseudotuberculosis Blocks Neutrophil Degranulation.
A Salmonella Typhi RNA thermosensor regulates virulence factors and innate immune evasion in response to host temperature.
Strategies used by Yersinia enterocolitica to evade killing by the host: thinking beyond Yops.
Diversity of endotoxin and its impact on pathogenesis.
Contribution of toll-like receptors 2 and 4 in an oral Yersinia enterocolitica mouse infection model.
Selective culling of high avidity antigen-specific CD4+ T cells after virulent Salmonella infection.
Municipal Wastewater Surveillance Revealed a High Community Disease Burden of a Rarely Reported and Possibly Subclinical Salmonella enterica Serovar Derby Strain.
Biogenesis and function of the autotransporter adhesins YadA, intimin and invasins.
Virulence properties of asymptomatic bacteriuria Escherichia coli.
New insights into the bacterial fitness-associated mechanisms revealed by the characterization of large plasmids of an avian pathogenic E. coli.

Polymorphism in the Yersinia LcrV Antigen Enables Immune Escape From the Protection Conferred by an LcrV-Secreting Lactococcus Lactis in a Pseudotuberculosis Mouse Model.
Decreased potency of the Vibrio cholerae sheathed flagellum to trigger host innate immunity.
The serine protease Pic as a virulence factor of atypical enteropathogenic Escherichia coli.
A rapid change in virulence gene expression during the transition from the intestinal lumen into tissue promotes systemic dissemination of Salmonella.
Bacterial evasion of host immune defense: Yersinia enterocolitica encodes a suppressor for tumor necrosis factor alpha expression.
Enterococcus faecalis subverts and invades the host urothelium in patients with chronic urinary tract infection.
Translocation of enterohemorrhagic Escherichia coli effector Tir to the plasma membrane via host Golgi apparatus.
Active modification of host inflammation by Salmonella.
Biosynthesis, transport, and modification of lipid A.
Biochemical, structural and molecular dynamics analyses of the potential virulence factor RipA from Yersinia pestis.
Isolates of Salmonella typhimurium circumvent NLRP3 inflammasome recognition in macrophages during the chronic phase of infection.
Selectively reduced intracellular proliferation of Salmonella enterica serovar typhimurium within APCs limits antigen presentation and development of a rapid CD8 T cell response.
Transcriptional regulation of drug resistance mechanisms in Salmonella: where we stand and what we need to know.
Yersinia enterocolitica evasion of the host innate immune response by V antigen-induced IL-10 production of macrophages is abrogated in IL-10-deficient mice.
Rapid Isolation of intact Salmonella-containing vacuoles using paramagnetic nanoparticles.
Interleukin-10 induction is an important virulence function of the Yersinia pseudotuberculosis type III effector YopM.
Invasive Salmonella exploits divergent immune evasion strategies in infected and bystander dendritic cell subsets.
Unusual molecular architecture of the Yersinia pestis cytotoxin YopM: a leucine-rich repeat protein with the shortest repeating unit.
Characterization of differential Toll-like receptor responses below the optical diffraction limit.
S. Typhimurium strategies to resist killing by cationic antimicrobial peptides.
Sialic acid and N-acetylglucosamine Regulate type 1 Fimbriae Synthesis.
Enterococcus faecalis Promotes Innate Immune Suppression and Polymicrobial Catheter-Associated Urinary Tract Infection.
The Yersinia virulence effector YopM binds caspase-1 to arrest inflammasome assembly and processing.
Bacterial virulence factor inhibits caspase-4/11 activation in intestinal epithelial cells.
A hypervariable N-terminal region of Yersinia LcrV determines Toll-like receptor 2-mediated IL-10 induction and mouse virulence.
Role of Yersinia pestis toxin complex family proteins in resistance to phagocytosis by polymorphonuclear leukocytes.
Canine bacterial urinary tract infections: new developments in old pathogens.
Vibrio parahaemolyticus effector proteins suppress inflammasome activation by interfering with host autophagy signaling.



Peptide MSI-1 inhibited MCR-1 and regulated outer membrane vesicles to combat immune evasion of <i>Escherichia coli</i> .
Modulation of Rho GTPases and the actin cytoskeleton by YopT of <i>Yersinia</i> .
<i>Shigella flexneri</i> T3SS effector IpaH4.5 modulates the host inflammatory response via interaction with NF- $\kappa$ B p65 protein.
Autophagosomes can support <i>Yersinia pseudotuberculosis</i> replication in macrophages.
Suppression of NF- $\kappa$ B-mediated beta-defensin gene expression in the mammalian airway by the <i>Bordetella</i> type III secretion system.
Current activities of the <i>Yersinia</i> effector protein YopM.
Absence of intestinal PPAR $\gamma$ aggravates acute infectious colitis in mice through a lipocalin-2-dependent pathway.
<i>Yersinia enterocolitica</i> YopH-Deficient Strain Activates Neutrophil Recruitment to Peyer's Patches and Promotes Clearance of the Virulent Strain.
Virulence traits and expression of <i>bstA</i> , <i>fliC</i> and <i>sopE2</i> in <i>Salmonella</i> Dublin strains isolated from humans and animals in Brazil.
Nanoflow LC-MS Method Allowing In-Depth Characterization of Natural Heterogeneity of Complex Bacterial Lipopolysaccharides.
Subversion of human intestinal mucosa innate immunity by a Crohn's disease-associated <i>E. coli</i> .
A microfluidic-based genetic screen to identify microbial virulence factors that inhibit dendritic cell migration.
AlgW regulates multiple <i>Pseudomonas syringae</i> virulence strategies.
Critical role for <i>Salmonella</i> effector SopB in regulating inflammasome activation.
The human-bacterial pathogen protein interaction networks of <i>Bacillus anthracis</i> , <i>Francisella tularensis</i> , and <i>Yersinia pestis</i> .
<i>Yersinia pestis</i> activates both IL-1 $\beta$ and IL-1 receptor antagonist to modulate lung inflammation during pneumonic plague.
Spatiotemporal Monitoring of <i>Pseudomonas syringae</i> Effectors via Type III Secretion Using Split Fluorescent Protein Fragments.
Deciphering the acylation pattern of <i>Yersinia enterocolitica</i> lipid A.
Phosphomimetic Tyrosine Mutations in Spa47 Inhibit Type Three Secretion ATPase Activity and <i>Shigella</i> Virulence Phenotype.
Few Differences in Metabolic Network Use Found Between <i>Salmonella enterica</i> Colonization of Plants and Typhoidal Mice.
Features of urinary <i>Escherichia coli</i> isolated from children with complicated and uncomplicated urinary tract infections in Mexico.
An SopB-mediated immune escape mechanism of <i>Salmonella enterica</i> can be subverted to optimize the performance of live attenuated vaccine carrier strains.
The DNA adenine methylase of <i>Salmonella Enteritidis</i> promotes their intracellular replication by inhibiting arachidonic acid metabolism pathway in macrophages.
Bacteria and protozoa differentially modulate the expression of Rab proteins.
The Genetics of Enteropathogenic <i>Escherichia coli</i> Virulence.
YciR, a Specific 3'-Phosphodiesterase, Plays a Role in the Pathogenesis of Uropathogenic <i>Escherichia coli</i> CFT073.
A comparative study of the clonal diversity and virulence characteristics of uropathogenic <i>Escherichia coli</i> isolated from Australian and Turkish (Turkey) children and adults with urinary tract infections.
<i>Burkholderia pseudomallei</i> type III secretion system cluster 3 ATPase BsaS, a chemotherapeutic target for small-molecule ATPase inhibitors.

Escherichia coli isolated from bovine mastitis invade mammary cells by a modified endocytic pathway.
TcpA, a novel Yersinia ruckeri TIR-containing virulent protein mediates immune evasion by targeting MyD88 adaptors.
Immunity to Burkholderia pseudomallei.
Substances released from probiotic Lactobacillus rhamnosus GR-1 potentiate NF- $\kappa$ B activity in Escherichia coli-stimulated urinary bladder cells.
$\gamma\delta$ T cell IFN $\gamma$ production is directly subverted by Yersinia pseudotuberculosis outer protein YopJ in mice and humans.
[Effect of spvB/spvC gene on Salmonella virulence and the host immune function].
Serine protease autotransporters of Enterobacteriaceae (SPATEs) are largely distributed among Escherichia coli isolated from the bloodstream.
Characterization of Escherichia coli bloodstream isolates associated with mortality.
The O-Antigen Capsule of Salmonella enterica Serovar Typhimurium Facilitates Serum Resistance and Surface Expression of FliC.
Salmonella Typhi Colonization Provokes Extensive Transcriptional Changes Aimed at Evading Host Mucosal Immune Defense During Early Infection of Human Intestinal Tissue.
Local Generation of Kynurenines Mediates Inhibition of Neutrophil Chemotaxis by Uropathogenic Escherichia coli.
Whole genome sequencing reveals high clonal diversity of Escherichia coli isolated from patients in a tertiary care hospital in Moshi, Tanzania.
Structural and biological diversity of lipopolysaccharides from Burkholderia pseudomallei and Burkholderia thailandensis.
SDS-PAGE Analysis of the Outer Membrane Proteins of Uropathogenic Escherichia coli Isolated from Patients in Different Wards of Nemazee Hospital, Shiraz, Iran.
Uropathogenic Escherichia coli employs both evasion and resistance to subvert innate immune-mediated zinc toxicity for dissemination.
Identification and functional characterization of EseH, a new effector of the type III secretion system of Edwardsiella piscicida.
The Burkholderia pseudomallei type III secretion system and BopA are required for evasion of LC3-associated phagocytosis.
Peptidoglycan editing in non-proliferating intracellular Salmonella as source of interference with immune signaling.
Bacterial flagella: twist and stick, or dodge across the kingdoms.
Salmonella Coiled-Coil- and TIR-Containing TcpS Evades the Innate Immune System and Subdues Inflammation.
Siderophore vaccine conjugates protect against uropathogenic Escherichia coli urinary tract infection.
O-Acetylation of Capsular Polysialic Acid Enables Escherichia coli K1 Escaping from Siglec-Mediated Innate Immunity and Lysosomal Degradation of E. coli-Containing Vacuoles in Macrophage-Like Cells.
Adherent-invasive Escherichia coli blocks interferon- $\gamma$ -induced signal transducer and activator of transcription (STAT)-1 in human intestinal epithelial cells.
The Madagascar hissing cockroach as a novel surrogate host for Burkholderia pseudomallei, B. mallei and B. thailandensis.
Neutrophil antimicrobial proteins enhance Shigella flexneri adhesion and invasion.
Enterohemorrhagic and enteropathogenic Escherichia coli evolved different strategies to resist antimicrobial peptides.
Bacterial serine proteases secreted by the autotransporter pathway: classification, specificity, and role in virulence.

Induction of indoleamine 2,3-dioxygenase by uropathogenic bacteria attenuates innate responses to epithelial infection.
Expression of suppressor of cytokine signalling 3 (SOCS3) in human bladder epithelial cells infected with uropathogenic <i>Escherichia coli</i> .
<i>Granulibacter bethesdensis</i> , a Pathogen from Patients with Chronic Granulomatous Disease, Produces a Penta-Acylated Hypostimulatory Glycero-D-talo-oct-2-ulosonic Acid-Lipid A Glycolipid (Ko-Lipid A).
<i>Yersinia enterocolitica</i> induces apoptosis and inhibits surface molecule expression and cytokine production in murine dendritic cells.
Neuropathogenic <i>Escherichia coli</i> K1 does not exhibit proteolytic activities to exert its pathogenicity.
Pushing the envelope: LPS modifications and their consequences.
General Utilization of Fluorescent Polyisoprenoids with Sugar Selective Phosphoglycosyltransferases.
<i>Salmonella</i> in Swine: Microbiota Interactions.
<i>Salmonella enterica</i> serovar enteritidis antimicrobial peptide resistance genes aid in defense against chicken innate immunity, fecal shedding, and egg deposition.
2'-O-methylation within prokaryotic and eukaryotic tRNA inhibits innate immune activation by endosomal Toll-like receptors but does not affect recognition of whole organisms.
Influences of <i>Vibrio cholerae</i> Lipid A Types on LPS Bilayer Properties.
Dominant negative effects by inactive Spa47 mutants inhibit T3SS function and <i>Shigella</i> virulence.
Comparative Pathogenomics of <i>Escherichia coli</i> : Polyvalent Vaccine Target Identification through Virulome Analysis.
SaaS sRNA promotes <i>Salmonella</i> intestinal invasion via modulating MAPK inflammatory pathway.
Lipid A Variants Activate Human TLR4 and the Noncanonical Inflammasome Differently and Require the Core Oligosaccharide for Inflammasome Activation.
Murine Gammaherpesvirus 68 Pathogenesis Is Independent of Caspase-1 and Caspase-11 in Mice and Impairs Interleukin-1 $\beta$ Production upon Extrinsic Stimulation in Culture.
Natural history of <i>Yersinia pestis</i> pneumonia in aerosol-challenged BALB/c mice.
Induction of the immunoprotective coat of <i>Yersinia pestis</i> at body temperature is mediated by the Caf1R transcription factor.
Evading plant immunity: feedback control of the T3SS in <i>Pseudomonas syringae</i> .
Enteropathogenic <i>Escherichia coli</i> Tir recruits cellular SHP-2 through ITIM motifs to suppress host immune response.
Regulatory Evolution Drives Evasion of Host Inflammasomes by <i>Salmonella</i> Typhimurium.
Host transcriptomic responses to pneumonic plague reveal that <i>Yersinia pestis</i> inhibits both the initial adaptive and innate immune responses in mice.
Ferritinophagy-mediated iron competition in RUTs: Tug-of-war between UPEC and host.
Inhibition of <i>Salmonella</i> -induced apoptosis as a marker of the protective efficacy of virulence gene-deleted live attenuated vaccine.
<i>Yersinia</i> virulence factor YopJ acts as a deubiquitinase to inhibit NF-kappa B activation.
Leptospiral LPS escapes mouse TLR4 internalization and TRIF-associated antimicrobial responses through O antigen and associated lipoproteins.
Pic-Producing <i>Escherichia coli</i> Induces High Production of Proinflammatory Mediators by the Host Leading to Death by Sepsis.
Mechanisms of <i>Yersinia</i> YopO kinase substrate specificity.
<i>Salmonella</i> escapes adaptive immune response via SIRT2 mediated modulation of innate immune response in dendritic cells.
A bacterial cysteine protease effector protein interferes with photosynthesis to suppress plant innate immune responses.

Yersinia enterocolitica exploits different pathways to accomplish adhesion and toxin injection into host cells.
Feeding a high dosage of zinc oxide affects suppressor of cytokine gene expression in Salmonella Typhimurium infected piglets.
UPEC hemolysin: more than just for making holes.
Shigella evades pyroptosis by arginine ADP-ribosylation of caspase-11.
Shigella Pathogenesis: New Insights through Advanced Methodologies.
LI1035, a putative effector secreted by Lawsonia intracellularis, targets the MAPK pathway and regulates actin organization in yeast and mammalian cells.
Probiotic bacteria prevent Salmonella - induced suppression of lymphoproliferation in mice by an immunomodulatory mechanism.
Enterohemorrhagic Escherichia coli Tir inhibits TAK1 activation and mediates immune evasion.
Immune response to Burkholderia pseudomallei.
New Insights Into DAEC and EAEC Pathogenesis and Phylogeny.
Molecular mechanisms and implications for infection of lipopolysaccharide variation in Neisseria.
Involvement of signal regulatory protein $\alpha$ , a negative regulator of Toll-like receptor signaling, in impairing the MyD88-independent pathway and intracellular killing of Burkholderia pseudomallei-infected mouse macrophages.
A Single Bacterial Immune Evasion Strategy Dismantles Both MyD88 and TRIF Signaling Pathways Downstream of TLR4.
Structural insight into the bacterial mucinase StcE essential to adhesion and immune evasion during enterohemorrhagic E. coli infection.
Structure of CFA/I fimbriae from enterotoxigenic Escherichia coli.
Targeting Rac1 by the Yersinia effector protein YopE inhibits caspase-1-mediated maturation and release of interleukin-1 $\beta$ .
What's on the Outside Matters: The Role of the Extracellular Polymeric Substance of Gram-negative Biofilms in Evading Host Immunity and as a Target for Therapeutic Intervention.
Conditioned medium from enterohemorrhagic Escherichia coli-infected T84 cells inhibits signal transducer and activator of transcription 1 activation by gamma interferon.
Protecting against plague: towards a next-generation vaccine.
Identifying Yersinia YopH-targeted signal transduction pathways that impair neutrophil responses during in vivo murine infection.
Surface Anchoring of the Yersinia enterocolitica Galactan Is Dependent on the Lipopolysaccharide O-Antigen.
The polybasic region of Rho GTPases defines the cleavage by Yersinia enterocolitica outer protein T (YopT).
Experimental Persistent Infection of BALB/c Mice with Small-Colony Variants of Burkholderia pseudomallei Leads to Concurrent Upregulation of PD-1 on T Cells and Skewed Th1 and Th17 Responses.
The UPEC pore-forming toxin $\alpha$ -hemolysin triggers proteolysis of host proteins to disrupt cell adhesion, inflammatory, and survival pathways.
A Yersinia ruckeri TIR Domain-Containing Protein (STIR-2) Mediates Immune Evasion by Targeting the MyD88 Adaptor.
Characterization of Pyrin Dephosphorylation and Inflammasome Activation in Macrophages as Triggered by the Yersinia Effectors YopE and YopT.
Enterohemorrhagic Escherichia coli O157:H7 Shiga toxins inhibit gamma interferon-mediated cellular activation.
Yersinia effector YopO uses actin as bait to phosphorylate proteins that regulate actin polymerization.

Predominant phosphorylation patterns in <i>Neisseria meningitidis</i> lipid A determined by top-down MS/MS.
The multifaceted virulence of adherent-invasive <i>Escherichia coli</i> .
Genomic Modification of TonB and Emergence of Small-Colony Phenotype in VIM- and NDM-Producing <i>Escherichia coli</i> following Cefiderocol Exposure In Vitro.
Modulation of <i>Caenorhabditis elegans</i> immune response and modification of <i>Shigella</i> endotoxin upon interaction.
Structure of the N-terminal domain of <i>Yersinia pestis</i> YopH at 2.0 Å resolution.
PI3P Triggers ICAM-1 Degradation in <i>Shigella</i> Infected Cells, Thus Dampening Immune Cell Recruitment.
Characterization of transcriptional activities at a divergent promoter of the type VI secretion system in enterohemorrhagic <i>Escherichia coli</i> O157:H7.
<i>Vibrio</i> variations on a type three theme.
Toll-like receptor 6 drives differentiation of tolerogenic dendritic cells and contributes to LcrV-mediated plague pathogenesis.
Cellular distribution of lipid A and LPS R595 after in vitro application to isolated human monocytes by freeze-fracture replica immunogold-labelling.
Expression of IroN, the salmochelin siderophore receptor, requires mRNA activation by RyhB small RNA homologues.
Role of <i>Escherichia coli</i> endopeptidases and dd-carboxypeptidases in infection and regulation of innate immune response.
Identification of a novel lipopolysaccharide core biosynthesis gene cluster in <i>Bordetella pertussis</i> , and influence of core structure and lipid A glucosamine substitution on endotoxic activity.
Chlamydial Protease-Like Activity Factor and Type III Secreted Effectors Cooperate in Inhibition of p65 Nuclear Translocation.
<i>Shigella flexneri</i> infection: pathogenesis and vaccine development.
Vitamin D (1 $\alpha$ ,25(OH) <sub>2</sub> D <sub>3</sub> ) supplementation minimized multinucleated giant cells formation and inflammatory response during <i>Burkholderia pseudomallei</i> infection in human lung epithelial cells.
Survival and Intra-Nuclear Trafficking of <i>Burkholderia pseudomallei</i> : Strategies of Evasion from Immune Surveillance?
<i>Yersinia ruckeri</i> strain SC09 disrupts proinflammatory activation via Toll/IL-1 receptor-containing protein STIR-3.
Elucidation of the 3-O-deacylase gene, pagL, required for the removal of primary $\beta$ -hydroxy fatty acid from the lipid A in the nitrogen-fixing endosymbiont <i>Rhizobium etli</i> CE3.
Prevention and treatment of colitis with <i>Lactococcus lactis</i> secreting the immunomodulatory <i>Yersinia</i> LcrV protein.
Role of $\beta$ 1 integrins and bacterial adhesins for Yop injection into leukocytes in <i>Yersinia enterocolitica</i> systemic mouse infection.
Enterotoxigenic <i>Escherichia coli</i> Flagellin Inhibits TNF-Induced NF- $\kappa$ B Activation in Intestinal Epithelial Cells.
Porcine extraintestinal pathogenic <i>Escherichia coli</i> delivers two serine protease autotransporters coordinately optimizing the bloodstream infection.
A phosphoethanolamine transferase specific for the 4'-phosphate residue of <i>Cronobacter sakazakii</i> lipid A.
Genomic characterization of molecular markers associated with antimicrobial resistance and virulence of the prevalent <i>Campylobacter coli</i> isolated from retail chicken meat in the United Arab Emirates.

Occurrence of glycine in the core oligosaccharides of <i>Hafnia alvei</i> lipopolysaccharides--identification of disubstituted glycoform.
Identification of a bovine mastitis <i>Escherichia coli</i> subset.
Interdependence of <i>Shigella flexneri</i> O Antigen and Enterobacterial Common Antigen Biosynthetic Pathways.
Flashy flagella: flagellin modification is relatively common and highly versatile among the Enterobacteriaceae.
<i>Yersinia</i> effector protein (YopO)-mediated phosphorylation of host gelsolin causes calcium-independent activation leading to disruption of actin dynamics.
Evasion of MAIT cell recognition by the African <i>Salmonella</i> Typhimurium ST313 pathovar that causes invasive disease.
Five <i>Xanthomonas</i> type III effectors suppress cell death induced by components of immunity-associated MAP kinase cascades.
TcpC inhibits toll-like receptor signaling pathway by serving as an E3 ubiquitin ligase that promotes degradation of myeloid differentiation factor 88.
Pathogenomic analyses of <i>Shigella</i> isolates inform factors limiting shigellosis prevention and control across LMICs.
The molecular mechanism of species-specific recognition of lipopolysaccharides by the MD-2/TLR4 receptor complex.
<i>Shigella flexneri</i> suppresses NF- $\kappa$ B activation by inhibiting linear ubiquitin chain ligation.
Phosphatidylcholine in membrane of <i>Escherichia coli</i> changes bacterial antigenicity.
TcpC Inhibits M1 but Promotes M2 Macrophage Polarization via Regulation of the MAPK/NF- $\kappa$ B and Akt/STAT6 Pathways in Urinary Tract Infection.
<i>Shigella</i> : A Highly Virulent and Elusive Pathogen.
TcpC inhibits neutrophil extracellular trap formation by enhancing ubiquitination mediated degradation of peptidylarginine deiminase 4.
Substitution of the <i>Bordetella pertussis</i> lipid A phosphate groups with glucosamine is required for robust NF- $\kappa$ B activation and release of proinflammatory cytokines in cells expressing human but not murine Toll-like receptor 4-MD-2-CD14.
Mammalian Lipopolysaccharide Receptors Incorporated into the Retroviral Envelope Augment Virus Transmission.
Pathogenesis of <i>Proteus mirabilis</i> in Catheter-Associated Urinary Tract Infections.
Identification of the flagellin glycosylation system in <i>Burkholderia cenocepacia</i> and the contribution of glycosylated flagellin to evasion of human innate immune responses.
Route of entry and tissue distribution of <i>Yersinia ruckeri</i> in experimentally infected rainbow trout <i>Oncorhynchus mykiss</i> .
Virulence studies of <i>Enterobacter sakazakii</i> isolates associated with a neonatal intensive care unit outbreak.
Escape from immune surveillance by <i>Capnocytophaga canimorsus</i> .
Proteomic Adaptation of Australian Epidemic <i>Bordetella pertussis</i> .
<i>Echinococcus granulosus</i> cyst fluid suppresses inflammatory responses by inhibiting TRAF6 signalling in macrophages.
A plague of actin disassembly.
Key diffusion mechanisms involved in regulating bidirectional water permeation across <i>E. coli</i> outer membrane lectin.
[Regulation of the O-antigen polysaccharide chain length by Wzz- a review ].
Bacterial effector restricts liquid-liquid phase separation of ZPR1 to antagonize host UPR(ER).

A novel mechanism of bacterial toxin transfer within host blood cell-derived microvesicles.
Loss of specificity variants of WzxC suggest that substrate recognition is coupled with transporter opening in MOP-family flippases.
$\beta$ -arrestin 2 quenches TLR signaling to facilitate the immune evasion of EPEC.
DNA fingerprinting analysis of breakthrough outbreaks in vaccine-protected poultry stocks.
Phosphocholine decoration of <i>Proteus mirabilis</i> O18 LPS induces hydrophobicity of the cell surface and electrokinetic potential, but does not alter the adhesion to solid surfaces.
TNF $\alpha$ and IL-1 $\beta$ influence the differentiation and migration of murine MSCs independently of the NF- $\kappa$ B pathway.
In vivo monitoring of dynamic interaction between neutrophil and human umbilical cord blood-derived mesenchymal stem cell in mouse liver during sepsis.
cAMP receptor protein regulates mouse colonization, motility, fimbria-mediated adhesion, and stress tolerance in uropathogenic <i>Proteus mirabilis</i> .
Merging mythology and morphology: the multifaceted lifestyle of <i>Proteus mirabilis</i> .
Phosphocholine-Modified Lipooligosaccharides of <i>Haemophilus influenzae</i> Inhibit ATP-Induced IL-1 $\beta$ Release by Pulmonary Epithelial Cells.
Effects of a recombinant schistosomal-derived anti-inflammatory molecular (rSj16) on the lipopolysaccharide (LPS)-induced activated RAW264.7.
Copper tolerance in bacteria requires the activation of multiple accessory pathways.
<i>Aeromonas salmonicida</i> subsp. <i>salmonicida</i> in the light of its type-three secretion system.
A hydrocarbon ruler measures palmitate in the enzymatic acylation of endotoxin.
Inactivation of human coagulation factor X by a protease of the pathogen <i>Capnocytophaga canimorsus</i> .
Phenotype, Virulence and Immunogenicity of <i>Edwardsiella piscicida</i> Cyclic AMP Receptor Protein (Crp) Mutants in Catfish Host.
Modification of the structure of peptidoglycan is a strategy to avoid detection by nucleotide-binding oligomerization domain protein 1.
Mycobacterial PIMs inhibit host inflammatory responses through CD14-dependent and CD14-independent mechanisms.
<i>Capnocytophaga canimorsus</i> Sepsis Following a Minor Dog Bite to the Finger: Case Report.
Effect of zinc-treated <i>Entamoeba histolytica</i> on the human polymorphonuclear respiratory burst.

Table A2-55, Cluster 54

Cluster 54 focuses on influenza viruses, emphasizing the role of mutations in immune escape variants (643)
Deep Sequencing Reveals Potential Antigenic Variants at Low Frequencies in Influenza A Virus-Infected Humans.
Differential Recognition of Influenza A Viruses by M158-66 Epitope-Specific CD8 <sup>+</sup> T Cells Is Determined by Extraepitopic Amino Acid Residues.
Immune Escape Adaptive Mutations in Hemagglutinin Are Responsible for the Antigenic Drift of Eurasian Avian-Like H1N1 Swine Influenza Viruses.
Selection of Antigenically Advanced Variants of Influenza Viruses.
Immune Escape Variants of H9N2 Influenza Viruses Containing Deletions at the Hemagglutinin Receptor Binding Site Retain Fitness In Vivo and Display Enhanced Zoonotic Characteristics.
Software for Characterizing the Antigenic and Genetic Evolution of Human Influenza Viruses.

Universal immunity to influenza must outwit immune evasion.
Structural basis of influenza virus neutralization.
Characterization of a broadly neutralizing monoclonal antibody that targets the fusion domain of group 2 influenza A virus hemagglutinin.
A Perspective on the Structural and Functional Constraints for Immune Evasion: Insights from Influenza Virus.
Genetic Characterization of Influenza A (H1N1) Pandemic 2009 Virus Isolates from Mumbai.
A human antibody recognizing a conserved epitope of H5 hemagglutinin broadly neutralizes highly pathogenic avian influenza H5N1 viruses.
Pandemic influenza is a zoonosis, as it requires introduction of avian-like gene segments in the human population.
Immune responses to influenza virus infection.
Human Monoclonal Antibody Derived from Transchromosomal Cattle Neutralizes Multiple H1 Clades of Influenza A Virus by Recognizing a Novel Conformational Epitope in the Hemagglutinin Head Domain.
Rules of co-occurring mutations characterize the antigenic evolution of human influenza A/H3N2, A/H1N1 and B viruses.
Monitoring the antigenic evolution of human influenza A viruses to understand how and when viruses escape from existing immunity.
Sera from Individuals with Narrowly Focused Influenza Virus Antibodies Rapidly Select Viral Escape Mutations In Ovo.
Variation at Extra-epitopic Amino Acid Residues Influences Suppression of Influenza Virus Replication by M1(58-66) Epitope-Specific CD8(+) T Lymphocytes.
A non-VH1-69 heterosubtypic neutralizing human monoclonal antibody protects mice against H1N1 and H5N1 viruses.
Diversifying Selection Analysis Predicts Antigenic Evolution of 2009 Pandemic H1N1 Influenza A Virus in Humans.
Structural Consequences of Antigenic Variants of Human A/H3N2 Influenza Viruses.
Selection of antigenically advanced variants of seasonal influenza viruses.
The role of genomics in tracking the evolution of influenza A virus.
[Effect of mutations changing the antigenic specificity on the receptor-binding activity of the influenza virus hemagglutinin of H1 and H5 subtypes].
Molecular characterization of influenza viruses circulating in Northern Italy during two seasons (2005/2006 and 2006/2007) of low influenza activity.
Genetic characterization of seasonal influenza A (H3N2) viruses in Ontario during 2010-2011 influenza season: high prevalence of mutations at antigenic sites.
Memory T Cell Dynamics in the Lung during Influenza Virus Infection.
The Effect of I155T, K156Q, K156E and N186K Mutations in Hemagglutinin on the Virulence and Reproduction of Influenza A/H5N1 Viruses.
Analysis of antigenically important residues in human influenza A virus in terms of B-cell epitopes.
A Role of Influenza Virus Exposure History in Determining Pandemic Susceptibility and CD8+ T Cell Responses.
Conservation and diversity of influenza A H1N1 HLA-restricted T cell epitope candidates for epitope-based vaccines.
Influenza vaccine strain selection and recent studies on the global migration of seasonal influenza viruses.



[Prediction and evolution of B cell epitopes of hemagglutinin in human-infecting H6N1 avian influenza virus].
Glycosylation of Hemagglutinin and Neuraminidase of Influenza A Virus as Signature for Ecological Spillover and Adaptation among Influenza Reservoirs.
Glycan shielding of the influenza virus hemagglutinin contributes to immunopathology in mice.
Genetic characterization of an H5N1 avian influenza virus from a vaccinated duck flock in Vietnam.
Influenza H5 virus escape mutants: immune protection and antibody production in mice.
Dynamically correlated mutations drive human Influenza A evolution.
The evolution of seasonal influenza viruses.
Influenza 2009 pandemic: Cellular immunemediated surveillance modulated by TH17 & Tregs.
Evolution and dynamics of the pandemic H1N1 influenza hemagglutinin protein from 2009 to 2017.
The Influenza B Virus Hemagglutinin Head Domain Is Less Tolerant to Transposon Mutagenesis than That of the Influenza A Virus.
Generation of Escape Variants of Neutralizing Influenza Virus Monoclonal Antibodies.
Structure and Function Analysis of an Antibody Recognizing All Influenza A Subtypes.
A mutation in the HLA-B*2705-restricted NP383-391 epitope affects the human influenza A virus-specific cytotoxic T-lymphocyte response in vitro.
Molecular characterization and phylogenetic analysis of human influenza A viruses in three consecutive seasons with different epidemiological profiles.
Epitope mapping of the 2009 pandemic and the A/Brisbane/59/2007 seasonal (H1N1) influenza virus haemagglutinins using mAbs and escape mutants.
Protective immunity to lethal challenge of the 1918 pandemic influenza virus by vaccination.
Cross-clade antibody reactivity may attenuate the ability of influenza virus to evade the immune response.
HA gene amino acid mutations contribute to antigenic variation and immune escape of H9N2 influenza virus.
Estimating Vaccine-Driven Selection in Seasonal Influenza.
Universal or Specific? A Modeling-Based Comparison of Broad-Spectrum Influenza Vaccines against Conventional, Strain-Matched Vaccines.
Genotypic Variants of Pandemic H1N1 Influenza A Viruses Isolated from Severe Acute Respiratory Infections in Ukraine during the 2015/16 Influenza Season.
Selection and antigenic characterization of immune-escape mutants of H7N2 low pathogenic avian influenza virus using homologous polyclonal sera.
Substantial Antigenic Drift in the Hemagglutinin Protein of Swine Influenza A Viruses.
Stability of the Influenza Virus Hemagglutinin Protein Correlates with Evolutionary Dynamics.
Role of positive selection pressure on the evolution of H5N1 hemagglutinin.
Long intervals of stasis punctuated by bursts of positive selection in the seasonal evolution of influenza A virus.
Molecular characterization of a H5N1 highly pathogenic avian influenza virus clade 2.3.2.1b circulating in Vietnam in 2011.
Highly conserved influenza A sequences as T cell epitopes-based vaccine targets to address the viral variability.
Quantifying the impact of immune escape on transmission dynamics of influenza.
Human monoclonal antibodies broadly neutralizing against influenza B virus.
Effect of seasonal vaccination on the selection of influenza A/H3N2 epidemic variants.
Influenza Virus Vaccination Elicits Poorly Adapted B Cell Responses in Elderly Individuals.

Molecular characterization of the predominant influenza A(H1N1)pdm09 virus in Mexico, December 2011-February 2012.
Human Influenza A Virus Hemagglutinin Glycan Evolution Follows a Temporal Pattern to a Glycan Limit.
Phylogenetic analysis of the neuraminidase gene of pandemic H1N1 influenza A virus circulating in the South American region.
Measuring Site-specific Glycosylation Similarity between Influenza a Virus Variants with Statistical Certainty.
Amino Acid Residue 217 in the Hemagglutinin Glycoprotein Is a Key Mediator of Avian Influenza H7N9 Virus Antigenicity.
The molecular determinants of antigenic drift in a novel avian influenza A (H9N2) variant virus.
Contrasting substitution patterns between HA proteins of avian and human influenza viruses: Implication for monitoring human influenza epidemics.
Aflunov®: a prepandemic influenza vaccine.
Fitness costs limit influenza A virus hemagglutinin glycosylation as an immune evasion strategy.
Antibody pressure by a human monoclonal antibody targeting the 2009 pandemic H1N1 virus hemagglutinin drives the emergence of a virus with increased virulence in mice.
Antibodies against conserved antigens provide opportunities for reform in influenza vaccine design.
Influenza A virus hemagglutinin glycosylation compensates for antibody escape fitness costs.
Aberrant Cellular Glycosylation May Increase the Ability of Influenza Viruses to Escape Host Immune Responses through Modification of the Viral Glycome.
Defining influenza A virus hemagglutinin antigenic drift by sequential monoclonal antibody selection.
Genetic drift evolution under vaccination pressure among H5N1 Egyptian isolates.
Antigenic mapping of an H9N2 avian influenza virus reveals two discrete antigenic sites and a novel mechanism of immune escape.
Universally Immune: How Infection Permissive Next Generation Influenza Vaccines May Affect Population Immunity and Viral Spread.
Modeling the global transmission of antiviral-resistant influenza viruses.
Genetic and antigenic characterisation of influenza A(H3N2) viruses isolated in Yokohama during the 2016/17 and 2017/18 influenza seasons.
Evolution of the hemagglutinin protein of the new pandemic H1N1 influenza virus: maintaining optimal receptor binding by compensatory substitutions.
Limited Predictability of Amino Acid Substitutions in Seasonal Influenza Viruses.
Differences in the ease with which mutant viruses escape from human monoclonal antibodies against the HA stem of influenza A virus.
Mapping person-to-person variation in viral mutations that escape polyclonal serum targeting influenza hemagglutinin.
The molecular basis of antigenic variation among A(H9N2) avian influenza viruses.
Evaluation of recombinant influenza virus-simian immunodeficiency virus vaccines in macaques.
Molecular evolution of HA gene of the influenza A H1N1 pdm09 strain during the consecutive seasons 2009-2011 in Hangzhou, China: several immune-escape variants without positively selected sites.
Harnessing the Power of T Cells: The Promising Hope for a Universal Influenza Vaccine.
Genome Hotspots for Nucleotide Substitutions and the Evolution of Influenza A (H1N1) Human Strains.
Two Escape Mechanisms of Influenza A Virus to a Broadly Neutralizing Stalk-Binding Antibody.
Broadly Reactive IgG Responses to Heterologous H5 Prime-Boost Influenza Vaccination Are Shaped by Antigenic Relatedness to Priming Strains.

Hemagglutinin stalk-based monoclonal antibody elicits broadly reactivity against group 1 influenza A virus.
Structural and functional bases for broad-spectrum neutralization of avian and human influenza A viruses.
Impaired dendritic cell maturation in response to pandemic H1N109 influenza virus.
Inferring the antigenic epitopes for highly pathogenic avian influenza H5N1 viruses.
Recent avian influenza virus A/H5N1 evolution in vaccinated and unvaccinated poultry from farms in Southern Vietnam, January-March 2010.
Impact of cross-protective vaccines on epidemiological and evolutionary dynamics of influenza.
Epistatically interacting substitutions are enriched during adaptive protein evolution.
Pleiotropic effects of hemagglutinin amino acid substitutions of influenza A(H1N1)pdm09 virus escape mutants.
Preemptive priming readily overcomes structure-based mechanisms of virus escape.
Structural Basis for the Broad, Antibody-Mediated Neutralization of H5N1 Influenza Virus.
Functional constraints of influenza A virus epitopes limit escape from cytotoxic T lymphocytes.
Dynamic interactions of influenza viruses in Hong Kong during 1998-2018.
Selection pressure on the hemagglutinin gene of influenza A (H1N1) virus: adaptation to human and swine hosts in Asia.
Altering the Immunogenicity of Hemagglutinin Immunogens by Hyperglycosylation and Disulfide Stabilization.
Viral Capsid, Antibody, and Receptor Interactions: Experimental Analysis of the Antibody Escape Evolution of Canine Parvovirus.
Intra- and interhost evolutionary dynamics of equine influenza virus.
Use of antigenic cartography in vaccine seed strain selection.
Why Glycosylation Matters in Building a Better Flu Vaccine.
Epistasis reduces fitness costs of influenza A virus escape from stem-binding antibodies.
How single mutations affect viral escape from broad and narrow antibodies to H1 influenza hemagglutinin.
A universal influenza virus vaccine candidate confers protection against pandemic H1N1 infection in preclinical ferret studies.
Immune Escape Adaptive Mutations in the H7N9 Avian Influenza Hemagglutinin Protein Increase Virus Replication Fitness and Decrease Pandemic Potential.
Antigenic Drift of the Influenza A(H1N1)pdm09 Virus Neuraminidase Results in Reduced Effectiveness of A/California/7/2009 (H1N1pdm09)-Specific Antibodies.
Antigenic Characterization of Human Monoclonal Antibodies for Therapeutic Use against H7N9 Avian Influenza Virus.
Pre-existing immunity to influenza virus hemagglutinin stalk might drive selection for antibody-escape mutant viruses in a human challenge model.
Viral capsid, antibody, and receptor interactions: experimental analysis of the antibody escape evolution of canine parvovirus.
The N-linked glycosylation site at position 158 on the head of hemagglutinin and the virulence of H5N1 avian influenza virus in mice.
Isolation and complete genomic characterization of pandemic H1N1/2009 influenza viruses from Cuban swine herds.
Modulation of immunodominant sites in influenza hemagglutinin compromise antigenic variation and select receptor-binding variant viruses.

H3N2 Influenza Viruses with 12- or 16-Amino Acid Deletions in the Receptor-Binding Region of Their Hemagglutinin Protein.
Humanized antibodies with broad-spectrum neutralization to avian influenza virus H5N1.
Structure of the apo anti-influenza CH65 Fab.
The Highly Productive Thermothelomyces heterothallica C1 Expression System as a Host for Rapid Development of Influenza Vaccines.
Antibody epitopes on the neuraminidase of a recent H3N2 influenza virus (A/Memphis/31/98).
Complete mapping of viral escape from neutralizing antibodies.
Natural and directed antigenic drift of the H1 influenza virus hemagglutinin stalk domain.
Prevalence of epistasis in the evolution of influenza A surface proteins.
To dream the impossible dream: universal influenza vaccination.
Re-emergence of H3N2 strains carrying potential neutralizing mutations at the N-linked glycosylation site at the hemagglutinin head, post the 2009 H1N1 pandemic.
Immunity-induced criticality of the genotype network of influenza A (H3N2) hemagglutinin.
Immune escape mutants of Highly Pathogenic Avian Influenza H5N1 selected using polyclonal sera: identification of key amino acids in the HA protein.
Viral Fitness and Antigenic Determinants of Porcine Parvovirus at the Amino Acid Level of the Capsid Protein.
Regulation of antinucleoprotein IgG by systemic vaccination and its effect on influenza virus clearance.
Pandemic Avian Influenza and Intra/Interhaemagglutinin Subtype Electrostatic Variation among Viruses Isolated from Avian, Mammalian, and Human Hosts.
Improving Statistical Certainty of Glycosylation Similarity between Influenza A Virus Variants Using Data-Independent Acquisition Mass Spectrometry.
Phylogenetic and evolutionary history of influenza B viruses, which caused a large epidemic in 2011-2012, Taiwan.
Contribution of virus-specific CD8+ cytotoxic T cells to virus clearance or pathologic manifestations of influenza virus infection in a T cell receptor transgenic mouse model.
[Epitope analysis of the hemagglutinin molecule of the Victoria lineage influenza B viruses].
Monitoring genetic diversity of influenza A(H1N1)pdm09 virus circulating during the post-pandemic period in Turkey.
Hepatitis C virus with a naturally occurring single amino-acid substitution in the E2 envelope protein escapes neutralization by naturally-induced and vaccine-induced antibodies.
The genomic rate of molecular adaptation of the human influenza A virus.
Identification of epistatic mutations and insights into the evolution of the influenza virus using a mass-based protein phylogenetic approach.
Antibody Immunodominance: The Key to Understanding Influenza Virus Antigenic Drift.
Understanding Antibody Responses in Early Life: Baby Steps towards Developing an Effective Influenza Vaccine.
A computational analysis of the antigenic properties of haemagglutinin in influenza A H3N2.
Multiple clusters of A(H1N1)pdm09 virus circulating in severe cases of influenza during the 2010-2011 season: a phylogenetic and molecular analysis of the neuraminidase gene.
Heterosubtypic immune pressure accelerates emergence of influenza A virus escape phenotypes in mice.
The 2009 H1N1 pandemic influenza virus: what next?
Viral safety of blood derivatives by immune neutralization.

Acute emergence and reversion of influenza A virus quasispecies within CD8+ T cell antigenic peptides.
Mutation of D201G near the receptor binding site significantly drives antigenic drift of circulating H9N2 subtype avian influenza virus.
Selective pressure to increase charge in immunodominant epitopes of the H3 hemagglutinin influenza protein.
New Insights into the Generation of CD4 Memory May Shape Future Vaccine Strategies for Influenza.
Mapping the antigenic and genetic evolution of influenza virus.
Population implications of the deployment of novel universal vaccines against epidemic and pandemic influenza.
Mutations in the haemagglutinin protein and their effect in transmission of highly pathogenic avian influenza (HPAI) H5N1 virus in sub-optimally vaccinated chickens.
Effects of hemagglutinin amino acid substitutions in H9 influenza A virus escape mutants.
Influenza Vaccine Effectiveness: Defining the H3N2 Problem.
Potential recombinant vaccine against influenza A virus based on M2e displayed on nodaviral capsid nanoparticles.
Hemagglutinin 222D/G polymorphism facilitates fast intra-host evolution of pandemic (H1N1) 2009 influenza A viruses.
Cross-immunity Against Avian Influenza A(H7N9) Virus in the Healthy Population Is Affected by Antigenicity-Dependent Substitutions.
Broadly neutralizing influenza hemagglutinin stem-specific antibody CR8020 targets residues that are prone to escape due to host selection pressure.
Host Adaptive Evolution of Avian-Origin H3N2 Canine Influenza Virus.
Mapping an Adeno-associated Virus 9-Specific Neutralizing Epitope To Develop Next-Generation Gene Delivery Vectors.
N-linked glycosylation in the hemagglutinin of influenza A viruses.
A complex epistatic network limits the mutational reversibility in the influenza hemagglutinin receptor-binding site.
Adaptive evolution and environmental durability jointly structure phylodynamic patterns in avian influenza viruses.
Comparative structural analysis of haemagglutinin proteins from type A influenza viruses: conserved and variable features.
The emergence of new antigen branches of H9N2 avian influenza virus in China due to antigenic drift on hemagglutinin through antibody escape at immunodominant sites.
In Vivo Therapy with M2e-Specific IgG Selects for an Influenza A Virus Mutant with Delayed Matrix Protein 2 Expression.
Diversity of Functionally Permissive Sequences in the Receptor-Binding Site of Influenza Hemagglutinin.
Subclade 2.2.1-Specific Human Monoclonal Antibodies That Recognize an Epitope in Antigenic Site A of Influenza A(H5) Virus HA Detected between 2015 and 2018.
The evolutionary dynamics of receptor binding avidity in influenza A: a mathematical model for a new antigenic drift hypothesis.
Electrostatic Variation of Haemagglutinin as a Hallmark of the Evolution of Avian Influenza Viruses.
Influenza A gradual and epochal evolution: insights from simple models.
N-linked glycans on influenza A H3N2 hemagglutinin constrain binding of host antibodies, but shielding is limited.
Glycosylation at Asn91 of H1N1 haemagglutinin affects binding to glycan receptors.

Characterization of epitopes on the rabies virus glycoprotein by selection and analysis of escape mutants.
Antigenic Determinants of Possible Vaccine Escape by Porcine Circovirus Subtype 2b Viruses.
Vaccination with inactivated virus against low pathogenic avian influenza subtype H9N2 does not prevent virus transmission in chickens.
Prevalence and Significance of Substitutions in the Fusion Protein of Respiratory Syncytial Virus Resulting in Neutralization Escape From Antibody MEDI8897.
Escape of pathogens from the host immune response by mutations and mimicry. Possible means to improve vaccine performance.
Antibody responses against wild-type yellow fever virus and the 17D vaccine strain: characterization with human monoclonal antibody fragments and neutralization escape variants.
Structural considerations in the fitness landscape of a virus.
Comparison of influenza virus replication fidelity in vitro using selection pressure with monoclonal antibodies.
Modelling the role of immunity in reversion of viral antigenic sites.
Emergence of novel avian origin H7N9 viruses after introduction of H7-Re3 and rLN79 vaccine strains to China.
Passage in egg culture is a major cause of apparent positive selection in influenza B hemagglutinin.
Non-random escape pathways from a broadly neutralizing human monoclonal antibody map to a highly conserved region on the hepatitis C virus E2 glycoprotein encompassing amino acids 412-423.
Novel Mutations Evading Avian Immunity around the Receptor Binding Site of the Clade 2.3.2.1c Hemagglutinin Gene Reduce Viral Thermostability and Mammalian Pathogenicity.
Role of vaccination-induced immunity and antigenic distance in the transmission dynamics of highly pathogenic avian influenza H5N1.
Evolution in the influenza A H3 stalk - a challenge for broad-spectrum vaccines?
Multiplexed screening of natural humoral immunity identifies antibodies at fine specificity for complex and dynamic viral targets.
Bird to human transmission biases and vaccine escape mutants in H5N1 infections.
Avian influenza--a pandemic waiting to happen?
Human Monoclonal Antibodies Potently Neutralize Zika Virus and Select for Escape Mutations on the Lateral Ridge of the Envelope Protein.
Large-scale evolutionary surveillance of the 2009 H1N1 influenza A virus using resequencing arrays.
Poultry vaccination directed evolution of H9N2 low pathogenicity avian influenza viruses in Korea.
Molecular dynamics analysis of antibody recognition and escape by human H1N1 influenza hemagglutinin.
New Immunoinformatics Tools for Swine: Designing Epitope-Driven Vaccines, Predicting Vaccine Efficacy, and Making Vaccines on Demand.
GP5 of porcine reproductive and respiratory syndrome virus (PRRSV) as a target for homologous and broadly neutralizing antibodies.
[The impact of conservative and hypervariable immunodominant epitopes in internal proteins of the influenza A virus on cytotoxic T-cell immune responses].
Emergence of new influenza A viruses which carry an escape mutation of the HLA-B27-restricted CTL epitope of NP in Japan.
Increasing Efficacy of Enveloped Whole-Virus Vaccines by In situ Immune-Complexing with the Natural Anti-Gal Antibody.
Neuraminidase receptor binding variants of human influenza A(H3N2) viruses resulting from substitution of aspartic acid 151 in the catalytic site: a role in virus attachment?

A Single Mutation N166D in Hemagglutinin Affects Antigenicity and Pathogenesis of H9N2 Avian Influenza Virus.
Immune escape by human immunodeficiency virus type 1 from neutralizing antibodies: evidence for multiple pathways.
Antigenic evolution of viruses in host populations.
Influenza A virus hemagglutinin antibody escape promotes neuraminidase antigenic variation and drug resistance.
Flexibility in surface-exposed loops in a virus capsid mediates escape from antibody neutralization.
Antigenic site variation in foot-and-mouth disease virus serotype O grown under vaccinal serum antibodies in vitro.
Identification of a conserved linear neutralizing epitope recognized by monoclonal antibody 9A9 against serotype A foot-and-mouth disease virus.
Dynamics of glycoprotein charge in the evolutionary history of human influenza.
Epistatic Interactions within the Influenza A Virus Polymerase Complex Mediate Mutagen Resistance and Replication Fidelity.
Structural basis of HCV neutralization by human monoclonal antibodies resistant to viral neutralization escape.
Diversification of specificity after maturation of the antibody response to the HIV gp41 epitope ELDKWA.
Pandemic influenza A viruses escape from restriction by human MxA through adaptive mutations in the nucleoprotein.
Molecular Evolution of Attachment Glycoprotein (G) and Fusion Protein (F) Genes of Respiratory Syncytial Virus ON1 and BA9 Strains in Xiamen, China.
Dynamic immunodominance hierarchy of neutralizing antibody responses to evolving GII.4 noroviruses.
Antigenic Drift Defines a New D4 Subgenotype of Measles Virus.
Viruses selectively mutate their CD8+ T-cell epitopes--a large-scale immunomic analysis.
Live Attenuated Influenza Vaccines engineered to express the nucleoprotein of a recent isolate stimulate human influenza CD8(+) T cells more relevant to current infections.
Recent structural solutions for antibody neutralization of viruses.
Longitudinal Antigenic Sequences and Sites from Intra-Host Evolution (LASSIE) Identifies Immune-Selected HIV Variants.
Naturally occurring mutations within 39 amino acids in the envelope glycoprotein of maedi-visna virus alter the neutralization phenotype.
Mutational analysis of a principal neutralization domain of visna/maedi virus envelope glycoprotein.
Loss of antigenic epitopes as the result of env gene recombination in retrovirus-induced leukemia in immunocompetent mice.
Homology modeling study toward identifying structural properties in the HA2 B-loop that would influence the HA1 receptor-binding site.
Quantifying selection and diversity in viruses by entropy methods, with application to the haemagglutinin of H3N2 influenza.
Isolation and genetic characterization of a novel 2.2.1.2a H5N1 virus from a vaccinated meat-turkeys flock in Egypt.
Identification of genetic diversity by cultivating influenza A(H3N2) virus in vitro in the presence of post-infection sera from small children.
Isolation of highly pathogenic avian influenza H5N1 from table eggs after vaccinal break in commercial layer flock.

Analysis of Amino Acid Changes in the Fusion Protein of Virulent Newcastle Disease Virus from Vaccinated Poultry in Nigerian Isolates.
Experimental challenge of chicken vaccinated with commercially available H5 vaccines reveals loss of protection to some highly pathogenic avian influenza H5N1 strains circulating in Hong Kong/China.
Parallel T Cell Immunogenic Regions in Influenza B and A Viruses with Distinct Nuclear Export Signal Functions: The Balance between Viral Life Cycle and Immune Escape.
Neutralization of zoonotic retroviruses by human antibodies: Genotype-specific epitopes within the receptor-binding domain from simian foamy virus.
Intrahost evolutionary dynamics of canine influenza virus in naive and partially immune dogs.
Potent antibody-mediated neutralization and evolution of antigenic escape variants of simian immunodeficiency virus strain SIVmac239 in vivo.
Emerging of H5N6 Subtype Influenza Virus with 129-Glycosylation Site on Hemagglutinin in Poultry in China Acquires Immune Pressure Adaptation.
Implications of localized charge for human influenza A H1N1 hemagglutinin evolution: Insights from deep mutational scans.
Investigating Substitutions in Antibody-Antigen Complexes Using Molecular Dynamics: A Case Study with Broad-spectrum, Influenza A Antibodies.
Assessing the effects of a two-amino acid flexibility in the Hemagglutinin 220-loop receptor-binding domain on the fitness of Influenza A(H9N2) viruses.
Structural Insights into Reovirus $\sigma 1$ Interactions with Two Neutralizing Antibodies.
Molecular epidemiology of echovirus 30: temporal circulation and prevalence of single lineages.
Adaptive evolution during the establishment of European avian-like H1N1 influenza A virus in swine.
Epidemic dynamics and antigenic evolution in a single season of influenza A.
A comprehensive analysis of reassortment in influenza A virus.
Influence of maternal immunity on vaccine efficacy and susceptibility of one day old chicks against Egyptian highly pathogenic avian influenza H5N1.
Modeling Infectious Bursal Disease Virus (IBDV) Antigenic Drift In Vitro.
Single mutation induced H3N2 hemagglutinin antibody neutralization: a free energy perturbation study.
A Deep Learning Approach for Predicting Antigenic Variation of Influenza A H3N2.
The N-terminal region of the human papillomavirus L2 protein contains overlapping binding sites for neutralizing, cross-neutralizing and non-neutralizing antibodies.
Limited naturally occurring escape in broadly neutralizing antibody epitopes in hepatitis C glycoprotein E2 and constrained sequence usage in acute infection.
Evolution of an HIV glycan-dependent broadly neutralizing antibody epitope through immune escape.
Glycosylation as a tool for rational vaccine design.
Optimal viral immune surveillance evasion strategies.
Antigenic variation in hemagglutinin-neuraminidase of Newcastle disease virus isolated from Tibet, China.
Peptide Antiviral Strategies as an Alternative to Treat Lower Respiratory Viral Infections.
Understanding the complex evolution of rapidly mutating viruses with deep sequencing: Beyond the analysis of viral diversity.
Deceptive imprinting and immune refocusing in vaccine design.
Envelope determinants of equine lentiviral vaccine protection.
B-Cell Memory Responses to Variant Viral Antigens.
T cell memory to evolutionarily conserved and shared hemagglutinin epitopes of H1N1 viruses: a pilot scale study.



Porcine circovirus type 2 (PCV2) evolution before and after the vaccination introduction: A large scale epidemiological study.
Viral proteome size and CD8+ T cell epitope density are correlated: the effect of complexity on selection.
Modification of the Spike Protein for Vaccines against Enveloped RNA Viruses.
High mutant frequency in populations of a DNA virus allows evasion from antibody therapy in an immunodeficient host.
Whole genome and in-silico analyses of G1P[8] rotavirus strains from pre- and post-vaccination periods in Rwanda.
Evidence for positive selection of hepatitis A virus antigenic variants in vaccinated men-having-sex-with men patients: Implications for immunization policies.
Do commercial avian influenza H5 vaccines induce cross-reactive antibodies against contemporary H5N1 viruses in Egypt?
T cell epitope clustering in the highly immunogenic BZLF1 antigen of Epstein-Barr virus.
Geometric Constraints Dominate the Antigenic Evolution of Influenza H3N2 Hemagglutinin.
A highly divergent antigenic site of foot-and-mouth disease virus retains its immunodominance.
Characterization of a Novel Conformational GII.4 Norovirus Epitope: Implications for Norovirus-Host Interactions.
Mutations in foot and mouth disease virus types A and O isolated from vaccinated animals.
Evolutionary Analysis and Prediction of Peptide Vaccine Candidates for Foot-and-Mouth-Disease Virus Types A and O in Bangladesh.
Stereophysicochemical variability plots highlight conserved antigenic areas in Flaviviruses.
Linear B-Cell Epitopes in Human Norovirus GII.4 Capsid Protein Elicit Blockade Antibodies.
Deconstructing the Antiviral Neutralizing-Antibody Response: Implications for Vaccine Development and Immunity.
A role for carbohydrates in immune evasion in AIDS.
Into the Deep (Sequence) of the Foot-and-Mouth Disease Virus Gene Pool: Bottlenecks and Adaptation during Infection in Naïve and Vaccinated Cattle.
Conformational Occlusion of Blockade Antibody Epitopes, a Novel Mechanism of GII.4 Human Norovirus Immune Evasion.
Host contact structure is important for the recurrence of Influenza A.
Sequencing of emerging canine distemper virus strain reveals new distinct genetic lineage in the United States associated with disease in wildlife and domestic canine populations.
Selection and characterization of human respiratory syncytial virus escape mutants resistant to a polyclonal antiserum raised against the F protein.
Generation and characterization of a protective mouse monoclonal antibody against human enterovirus 71.
Glycan masking in vaccine design: Targets, immunogens and applications.
Genetic variability of measles virus in acute and persistent infections.
Genomic evolution in a virus under specific selection for host recognition.
Potential Novel N-Glycosylation Patterns Associated with the Emergence of New Genetic Variants of PRRSV-2 in the U.S.
T-cell epitopes predicted from the Nucleocapsid protein of Sin Nombre virus restricted to 30 HLA alleles common to the North American population.
Molecular characterization of the monoclonal antibodies composing ZMAb: a protective cocktail against Ebola virus.
Virus glycosylation: role in virulence and immune interactions.

Antigenic evolution of vaccine-derived polioviruses: changes in individual epitopes and relative stability of the overall immunological properties.
Antigenic heterogeneity of capsid protein VP1 in foot-and-mouth disease virus (FMDV) serotype Asia 1.
Quasispecies made simple.
Peek-a-boo: membrane hijacking and the pathogenesis of viral hepatitis.
Oncolytic herpes simplex virus tumor targeting and neutralization escape by engineering viral envelope glycoproteins.
Monitoring Viral Genetic Variation as a Tool To Improve Molecular Diagnostics for Mumps Virus.
Mechanism of Cross-Resistance to Fusion Inhibitors Conferred by the K394R Mutation in Respiratory Syncytial Virus Fusion Protein.
Experimental analysis of sources of error in evolutionary studies based on Roche/454 pyrosequencing of viral genomes.
Overcoming antigenic diversity and improving vaccines using DNA shuffling and screening technologies.
Mutation or loss of Wilms' tumor gene 1 (WT1) are not major reasons for immune escape in patients with AML receiving WT1 peptide vaccination.
Dengue virus neutralization is modulated by IgG antibody subclass and Fcγ receptor subtype.
A decade of sustained selection pressure on two surface sites of the VP1 protein of Enterovirus A71 suggests that immune evasion may be an indirect driver for virulence.
Persistent BK Polyomavirus Viruria is Associated with Accumulation of VP1 Mutations and Neutralization Escape.
Limited inter- and intra-patient sequence diversity of the genetic lineage A human metapneumovirus fusion gene.
Predominance and first complete genomic characterization of canine parvovirus 2b in Turkey.
The hypervariable region of Anaplasma marginale major surface protein 2 (MSP2) contains multiple immunodominant CD4+ T lymphocyte epitopes that elicit variant-specific proliferative and IFN-γ responses in MSP2 vaccinates.
JCPyV VP1 Mutations in Progressive Multifocal Leukoencephalopathy: Altering Tropism or Mediating Immune Evasion?
Neutralization interfering antibodies: a "novel" example of humoral immune dysfunction facilitating viral escape?
Infant antibody and B-cell responses following confirmed pediatric GII.17 norovirus infections functionally distinguish GII.17 genetic clusters.
Hepacivirus A Infection in Horses Defines Distinct Envelope Hypervariable Regions and Elucidates Potential Roles of Viral Strain and Adaptive Immune Status in Determining Envelope Diversity and Infection Outcome.
The 2.8 Å Electron Microscopy Structure of Adeno-Associated Virus-DJ Bound by a Heparinoid Pentasaccharide.
Neoantigen Cancer Vaccines: Real Opportunity or Another Illusion?
Co-circulation of dengue virus serotypes in Central India: Evidence of prolonged viremia in DENV-2.
Population Impact and Effectiveness of Monovalent Rotavirus Vaccination in Urban Malawian Children 3 Years After Vaccine Introduction: Ecological and Case-Control Analyses.
Silencing and activating anergic B cells.
Human lymph node immune dynamics as driver of vaccine efficacy: an understudied aspect of immune responses.
Sequential mutations associated with adaptation of human cytomegalovirus to growth in cell culture.

Combining genomics and epidemiology to track mumps virus transmission in the United States.
Filovirus Neutralising Antibodies: Mechanisms of Action and Therapeutic Application.
Molecular analysis of rearranged VH genes during B cell chronic lymphocytic leukemia: intraclonal stability is frequent but not constant.
Genetic analysis of bovine respiratory syncytial virus in Croatia.
Multiple sites in the N-terminal half of simian immunodeficiency virus capsid protein contribute to evasion from rhesus monkey TRIM5 $\alpha$ -mediated restriction.
Full-Genome Sequence of Infectious Laryngotracheitis Virus (Gallid Alphaherpesvirus 1) Strain VFAR-043, Isolated in Peru.
N-Glycans Mediate the Ebola Virus-GP1 Shielding of Ligands to Immune Receptors and Immune Evasion.
Evidence of natural co-infection with PCV2b subtypes in vivo.
Near-Complete Genome Sequences of Measles Virus Strains from 10 Years of Uganda Country-wide Surveillance.
Insights into immune evasion of human metapneumovirus: novel 180- and 111-nucleotide duplications within viral G gene throughout 2014-2017 seasons in Barcelona, Spain.
Origin, genetic diversity, adaptive evolution and transmission dynamics of Getah virus.
Antibody-mediated depletion of immunosuppressive factors from ovarian carcinoma-associated ascites for investigation of paracrine versus autocrine effects.
Rapid emergence and predominance of a broadly recognizing and fast-evolving norovirus GII.17 variant in late 2014.
Insights From Analysis of Human Antigen-Specific Memory B Cell Repertoires.
Molecular basis of clonal evolution in multiple myeloma.
Molecular epidemiological study on Infectious Pancreatic Necrosis Virus isolates from aquafarms in Scotland over three decades.
[Study of molecular epidemiology and genetic diversity of human bocvirus in children with respiratory tract infection].
A detailed comparative analysis of codon usage bias in Alongshan virus.
Phylogenetic analysis and structural predictions of human adenovirus penton proteins as a basis for tissue-specific adenovirus vector design.
Genome sequence of an unusual human G10P[8] rotavirus detected in Vietnam.
Geographically variable selection in Ambystoma tigrinum virus (Iridoviridae) throughout the western USA.
JC polyomavirus mutants escape antibody-mediated neutralization.
Modeling of the human rhinovirus C capsid suggests a novel topography with insights on receptor preference and immunogenicity.
Expression of major surface protein 2 antigenic variants during acute Anaplasma marginale rickettsemia.
Evaluation of a multi-epitope subunit vaccine against avian leukosis virus subgroup J in chickens.
Molecular epidemiology analysis of symptomatic and asymptomatic norovirus infections in Chinese infants.
Deciphering the genetic landscape of pulmonary lymphomas.
Genetic variability in the major capsid L1 protein of human papillomavirus type 16 (HPV-16) and 18 (HPV-18).
Reconstruction of viral population structure from next-generation sequencing data using multicommodity flows.

Chikungunya virus neutralization antigens and direct cell-to-cell transmission are revealed by human antibody-escape mutants.
Oligomannose N-Glycans 3D Architecture and Its Response to the FcγRIIIa Structural Landscape.
In vivo studies on the effect of co-encapsulation of CpG DNA and antigen in acid-degradable microparticle vaccines.
A Norovirus Uses Bile Salts To Escape Antibody Recognition While Enhancing Receptor Binding.
The possible impact of novel mutations in human papillomavirus 52 on the infection characteristics.
Glycosylation of ALV-J Envelope Protein at Sites 17 and 193 Is Pivotal in the Virus Infection.
[Treatment strategies for multiple myeloma based on molecular pathogenesis].
Monkeypox virus (MPXV) genomics: A mutational and phylogenomic analyses of B.1 lineages.
Origin, antigenicity, and function of a secreted form of ORF2 in hepatitis E virus infection.
Natural selection differences detected in key protein domains between non-pathogenic and pathogenic feline coronavirus phenotypes.
Tracing the Genetic Evolution of Canine Parvovirus Type 2 (CPV-2) in Thailand.
Molecular characterization and clinical epidemiology of human respiratory syncytial virus (HRSV) A and B in hospitalized children, Southern Brazil.
Norovirus in healthcare settings.
Human-monoclonal-antibody therapy protects nonhuman primates against advanced Lassa fever.
Peptide cross-reactivity: the original sin of vaccines.
Tetramer Immunization and Selection Followed by CELLISA Screening to Generate Monoclonal Antibodies against the Mouse Cytomegalovirus m12 Immuno-evasin.
Rotavirus A Genome Segments Show Distinct Segregation and Codon Usage Patterns.
The murine gammaherpesvirus-68 gp150 acts as an immunogenic decoy to limit virion neutralization.
Molecular Characterization of BK Polyomavirus Replication in Allogeneic Hematopoietic Cell Transplantation Patients.
PVS: a web server for protein sequence variability analysis tuned to facilitate conserved epitope discovery.
Structural and phylogenetic analysis of adenovirus hexons by use of high-resolution x-ray crystallographic, molecular modeling, and sequence-based methods.
Analysis of genetic variation in human papillomavirus type 16 E1 and E2 in women with cervical infection in Xinjiang, China.
Frequency evaluation and molecular characterization of HHV-6 and HHV-7 among children under 5 years with fever and skin rash.
[Genome cloning and phylogenetic analysis of human bocavirus capsid gene].
Genetic variations in E6, E7 and the long control region of human papillomavirus type 16 among patients with cervical lesions in Xinjiang, China.
Hsp90 shapes protein and RNA evolution to balance trade-offs between protein stability and aggregation.
Diversity of Human Enterovirus Co-Circulations in Five Kindergartens in Bangkok between July 2019 and January 2020.
Genetic features of bovine viral diarrhoea virus subgenotype 1c in newborn calves at nucleotide and synonymous codon usages.
Optimizing DNA transduction by selection of mutations that evade bacterial defense systems.
Self-tolerance curtails the B cell repertoire to microbial epitopes.
Comprehensive analysis of the overall codon usage patterns in equine infectious anemia virus.
Genome-wide analysis of synonymous codon usage in Huaiyangshan virus and other bunyaviruses.

Lessons in nonhuman primate models for AIDS vaccine research: from minefields to milestones.
An HSV-2 Trivalent Vaccine Is Immunogenic in Rhesus Macaques and Highly Efficacious in Guinea Pigs.
Functional conservation of HTLV-1 rex balances the immune pressure for sequence variation in the rex gene.
Mutations of MAP2K1 are frequent in pediatric-type follicular lymphoma and result in ERK pathway activation.
High Genotypic Diversity, Putative New Types and Intra-Genotype Variants of Bovine Papillomavirus in Northeast Brazil.
Type-specific epitope locations revealed by X-ray crystallographic study of adenovirus type 5 hexon.
Efficient error correction for next-generation sequencing of viral amplicons.
Polio type 2 and 3 eradication: Relevance to the immunity status of individuals living in Germany, 2005-2020.
Lessons from mouse models of MPN.
Analysis of TET2 mutations in paroxysmal nocturnal hemoglobinuria (PNH).
The current toolbox for APOBEC drug discovery.
Emergence of clonal hematopoiesis in the majority of patients with acquired aplastic anemia.
Quantitative analysis of an anti-viral immune escape compound ML-7 in feline plasma using ultra performance liquid chromatography/electrospray ionization mass spectrometry.
The face of Ebola: changing frequency of haemorrhage in the West African compared with Eastern-Central African outbreaks.

Table A2-56, Cluster 55

Cluster 55 focuses on Schistosoma mansoni and other helminth parasites, emphasizing evasion of host immunity during infection (537)
[Proteases in helminthic parasites].
A combined proteomic and immunologic approach for the analysis of Schistosoma mansoni cercariae and adult worm protein extracts and the detection of one of the vaccine candidates, Sm28GST, from a Venezuelan parasite isolate.
Transcriptome of the parasitic flatworm Schistosoma mansoni during intra-mammalian development.
Serine proteases of parasitic helminths.
Stage- and gender-specific proteomic analysis of Brugia malayi excretory-secretory products.
Blunting the knife: development of vaccines targeting digestive proteases of blood-feeding helminth parasites.
Schistosoma mansoni alter transcription of immunomodulatory gene products following in vivo praziquantel exposure.
Vector-based RNA interference of cathepsin B1 in Schistosoma mansoni.
Immune evasion genes from filarial nematodes.
Characterization and differential expression of cathepsin L3 alleles from Fasciola hepatica.
Proteinases and associated genes of parasitic helminths.
Proteomic analysis of Fasciola hepatica excretory and secretory products (FhESPs) involved in interacting with host PBMCs and cytokines by shotgun LC-MS/MS.
Infection by the Helminth Parasite Fasciola hepatica Requires Rapid Regulation of Metabolic, Virulence, and Invasive Factors to Adjust to Its Mammalian Host.
Cytokine induction and exploitation in schistosome infections.

Antibody trapping: A novel mechanism of parasite immune evasion by the trematode <i>Echinostoma caproni</i> .
The Two Faces of Nematode Infection: Virulence and Immunomodulatory Molecules From Nematode Parasites of Mammals, Insects and Plants.
Parasitic Nematode Immunomodulatory Strategies: Recent Advances and Perspectives.
Expression, purification and characterization of two leucine aminopeptidases of the blood fluke, <i>Schistosoma mansoni</i> .
REVIEW OF THE ROLE OF PARASITIC NEMATODE EXCRETORY/SECRETORY PROTEINS IN HOST IMMUNOMODULATION.
Developmental gene expression profiles of the human pathogen <i>Schistosoma japonicum</i> .
Proteases in blood-feeding nematodes and their potential as vaccine candidates.
Infection of mice lacking interleukin-7 (IL-7) reveals an unexpected role for IL-7 in the development of the parasite <i>Schistosoma mansoni</i> .
<i>Schistosoma mansoni</i> host-parasite relationship: interaction of contrapsin with adult worms.
Cathepsin F of <i>Teladorsagia circumcincta</i> is a recently evolved cysteine protease.
Culture for genetic manipulation of developmental stages of <i>Schistosoma mansoni</i> .
<i>Fasciola hepatica</i> cathepsin L-like proteases: biology, function, and potential in the development of first generation liver fluke vaccines.
Evasion of Host Immunity During <i>Fasciola hepatica</i> Infection.
Serpins in <i>Fasciola hepatica</i> : insights into host-parasite interactions.
Schistosome transcriptomes: new insights into the parasite and schistosomiasis.
Expression profile of the <i>Schistosoma japonicum</i> degradome reveals differential protease expression patterns and potential anti-schistosomal intervention targets.
Molecular characterization and functional analysis of the <i>Schistosoma mekongi</i> Ca(2+)-dependent cysteine protease (calpain).
Exposed proteins of the <i>Schistosoma japonicum</i> tegument.
Do antioxidants play a role in schistosome host-parasite interactions?
Exploiting Comparative Omics to Understand the Pathogenic and Virulence-Associated Protease: Anti-Protease Relationships in the Zoonotic Parasites <i>Fasciola hepatica</i> and <i>Fasciola gigantica</i> .
Glycans from <i>Fasciola hepatica</i> Modulate the Host Immune Response and TLR-Induced Maturation of Dendritic Cells.
Detection and partial characterization of glycosylphosphatidylinositol-specific phospholipase activities from <i>Fasciola hepatica</i> and <i>Schistosoma mansoni</i> .
Adult and juvenile <i>Fasciola</i> cathepsin L proteases: different enzymes for different roles.
<i>Fasciola hepatica</i> -Derived Molecules as Regulators of the Host Immune Response.
Behind Enemy Lines: Immunomodulatory Armamentarium of the Schistosome Parasite.
Roles of Cysteine Proteases in Biology and Pathogenesis of Parasites.
Molecular characterization of a novel cathepsin L from <i>Trichinella spiralis</i> and its participation in invasion, development and reproduction.
Adult schistosomes have an epithelial bacterial population distinct from the surrounding mammalian host blood.
A proteomic approach to the identification of tegumental proteins of male and female <i>Schistosoma bovis</i> worms.
<i>Schistosoma mansoni</i> Infection of Mice, Rats and Humans Elicits a Strong Antibody Response to a Limited Number of Reduction-Sensitive Epitopes on Five Major Tegumental Membrane Proteins.
Helminth Glycans at the Host-Parasite Interface and Their Potential for Developing Novel Therapeutics.

Synthetic and natural protease inhibitors provide insights into parasite development, virulence and pathogenesis.
Vaccination with antioxidant enzymes confers protective immunity against challenge infection with <i>Schistosoma mansoni</i> .
Eosinophil-mediated tissue inflammatory responses in helminth infection.
A distinctive repertoire of cathepsins is expressed by juvenile invasive <i>Fasciola hepatica</i> .
Excretory-secretory products of larval <i>Fasciola hepatica</i> investigated using a two-dimensional proteomic approach.
[Research progress on cystatin of parasitic nematodes].
A Complex Proteomic Response of the Parasitic Nematode <i>Anisakis simplex</i> s.s. to <i>Escherichia coli</i> Lipopolysaccharide.
Stage-specifically expressed schistosome proteins as potential chemotherapeutic targets.
Helminth genomics: The implications for human health.
Characterization and expression of a novel cystatin gene from <i>Schistosoma japonicum</i> .
Schistosome proteomics: updates and clinical implications.
A preliminary study to understand the effect of <i>Fasciola hepatica</i> tegument on naïve macrophages and humoral responses in an ovine model.
A cercarial invadolysin interferes with the host immune response and facilitates infection establishment of <i>Schistosoma mansoni</i> .
Potential roles of <i>Toxocara canis</i> larval excretory secretory molecules in immunomodulation and immune evasion.
Discovery of Potent Inhibitors of <i>Schistosoma mansoni</i> NAD <sup>+</sup> Catabolizing Enzyme.
Characterization of <i>Haemonchus contortus</i> Excretory/Secretory Antigen (ES-15) and Its Modulatory Functions on Goat Immune Cells In Vitro.
Study of the cross-talk between <i>Fasciola hepatica</i> juveniles and the intestinal epithelial cells of the host by transcriptomics in an in vitro model.
Chapter 4. Peptidases of trematodes.
How schistosomes alter the human serum proteome.
Immune Evasion Strategies of Schistosomes.
<i>Schistosoma mansoni</i> host-exposed surface antigens characterized by sera and recombinant antibodies from schistosomiasis-resistant rats.
Proteomic analysis of secretory products from the model gastrointestinal nematode <i>Heligmosomoides polygyrus</i> reveals dominance of venom allergen-like (VAL) proteins.
Insights into the biological features of the antigenic determinants recognized by four monoclonal antibodies in redia and adult stages of the liver fluke <i>Fasciola hepatica</i> .
Excretory-secretory proteome of larval <i>Schistosoma mansoni</i> and <i>Echinostoma caproni</i> , two parasites of <i>Biomphalaria glabrata</i> .
Immunofluorescent Localization of Proteins in <i>Schistosoma mansoni</i> .
Secreted cathepsin L-like peptidases are involved in the degradation of trapped antibodies on the surface of <i>Echinostoma caproni</i> .
<i>Fasciola hepatica</i> fatty acid binding protein induces the alternative activation of human macrophages.
Functional Diversity of the Excretory/Secretory Proteins of Nematode Parasites.
Immunology of human helminth infection.
Developmentally regulated secretion of cathepsin L-like cysteine proteases by <i>Haemonchus contortus</i> .
<i>Fasciola hepatica</i> , TGF- $\beta$ and host mimicry: the enemy within.
The ultrastructural architecture of the adult <i>Schistosoma japonicum</i> tegument.

MicroRNAs in Helminth Parasites: A Systematic Review.
Parasitic Helminth-Derived microRNAs and Extracellular Vesicle Cargos as Biomarkers for Helminthic Infections.
Structural and functional analysis of cathepsin S of Heterodera spp: a promising candidate for its control.
Analysis of Schistosoma mansoni Extracellular Vesicles Surface Glycans Reveals Potential Immune Evasion Mechanism and New Insights on Their Origins of Biogenesis.
Functional characterization of single-domain cystatin-like cysteine proteinase inhibitors expressed by the trematode Fasciola hepatica.
Cattle co-infection of Echinococcus granulosus and Fasciola hepatica results in a different systemic cytokine profile than single parasite infection.
Cysteine proteases as potential antigens in antiparasitic DNA vaccines.
[Significance of schistosome lectins in the host allergic reaction evoked by penetration of cercaria].
Predictions of novel Schistosoma mansoni - human protein interactions consistent with experimental data.
Excretory/secretory products of anisakid nematodes: biological and pathological roles.
On the three-finger protein domain fold and CD59-like proteins in Schistosoma mansoni.
A quantitative proteomic analysis of the tegumental proteins from Schistosoma mansoni schistosomula reveals novel potential therapeutic targets.
Making sense of the schistosome surface.
Protective role of purified cysteine proteinases against Fasciola gigantica infection in experimental animals.
Recombinant micro-exon gene 3 (MEG-3) antigens from Schistosoma mansoni failed to induce protection against infection but show potential for serological diagnosis.
Resolving the origins of secretory products and anthelmintic responses in a human parasitic nematode at single-cell resolution.
Glycan gimmickry by parasitic helminths: a strategy for modulating the host immune response?
Cloning, expression and characterisation of a cysteine protease from Trichinella spiralis.
Production and processing of a recombinant Fasciola hepatica cathepsin B-like enzyme (FhcatB1) reveals potential processing mechanisms in the parasite.
Role of the bovine immune system and genome in resistance to gastrointestinal nematodes.
Early differential gene expression in haemocytes from resistant and susceptible Biomphalaria glabrata strains in response to Schistosoma mansoni.
Identification of host immune regulation candidate genes of Toxascaris leonina by expression sequenced tags (ESTs) analysis.
Cercarial transformation and in vitro cultivation of Schistosoma mansoni schistosomules.
Molecular characterization of Schistosoma japonicum tegument protein tetraspanin-2: sequence variation and possible implications for immune evasion.
Juvenile-specific cathepsin proteases in Fasciola spp.: their characteristics and vaccine efficacies.
Fasciola hepatica mucin-encoding gene: expression, variability and its potential relevance in host-parasite relationship.
Revisiting the Schistosoma japonicum life cycle transcriptome for new insights into lung schistosomula development.
Filarial nematode secreted product ES-62 is an anti-inflammatory agent: therapeutic potential of small molecule derivatives and ES-62 peptide mimetics.
Identification of a developmentally regulated Schistosoma mansoni serine protease homologous to mouse plasma kallikrein and human factor I.



Molecular cloning and characterization of serine protease inhibitor from food-borne nematode, <i>Gnathostoma spinigerum</i> .
Proteomic analysis of the schistosome tegument and its surface membranes.
Functional analysis of the cathepsin-like cysteine protease genes in adult <i>Brugia malayi</i> using RNA interference.
Cathepsins B1 and B2 of <i>Trichobilharzia</i> spp., bird schistosomes causing cercarial dermatitis.
Genomics of parasitic flatworms.
Examining the role of macrolides and host immunity in combatting filarial parasites.
Molecular cloning, characterization and functional analysis of a novel juvenile-specific cathepsin L of <i>Fasciola gigantica</i> .
Molecular characterization of a novel GSTO2 of <i>Fasciola hepatica</i> and its roles in modulating murine macrophages.
Novel mechanisms of immune evasion by <i>Schistosoma mansoni</i> .
Helminth C-type lectins and host-parasite interactions.
Isolation of cDNAs encoding secreted and transmembrane proteins from <i>Schistosoma mansoni</i> by a signal sequence trap method.
Functions of the tegument of schistosomes: clues from the proteome and lipidome.
Insect immune responses to nematode parasites.
Sugar Coating: Utilisation of Host Serum Sialoglycoproteins by <i>Schistosoma mansoni</i> as a Potential Immune Evasion Mechanism.
Mapping the binding between the tetraspanin molecule (Sj23) of <i>Schistosoma japonicum</i> and human non-immune IgG.
The tegument surface membranes of the human blood parasite <i>Schistosoma mansoni</i> : a proteomic analysis after differential extraction.
Transcriptomic Analysis of Ovine Hepatic Lymph Node Following <i>Fasciola hepatica</i> Infection - Inhibition of NK Cell and IgE-Mediated Signaling.
Molecular Characteristics and Potent Immunomodulatory Activity of <i>Fasciola hepatica</i> Cystatin.
Unisexual infection with <i>Schistosoma mansoni</i> in mice has the potential to boost the immune response against eggs after challenge infection.
Functional characterization of SjB10, an intracellular serpin from <i>Schistosoma japonicum</i> .
Transcriptome analysis of the acoelomate human parasite <i>Schistosoma mansoni</i> .
Nematode microRNAs can Individually Regulate Interferon Regulatory Factor 4 and mTOR in Differentiating T Helper 2 Lymphocytes and Modulate Cytokine Production in Macrophages.
Partial purification and characterization of a cysteine protease inhibitor from the plerocercoid of <i>Spirometra erinacei</i> .
Time series analysis of the transcriptional responses of <i>Biomphalaria glabrata</i> throughout the course of intramolluscan development of <i>Schistosoma mansoni</i> and <i>Echinostoma paraensei</i> .
The regulatory roles of <i>Fasciola hepatica</i> GSTO1 protein in inflammatory cytokine expression and apoptosis in murine macrophages.
<i>Fasciola hepatica</i> : rapid switching of stage-specific antigen expression after infection.
Identification of abundantly expressed novel and conserved genes from the infective larval stage of <i>Toxocara canis</i> by an expressed sequence tag strategy.
Effects of an <i>Onchocerca</i> -derived cysteine protease inhibitor on microfilariae in their simuliid vector.
Evolution of tetraspanin antigens in the zoonotic Asian blood fluke <i>Schistosoma japonicum</i> .
In vitro lectin binding to the outer surface of <i>Spirocerca lupi</i> at different life-stages.
Immunomodulation and Immune Escape Strategies of Gastrointestinal Helminths and Schistosomes.

Characterization of <i>Schistosoma japonicum</i> tetraspanning orphan receptor and its role in binding to complement C2 and immunoprotection against murine schistosomiasis.
Characterization and expression of cathepsin B2 in <i>Fasciola gigantica</i> .
Secreted proteomes of different developmental stages of the gastrointestinal nematode <i>Nippostrongylus brasiliensis</i> .
<i>Haemonchus contortus</i> P-glycoproteins interact with host eosinophil granules: a novel insight into the role of ABC transporters in host-parasite interaction.
Down-regulated lymphoproliferation coincides with parasite maturation and with the collapse of both gamma interferon and interleukin-4 responses in a bovine model of onchocerciasis.
Cysteine protease secreted by <i>Paragonimus westermani</i> attenuates effector functions of human eosinophils stimulated with immunoglobulin G.
Transcriptomics tool for the human <i>Schistosoma</i> blood flukes using microarray gene expression profiling.
Comparative proteome analysis of the tegument of male and female adult <i>Schistosoma mansoni</i> .
Human serum activates the tegument of female schistosomes and supports recovery from Praziquantel.
Myeloid-Derived Suppressor Cells: The Expanding World of Helminth Modulation of the Immune System.
Identification and expression of a transforming growth factor beta (TGF- $\beta$ ) homologue in the tropical liver fluke <i>Fasciola gigantica</i> .
Generation of a single-chain variable fragment phage display antibody library from naïve mice panned against <i>Fasciola hepatica</i> antigens.
Identifying novel candidates and configurations for human helminth vaccines.
Transcriptomes and pathways associated with infectivity, survival and immunogenicity in <i>Brugia malayi</i> L3.
Cysteine proteases secreted by the pinewood nematode, <i>Bursaphelenchus xylophilus</i> : In silico analysis.
Proteomic analysis of <i>Schistosoma mansoni</i> cercarial secretions.
Data-mining of the <i>Meloidogyne incognita</i> degradome and comparative analysis of proteases in nematodes.
Generating a core cluster of <i>Fasciola hepatica</i> virulence and immunomodulation-related genes using a comparative in silico approach.
Proteomic analysis of the <i>Echinococcus granulosus</i> metacestode during infection of its intermediate host.
Fibrinogen and fibrin are novel substrates for <i>Fasciola hepatica</i> cathepsin L peptidases.
[STRUCTURAL, MOLECULAR AND FUNCTIONAL FEATURES OF NEMATODE SURFACES AND THE POSSIBILITY OF ANTHELMINTIC DRUGS DEVELOPMENT (REVIEW)].
Let's talk about sexes: sex-related N-glycosylation in ecologically important invertebrates.
Molecular cloning and characterization of a cathepsin L-like cysteine protease of <i>Angiostrongylus cantonensis</i> .
Molecular characterization and determination of the biochemical properties of cathepsin L of <i>Trichinella spiralis</i> .
Cloning and Characterization of Two Potent Kunitz Type Protease Inhibitors from <i>Echinococcus granulosus</i> .
Nippocystatin, a cysteine protease inhibitor from <i>Nippostrongylus brasiliensis</i> , inhibits antigen processing and modulates antigen-specific immune response.

Fasciola hepatica - the pilot study of in vitro assessing immune response against native and recombinant antigens of the fluke.
Inhibition of carbonic anhydrase using aspirin is a novel method to block schistosomiasis infection of the parasitic trematode, Schistosoma mansoni, in the intermediate snail host, Biomphalaria glabrata.
Toxocara canis: molecular basis of immune recognition and evasion.
Commensal-pathogen interactions in the intestinal tract: lactobacilli promote infection with, and are promoted by, helminth parasites.
Immunosuppressive PAS-1 is an excretory/secretory protein released by larval and adult worms of the ascarid nematode Ascaris suum.
The esophageal gland mediates host immune evasion by the human parasite Schistosoma mansoni.
Secretion of RNA-Containing Extracellular Vesicles by the Porcine Whipworm, Trichuris suis.
Secretion of an anti-inflammatory, immunomodulatory factor by Schistosomulae of Schistosoma mansoni.
Enteral vaccination of rats against Fasciola hepatica using recombinant cysteine proteinase (cathepsin L1).
Targeting Cysteine Proteases and their Inhibitors to Combat Trypanosomiasis.
A single amino acid substitution affects substrate specificity in cysteine proteinases from Fasciola hepatica.
Proteases as Therapeutic Targets Against the Parasitic Cnidarian Ceratomyxa shasta: Characterization of Molecules Key to Parasite Virulence In Salmonid Hosts.
Angiostrongylus cantonensis cathepsin B-like protease (Ac-cathB-1) is involved in host gut penetration.
Trichinella-induced immunomodulation: Another tale of helminth success.
Immunological genomics of Brugia malayi: filarial genes implicated in immune evasion and protective immunity.
Biochemical and biophysical methodologies open the road for effective schistosomiasis therapy and vaccination.
Expression and characterization of glutathione peroxidase activity in the human blood fluke Schistosoma mansoni.
Developmentally regulated expression of surface carbohydrate residues on larval stages of the avian schistosome Trichostrongylus axei.
Revealing the inhibitory potential of Yersinia enterocolitica on cysteine proteases of the papain family.
A novel C-type lectin secreted by a tissue-dwelling parasitic nematode.
Cysteine protease activities during maturation stages of Paragonimus westermani.
Cystic echinococcosis: aspects of immune response, immunopathogenesis and immune evasion from the human host.
Evidence of opiates and opioid neuropeptides and their immune effects in parasitic invertebrates representing three different phyla: Schistosoma mansoni, Theromyzon tessulatum, Trichinella spiralis.
Liver cystic echinococcosis and human host immune and autoimmune follow-up: A review.
Schistosome vaccines: problems, pitfalls and prospects.
Schistosoma mansoni Tegument (Smteg) Induces IL-10 and Modulates Experimental Airway Inflammation.
Proteinases released in vitro by the parasitic stages of the bovine abomasal nematode Ostertagia ostertagi.
ES proteins analysis of Angiostrongylus cantonensis: products of the potential parasitism genes?
Proteins exposed at the adult schistosome surface revealed by biotinylation.

Pivotal roles of the parasite PGD2 synthase and of the host D prostanoid receptor 1 in schistosome immune evasion.
Impaired resistance in early secondary <i>Nippostrongylus brasiliensis</i> infections in mice with defective eosinophilopoiesis.
Characterization of excretory/secretory endopeptidase and metallo-aminopeptidases from <i>Taenia crassiceps</i> metacestodes.
Cleavage of immunoglobulin G by excretory-secretory cathepsin S-like protease of <i>Spirometra mansoni</i> plerocercoid.
Protein variation in blood-dwelling schistosome worms generated by differential splicing of micro-exon gene transcripts.
Purification of a chymotrypsin-like enzyme present on adult <i>Schistosoma mansoni</i> worms from infected mice and its characterization as a host carboxylesterase.
[Cathepsin L cysteine protease from <i>Taenia solium</i> : its biological role in the infection and potential use for the immunodiagnosis of neurocysticercosis].
The Potential Role of Schistosome-Associated Factors as Therapeutic Modulators of the Immune System.
Planarians to schistosomes: an overview of flatworm cell-types and regulators.
Emerging roles for extracellular vesicles in <i>Schistosoma</i> infection.
A single exon-encoded <i>Theileria parva</i> strain Muguga cysteine protease (ThpCP): Molecular modelling and characterisation.
Bovine Natural Antibody Relationships to Specific Antibodies and <i>Fasciola hepatica</i> Burdens after Experimental Infection and Vaccination with Glutathione S-Transferase.
Studies on phospholipid turnover argue against sloughing of tegumental membranes in adult <i>Schistosoma mansoni</i> .
Identification and characterization of a cathepsin L-like cysteine protease from <i>Rhipicephalus (Boophilus) annulatus</i> .
Non-immune immunoglobulins shield <i>Schistosoma japonicum</i> from host immunorecognition.
Multi-Omic Profiling, Structural Characterization, and Potent Inhibitor Screening of Evasion-Related Proteins of a Parasitic Nematode, <i>Haemonchus contortus</i> , Surviving Vaccine Treatment.
IL-12 and TNF-alpha production by dendritic cells stimulated with <i>Schistosoma mansoni</i> schistosomula tegument is TLR4- and MyD88-dependent.
Transforming growth factor-beta and Th17 responses in resistance to primary murine schistosomiasis <i>mansoni</i> .
<i>Rhipicephalus microplus</i> and <i>Ixodes ovatus</i> cystatins in tick blood digestion and evasion of host immune response.
Serine protease activity in excretory-secretory products of <i>Oestrus ovis</i> (Diptera: Oestridae) larvae.
Schistosomes in the Lung: Immunobiology and Opportunity.
Effects of gastrointestinal nematode infection on the ruminant immune system.
Identification of Two Meloidogyne hapla Genes and an Investigation of Their Roles in the Plant-Nematode Interaction.
<i>Fasciola gigantica</i> Cathepsin L1H: High Sensitivity and Specificity of Immunochromatographic Strip Test for Antibody Detection.
Effect of cysteine peptidase inhibitor of <i>Eudiplozoon nipponicum</i> (Monogenea) on cytokine expression of macrophages in vitro.
<i>Fasciola gigantica</i> tegumental calcium-binding EF-hand protein 4 exerts immunomodulatory effects on goat monocytes.

Cathepsin L proteinase secreted by <i>Fasciola hepatica</i> in vitro prevents antibody-mediated eosinophil attachment to newly excysted juveniles.
Looking beyond the induction of Th2 responses to explain immunomodulation by helminths.
<i>Fasciola hepatica</i> : parasite-secreted proteinases degrade all human IgG subclasses: determination of the specific cleavage sites and identification of the immunoglobulin fragments produced.
Neglected Agent Eminent Disease: Linking Human Helminthic Infection, Inflammation, and Malignancy.
A family of secreted mucins from the parasitic nematode <i>Toxocara canis</i> bears diverse mucin domains but shares similar flanking six-cysteine repeat motifs.
<i>Toxocara canis</i> larval excretory/secretory proteins impair eosinophil-dependent resistance of mice to <i>Nippostrongylus brasiliensis</i> .
Liver pathology and immune response in experimental <i>Fasciola hepatica</i> infections of goats.
Detection of cysteine protease in <i>Taenia solium</i> -induced brain granulomas in naturally infected pigs.
Immuno-evasive tactics by schistosomes identify an effective allergy preventative.
Regulation of immunity by Taeniids: lessons from animal models and in vitro studies.
Unveiling the immunomodulatory properties of <i>Haemonchus contortus</i> adhesion regulating molecule 1 interacting with goat T cells.
Tissue localization of collagenase and leucine aminopeptidase in the bovine filarial parasite <i>Setaria cervi</i> .
[Proteomic analysis of the excretion-secretion products of four <i>Trichinella spiralis</i> isolates obtained from accidental hosts].
Sensitization with anti-inflammatory BmAFI of <i>Brugia malayi</i> allows L3 development in the hostile peritoneal cavity of <i>Mastomys coucha</i> .
Novel cathepsin B and cathepsin B-like cysteine protease of <i>Naegleria fowleri</i> excretory-secretory proteins and their biochemical properties.
Therapeutic potential of helminths in autoimmune diseases: helminth-derived immune-regulators and immune balance.
Excretory/secretory proteins inhibit host immune responses by downregulating the TLR4/NF- $\kappa$ B/MAPKs signaling pathway: A possible mechanism of immune evasion in parasitic nematode <i>Haemonchus contortus</i> .
Functionally Expression of Metalloproteinase in <i>Taenia solium</i> Metacestode and Its Evaluation for Serodiagnosis of Cysticercosis.
The Role of the Intestinal Epithelium in the "Weep and Sweep" Response during Gastro-Intestinal Helminth Infections.
Degradation of human matrix metalloproteinase-9 by secretory metalloproteinases of <i>Angiostrongylus cantonensis</i> infective stage.
Sm-p80-based DNA vaccine provides baboons with levels of protection against <i>Schistosoma mansoni</i> infection comparable to those achieved by the irradiated cercarial vaccine.
Echinococcosis.
The genome of the hydatid tapeworm <i>Echinococcus granulosus</i> .
Transcriptome of <i>Sphaerospora molnari</i> (Cnidaria, Myxosporidia) blood stages provides proteolytic arsenal as potential therapeutic targets against sphaerosporosis in common carp.
<i>Eudiplozoon nipponicum</i> (Monogenea, Diplozoidae) and its adaptation to haematophagy as revealed by transcriptome and secretome profiling.
Crystal structure of <i>Schistosoma mansoni</i> arginase, a potential drug target for the treatment of schistosomiasis.
<i>Schistosoma mansoni</i> : the developmental regulation and immunolocalization of antioxidant enzymes.

Extracellular Vesicles Derived From <i>Trichinella spiralis</i> Muscle Larvae Ameliorate TNBS-Induced Colitis in Mice.
Characterization of Divalent Metal Transporter 1 (DMT1) in <i>Brugia malayi</i> suggests an intestinal-associated pathway for iron absorption.
TLRs, Treg, and B Cells, an Interplay of Regulation during Helminth Infection.
In vivo studies of the early, peritoneal, cellular and free radical response in rats infected with <i>Fasciola hepatica</i> by flow cytometric analysis.
Cross-reactivity of <i>Schistosoma mansoni</i> cytosolic superoxide dismutase, a protective vaccine candidate, with host superoxide dismutase and identification of parasite-specific B epitopes.
Proteins as Targets in Anti-Schistosomal Drug Discovery and Vaccine Development.
Molecular characterization and functional analysis of a glutathione peroxidase gene from <i>Aphelenchoides besseyi</i> (Nematoda: Aphelenchoididae).
Molecular cloning and expression of the cDNA sequence encoding a novel aspartic protease from <i>Uncinaria stenocephala</i> .
Virulence factors of schistosomes.
Bm-CPI-2, a cystatin from <i>Brugia malayi</i> nematode parasites, differs from <i>Caenorhabditis elegans</i> cystatins in a specific site mediating inhibition of the antigen-processing enzyme AEP.
Induction of protection in murine experimental models against <i>Trichinella spiralis</i> : an up-to-date review.
Prime-boost and recombinant protein vaccination strategies using Sm-p80 protects against <i>Schistosoma mansoni</i> infection in the mouse model to levels previously attainable only by the irradiated cercarial vaccine.
Recombinant dynein light intermediate chain of <i>Haemonchus contortus</i> affects the functions of goat immune cells in vitro.
SjCa8, a calcium-binding protein from <i>Schistosoma japonicum</i> , inhibits cell migration and suppresses nitric oxide release of RAW264.7 macrophages.
Murine nematode immunology in Australasia.
RNAi-mediated gene silencing to assess the role of synaptobrevin and cystatin in tick blood feeding.
An abundantly expressed mucin-like protein from <i>Toxocara canis</i> infective larvae: the precursor of the larval surface coat glycoproteins.
Differential release and phagocytosis of tegument glycoconjugates in neurocysticercosis: implications for immune evasion strategies.
<i>Fasciola hepatica</i> induces weak NETosis and low production of intra- and extracellular ROS in exposed bovine polymorphonuclear neutrophils.
Improving animal and human health through understanding liver fluke immunology.
Bacterial feeding nematodes ingest haemocytes in the haemocoel of the insect <i>Galleria mellonella</i> .
Local immune response to experimental <i>Fasciola hepatica</i> infection in sheep.
A phosphorylcholine-containing filarial nematode-secreted product disrupts B lymphocyte activation by targeting key proliferative signaling pathways.
Involvement of IL-10 and IL-4 in evasion strategies of <i>Echinococcus granulosus</i> to host immune response.
Proteomic analysis revealed T cell hyporesponsiveness induced by <i>Haemonchus contortus</i> excretory and secretory proteins.
[Expression of the cathepsin L1 gene of <i>Fasciola hepatica</i> eucaryotic cells].
Insect Immune Evasion by Dauer and Nondauer Entomopathogenic Nematodes.
A Novel $\alpha/\beta$ Hydrolase Domain Protein Derived From <i>Haemonchus contortus</i> Acts at the Parasite-Host Interface.

On the presence and immunoregulatory functions of extracellular microRNAs in the trematode <i>Fasciola hepatica</i> .
[Role of macrophages in schistosome infection].
<i>Fasciola hepatica</i> and <i>Fasciola gigantica</i> : comparison of cellular response to experimental infection in sheep.
Immune reactivity and host modulatory roles of two novel <i>Haemonchus contortus</i> cathepsin B-like proteases.
Ivermectin disrupts the function of the excretory-secretory apparatus in microfilariae of <i>Brugia malayi</i> .
Proteomic analysis of exosome-like vesicles from <i>Fasciola gigantica</i> adult worm provides support for new vaccine targets against fascioliasis.
Immune Response to <i>Brugia malayi</i> Asparaginyl-tRNA Synthetase in Balb/c Mice and Human Clinical Samples of Lymphatic Filariasis.
Identification of proteins from the secretory/excretory products (SEPs) of the branchiuran ectoparasite <i>Argulus foliaceus</i> (Linnaeus, 1758) reveals unique secreted proteins amongst haematophagous ecdysozoa.
Eliminating Schistosomes through Vaccination: What are the Best Immune Weapons?
Serine protease activity in developmental stages of <i>Eimeria tenella</i> .
Polymorphism and natural selection of antigen B1 of <i>Echinococcus granulosus</i> isolated from different host assemblages in India.
Screening of an <i>Echinococcus granulosus</i> cDNA library with IgG4 from patients with cystic echinococcosis identifies a new tegumental protein involved in the immune escape.
Soil-transmitted helminthic vaccines: Where are we now?
Hyporesponsiveness of murine B lymphocytes exposed to the filarial nematode secreted product ES-62 in vivo.
Characterization of a glycine-rich protein from <i>Rhipicephalus microplus</i> : tissue expression, gene silencing and immune recognition.
Homologues of human macrophage migration inhibitory factor from a parasitic nematode. Gene cloning, protein activity, and crystal structure.
Moderate protection is induced by a chimeric protein composed of leucine aminopeptidase and cathepsin L1 against <i>Fasciola hepatica</i> challenge in sheep.
[STRUCTURAL, MOLECULAR AND FUNCTIONAL FEATURES OF THE TREMATODE TEGUMENT AND OF ANTHELMINTIC DRUGS DEVELOPMENT (REVIEW)].
Cleavage of human IgE mediated by <i>Schistosoma mansoni</i> .
Adaptation of the secretome of <i>Echinostoma caproni</i> may contribute to parasite survival in a Th1 milieu.
Cercaria-schistosomulum surface transformation of <i>Trichobilharzia szidati</i> and its putative immunological impact.
Proteomic identification of endothelial cell surface proteins isolated from the hepatic portal vein of mice infected with <i>Schistosoma bovis</i> .
Insect Immunity to Entomopathogenic Nematodes and Their Mutualistic Bacteria.
Molecular changes in <i>Opisthorchis viverrini</i> (Southeast Asian liver fluke) during the transition from the juvenile to the adult stage.
Time-resolved transcriptional profiling of <i>Trichinella</i> -infected murine myocytes helps to elucidate host-pathogen interactions in the muscle stage.
Characterization of <i>Haemonchus contortus</i> calreticulin suggests its role in feeding and immune evasion by the parasite.

ICAM-1 and iNOS expression increased in the skin of mice after vaccination with gamma-irradiated cercariae of <i>Schistosoma mansoni</i> .
Intrapopulation dynamics of a wild strain of <i>Taenia crassiceps</i> (WFU) (Cestoda: Taeniidae) in BALB/cJ mice.
Macrophage migration inhibitory factor (mif) transcription is significantly elevated in <i>Caenorhabditis elegans</i> dauer larvae.
Ticks, Ixodes scapularis, Feed Repeatedly on White-Footed Mice despite Strong Inflammatory Response: An Expanding Paradigm for Understanding Tick-Host Interactions.
Proteomic analysis of soluble protein extract of adult <i>Toxocara cati</i> .
A new bacteria-free strategy induced by MaGal2 facilitates pinewood nematode escape immune response from its vector beetle.
<i>Trichinella spiralis</i> paramyosin binds to C8 and C9 and protects the tissue-dwelling nematode from being attacked by host complement.
<i>Trichuris suis</i> secrete products that reduce disease severity in a multiple sclerosis model.
Genetic diversity among natural populations of <i>Schistosoma haematobium</i> might contribute to inconsistent virulence and diverse clinical outcomes.
Enhancement of Sm-p80 (large subunit of calpain) induced protective immunity against <i>Schistosoma mansoni</i> through co-delivery of interleukin-2 and interleukin-12 in a DNA vaccine formulation.
[Larval stages of <i>Ascaris lumbricoides</i> : hyaluronan-binding capacity].
Interaction between the blood fluke, <i>Sanguinicola inermis</i> and humoral components of the immune response of carp, <i>Cyprinus carpio</i> .
Characterization of the immune response to DNA vaccination strategies for schistosomiasis candidate antigen, Sm-p80 in the baboon.
MicroRNA expression profile in the third- and fourth-stage larvae of <i>Angiostrongylus cantonensis</i> .
Novel host immune evasion strategy of the endoparasitoid <i>Drino inconspicuoides</i> .
Having a pair: the key to immune evasion for the diploid pathogen <i>Schistosoma japonicum</i> .
Induction of protective immunity against <i>Schistosoma mansoni</i> via DNA priming and boosting with the large subunit of calpain (Sm-p80): adjuvant effects of granulocyte-macrophage colony-stimulating factor and interleukin-4.
Cytokine expression in hamsters experimentally infected with <i>Opisthorchis viverrini</i> .
<i>Aphidius ervi</i> teratocytes release an extracellular enolase.
Mucins in the host defence against <i>Naegleria fowleri</i> and mucinolytic activity as a possible means of evasion.
Functional analysis of a highly conserved abundant larval transcript-2 (alt-2) intron 2 repeat region of lymphatic filarial parasites.
Cloning and partial characterization of a <i>Boophilus microplus</i> (Acari: Ixodidae) calreticulin.
Identification and characterisation of a novel repetitive antigen from <i>Onchocerca</i> spp.
A filarial nematode-secreted phosphorylcholine-containing glycoprotein uncouples the B cell antigen receptor from extracellular signal-regulated kinase-mitogen-activated protein kinase by promoting the surface Ig-mediated recruitment of Src homology 2 domain-containing tyrosine phosphatase-1 and Pac-1 mitogen-activated kinase-phosphatase.
A shift to Th2 immune response caused by constitutive expression of IPSE/alpha-1 in transfected pig fibroblasts in mice.
A Nematode Calreticulin, Rs-CRT, Is a Key Effector in Reproduction and Pathogenicity of <i>Radopholus similis</i> .
Penetration of the mosquito midgut is not required for <i>Brugia pahangi</i> microfilariae to avoid the melanotic encapsulation response of <i>Armigeres subalbatus</i> .



Tick saliva inhibits dendritic cell migration, maturation, and function while promoting development of Th2 responses.
Hepatic transcriptome study of <i>Taenia asiatica</i> infection in suckling pigs.
Effect of CD4+ CD25+ regulatory T cells on the immune evasion of <i>Schistosoma japonicum</i> .
Humoral and T cell-mediated immune response against trichomoniasis.
Production and characterization of monoclonal antibodies against cathepsin B and cathepsin B-Like proteins of <i>Naegleria fowleri</i> .
Tick saliva induces regulatory dendritic cells: MAP-kinases and Toll-like receptor-2 expression as potential targets.
Molecular and Functional Characterization of Inhibitor of Apoptosis Proteins (IAP, BIRP) in <i>Echinococcus granulosus</i> .
Proteomic mapping of the lung vascular endothelial cell surface in <i>Schistosoma bovis</i> -infected hamsters.
Recombinant Sj16 from <i>Schistosoma japonicum</i> contains a functional N-terminal nuclear localization signal necessary for nuclear translocation in dendritic cells and interleukin-10 production.
Concepts in immunology and diagnosis of hydatid disease.
Evidence for serine protease inhibitor activity in the ovarian calyx fluid of the endoparasitoid <i>Venturia canescens</i> .
Immunology of <i>Taenia solium</i> taeniasis and human cysticercosis.
Infective Larvae of <i>Brugia malayi</i> Induce Polarization of Host Macrophages that Helps in Immune Evasion.
<i>Echinococcus multilocularis</i> Calreticulin Interferes with C1q-Mediated Complement Activation.
Antibody response to glycan antigens of hydatid cyst fluid, laminated layer and protoscolex of <i>Echinococcus granulosus</i> .
Hemomucin, an O-glycosylated protein on embryos of the wasp <i>Macrocentrus cingulum</i> that protects it against encapsulation by hemocytes of the host <i>Ostrinia furnacalis</i> .
Whole blood transcriptome analysis in onchocerciasis.
Immunome and immune complex-forming components of <i>Brugia malayi</i> identified by microfilaremic human sera.
Effects of protoscoleces and AgB from <i>Echinococcus granulosus</i> on human neutrophils: possible implications on the parasite's immune evasion mechanisms.
A 53 KDa Glycan Antigen of Hydatid Cyst Wall May Involve in Evasion from Host Immune System.
Differential Evolutionary Selection and Natural Evolvability Observed in ALT Proteins of Human Filarial Parasites.
Endogenous morphine and parasitic helminthes.
Immunomodulation of ALT-2 and TLR may collude in antigen specific T cell hyporesponsiveness: proposed mechanism for elevated IL-10 levels in Balb/C mice.
In vitro effects of prostaglandin E2 on leucocytes from sticklebacks ( <i>Gasterosteus aculeatus</i> ) infected and not infected with the cestode <i>Schistocephalus solidus</i> .
Hemiurid trematode sporocysts are undetected by hemocytes of their intermediate host, the ark cockle <i>Anadara trapezia</i> : potential role of surface carbohydrates in successful parasitism.
Predictive Global Models of Cruzain Inhibitors with Large Chemical Coverage.
The experimental infection of pigs with different numbers of <i>Taenia solium</i> eggs: immune response and efficiency of establishment.
Efficient use of a small genome to generate antigenic diversity in tick-borne ehrlichial pathogens.
The vagina has reducing environment sufficient for activation of <i>Trichomonas vaginalis</i> cysteine proteinases.

Structural characterization of anti-CCL5 activity of the tick salivary protein evasin-4.
Mutation scan screening of <i>Echinococcus granulosus</i> isolates of Indian origin.
A trypsin inhibitor-like protein secreted by <i>Cotesia vestalis</i> teratocytes inhibits hemolymph prophenoloxidase activation of <i>Plutella xylostella</i> .
Molecular epidemiology of bovine anaplasmosis with a particular focus in Mexico.
Identification of critical residues of an immunodominant region of <i>Echinococcus granulosus</i> antigen B.
Integrative approach reveals composition of endoparasitoid wasp venoms.
Differences in the proliferation trend of ' <i>Microsporidium</i> ' sp. PL03 in <i>Culex pipiens</i> and <i>C. torrentium</i> larvae.
<i>Echinococcus multilocularis</i> metacystodes modulate cellular cytokine and chemokine release by peripheral blood mononuclear cells in alveolar echinococcosis patients.
Genome-wide prediction of the polymorphic Ser gene family in <i>Tetrahymena thermophila</i> based on motif analysis.
A possible relationship between Thromboxane B2 and Leukotriene B4 and the encapsulation of <i>Dirofilaria repens</i> worms in human subcutaneous dirofilariasis.
Human dendritic cell sequestration onto the <i>Necator americanus</i> larval sheath during ex-sheathing: a possible mechanism for immune privilege.
Cloning and expression of apyrase gene from <i>Ancylostoma caninum</i> in <i>Escherichia coli</i> .
An Ovarian Protein Involved in Passive Avoidance of an Endoparasitoid To Evade Its Host Immune Response.
In vitro effects of the neuroactive substances serotonin and $\gamma$ -aminobutyric acid on leucocytes from sticklebacks ( <i>Gasterosteus aculeatus</i> ).
The tick salivary protein sialostatin L inhibits the Th9-derived production of the asthma-promoting cytokine IL-9 and is effective in the prevention of experimental asthma.
Proteome analysis of excretory-secretory proteins of <i>Entamoeba histolytica</i> HM1:IMSS via LC-ESI-MS/MS and LC-MALDI-TOF/TOF.
Lack of an association between hydatid disease and autoimmunity.
Sequence analysis and expression of a virus-like particle protein, VLP2, from the parasitic wasp <i>Venturia canescens</i> .
Autoimmunity in Human CE: Correlative with The Fertility Status of The CE Cyst.
Tick saliva protein Evasin-3 modulates chemotaxis by disrupting CXCL8 interactions with glycosaminoglycans and CXCR2.
Scuticociliate proteinases may modulate turbot immune response by inducing apoptosis in pronephric leucocytes.
<i>Echinococcus granulosus</i> cyst fluid inhibits inflammatory responses through inducing histone demethylase KDM5B in macrophages.
Dynamic changes of DC and T cell subsets in mice during <i>Echinococcus multilocularis</i> infection.
Investigation of the transcriptomic response in Atlantic salmon ( <i>Salmo salar</i> ) gill exposed to <i>Paramoeba perurans</i> during early onset of disease.
An insight into the sialotranscriptome and virome of Amazonian anophelines.
Cloning, expression, purification, and kinetic characterization of mitochondrial thioredoxin (TsTrx2), cytosolic thioredoxin (TsTrx1), and glutaredoxin (TsGrx1) from <i>Taenia solium</i> .
Establishment of a secondary infection laboratory model of <i>Echinococcus shiquicus</i> metacystode using BALB/c mice and Mongolian jirds ( <i>Meriones unguiculatus</i> ).
Insight into the Sialome of the Bed Bug, <i>Cimex lectularius</i> .
Manipulation of the immune evasive properties of circulating cathodic antigen induces protective immunity against <i>Schistosomiasis mansoni</i> in C57BL/6 mice.

A new parasitic metacercaria from the land snail <i>Monacha obstructa</i> pfeiffer1842 with critical review on relevant metacercariae belonging to the genus <i>Brachylaima</i> Dujardin 1843.
Crystal structure of Cu/Zn superoxide dismutase from <i>Taenia solium</i> reveals metal-mediated self-assembly.
<i>Moniezia benedeni</i> Infection Restrain IgA(+), IgG(+), and IgM(+) Cells Residence in Sheep ( <i>Ovis aries</i> ) Small Intestine.
[Research progress on the immunity of hepatic echinococcosis in human].
Atomic fidelity of subunit-based chemically-synthesized antimalarial vaccine components.
Resurrection of plant disease resistance proteins via helper NLR bioengineering.
Characterisation of recombinant immunoreactive antigens of the scab mite <i>Sarcoptes scabiei</i> .
Characterization of a diagnostic Fab fragment binding trimeric Lewis X.
Unlike mammalian GRFIN, the zebrafish homologue (DrGRFIN) represents a functional carbohydrate-binding galectin.
Characterization of proteases in the skin mucus of Atlantic salmon ( <i>Salmo salar</i> ) infected with the salmon louse ( <i>Lepeophtheirus salmonis</i> ) and in whole-body louse homogenate.
Bioinformatic analysis suggests potential mechanisms underlying parasitoid venom evolution and function.
From mammals to viruses: the <i>Schlafen</i> genes in developmental, proliferative and immune processes.
Characterization of a multicopy family of genes encoding a surface-expressed serine endoprotease in rat <i>Pneumocystis carinii</i> .
DOCK family proteins: key players in immune surveillance mechanisms.
Presence and structure-activity relationship of intrinsically disordered regions across mucins.
<i>Uronema marinum</i> : identification and biochemical characterization of phosphatidylcholine-hydrolyzing phospholipase C.

Table A2-57, Cluster 56

Cluster 56 focuses on construction of a cancer-associated long non-coding RNA signature to predict prognosis and immune landscape in myriad cancers (794)
Analysis of Ferroptosis-Related LncRNAs Signatures Associated with Tumor Immune Infiltration and Experimental Validation in Clear Cell Renal Cell Carcinoma.
Integrated analysis of 1804 samples of six centers to construct and validate a robust immune-related prognostic signature associated with stromal cell abundance in tumor microenvironment for gastric cancer.
Guidelines on lung adenocarcinoma prognosis based on immuno-glycolysis-related genes.
The LncRNA signature associated with cuproptosis as a novel biomarker of prognosis in immunotherapy and drug screening for clear cell renal cell carcinoma.
Construction and Validation of a Reliable Disulfidptosis-Related LncRNAs Signature of the Subtype, Prognostic, and Immune Landscape in Colon Cancer.
Construction of a cancer-associated fibroblasts-related long non-coding RNA signature to predict prognosis and immune landscape in pancreatic adenocarcinoma.
Construction of a novel immune response prediction signature to predict the efficacy of immune checkpoint inhibitors in clear cell renal cell carcinoma patients.
A Ferroptosis-Related lncRNAs Signature Predicts Prognosis and Therapeutic Response of Gastric Cancer.

Integrating single-cell and bulk RNA sequencing to develop a cancer-associated fibroblast-related signature for immune infiltration prediction and prognosis in lung adenocarcinoma.
A five-cuproptosis-related lncRNA Signature: predicting prognosis, assessing immune function & drug sensitivity in lung squamous cell carcinoma.
Cuproptosis-related lncRNA predict prognosis and immune response of lung adenocarcinoma.
Histone acetylation modification regulator-mediated tumor microenvironment infiltration characteristics and prognostic model of lung adenocarcinoma patients.
N7-Methylguanosine-Related lncRNAs: Integrated Analysis Associated With Prognosis and Progression in Clear Cell Renal Cell Carcinoma.
Identification of cuproptosis-based molecular subtypes, construction of prognostic signature and characterization of immune landscape in colon cancer.
Interferon gamma-related gene signature based on anti-tumor immunity predicts glioma patient prognosis.
Identification of a Six-Immune-Related Long Non-coding RNA Signature for Predicting Survival and Immune Infiltrating Status in Breast Cancer.
Prognostic analysis of RAS-related lncRNAs in liver hepatocellular carcinoma.
Construction and verification of a novel hypoxia-related lncRNA signature related with survival outcomes and immune microenvironment of bladder urothelial carcinoma by weighted gene co-expression network analysis.
A novel signature incorporating lipid metabolism- and immune-related genes to predict the prognosis and immune landscape in hepatocellular carcinoma.
Identification of m6A/m5C/m1A-associated lncRNAs for prognostic assessment and immunotherapy in pancreatic cancer.
Prognosis and Characterization of Immune Microenvironment in Head and Neck Squamous Cell Carcinoma through a Pyroptosis-Related Signature.
Signature and Molecular Mechanism of Mitochondrial Energy Metabolism Pathway-Related Genes in Lung Adenocarcinoma.
A Novel Matrisomal-Related lncRNA Signature Associated With Survival Outcome and Immune Evasion in Patients With Gastric Cancer.
Novel insight on predicting prognosis of gastric cancer based on inflammation.
A novel cuproptosis-related lncRNA nomogram to improve the prognosis prediction of gastric cancer.
Characterization of sialylation-related long noncoding RNAs to develop a novel signature for predicting prognosis, immune landscape, and chemotherapy response in colorectal cancer.
Comprehensive analysis of immune ferroptosis gene in renal clear cell carcinoma: prognosis and influence of tumor microenvironment.
A novel basement membrane-related gene signature for prognosis of lung adenocarcinomas.
PADI1 and Its Co-Expressed Gene Signature Unveil Colorectal Cancer Prognosis and Immunotherapy Efficacy.
PLXDC1 Can Be a Biomarker for Poor Prognosis and Immune Evasion in Gastric Cancer.
Identification of tumor microenvironment-related signature for predicting prognosis and immunotherapy response in patients with bladder cancer.
Construction of a pancreatic cancer prediction model for oxidative stress-related lncRNA.
A new prognostic model for RHOV, ABCC2, and CYP4B1 to predict the prognosis and association with immune infiltration of lung adenocarcinoma.
SARS-CoV-2 Pattern Provides a New Scoring System and Predicts the Prognosis and Immune Therapeutic Response in Glioma.

Illustrating the biological functions and diagnostic value of transmembrane protein family members in glioma.
mRNAsi-related metabolic risk score model identifies poor prognosis, immunoevasive contexture, and low chemotherapy response in colorectal cancer patients through machine learning.
Cuproptosis-Related genes in the prognosis of colorectal cancer and their correlation with the tumor microenvironment.
Identification of necroptosis-related subtypes, development of a novel signature, and characterization of immune infiltration in colorectal cancer.
Development and validation of a novel endoplasmic reticulum stress-related lncRNA prognostic signature and candidate drugs in breast cancer.
Development and validation of an immune-related prognosis signature associated with hypoxia and ferroptosis in hepatocellular carcinoma.
Construction of a prognostic assessment model for colon cancer patients based on immune-related genes and exploration of related immune characteristics.
Comprehensive analysis of karyopherin alpha family expression in lung adenocarcinoma: Association with prognostic value and immune homeostasis.
A methylation- and immune-related lncRNA signature to predict ovarian cancer outcome and uncover mechanisms of chemoresistance.
A neutrophil extracellular traps-associated lncRNA signature predicts the clinical outcomes in patients with lung adenocarcinoma.
The role of lactate metabolism-related lncRNAs in the prognosis, mutation, and tumor microenvironment of papillary thyroid cancer.
Comprehensive genomics analysis of aging related gene signature to predict the prognosis and drug resistance of colon adenocarcinoma.
Classification of Lung Adenocarcinoma Based on Immune Checkpoint and Screening of Related Genes.
In silico development and validation of a novel glucose and lipid metabolism-related gene signature in gastric cancer.
Cuproptosis-Related lncRNA-Based Prediction of the Prognosis and Immunotherapy Response in Papillary Renal Cell Carcinoma.
Development and validation a prognostic model based on natural killer T cells marker genes for predicting prognosis and characterizing immune status in glioblastoma through integrated analysis of single-cell and bulk RNA sequencing.
An Immune-Related Gene Pairs Signature Predicts Prognosis and Immune Heterogeneity in Glioblastoma.
Establishment, immunological analysis, and drug prediction of a prognostic signature of ovarian cancer related to histone acetylation.
Pan-cancer analysis identifies migrasome-related genes as a potential immunotherapeutic target: A bulk omics research and single cell sequencing validation.
Computed tomography-detected extramural venous invasion-related gene signature: a potential negative biomarker of immune checkpoint inhibitor treatment in patients with gastric cancer.
Identifying Hypoxia Characteristics to Stratify Prognosis and Assess the Tumor Immune Microenvironment in Renal Cell Carcinoma.
Identification and Validation of an Immune Evasion Molecular Subgroup of Patients With Colon Cancer for Implications of Immunotherapy.
Identification and validation of a hypoxia-related prognostic and immune microenvironment signature in bladder cancer.

Identification and Validation of a Novel Glycolysis-Related Gene Signature for Predicting the Prognosis and Therapeutic Response in Triple-Negative Breast Cancer.
The immune escape signature predicts the prognosis and immunotherapy sensitivity for pancreatic ductal adenocarcinoma.
Screening and Validation of the Hypoxia-Related Signature of Evaluating Tumor Immune Microenvironment and Predicting Prognosis in Gastric Cancer.
Anoikis-related long non-coding RNA signatures to predict prognosis and small molecular drug response in cervical cancer.
Identification and validation of a prognostic model based on ferroptosis-associated genes in head and neck squamous cancer.
Prognosis signature for predicting the survival and immunotherapy response in esophageal carcinoma based on cellular senescence-related genes.
Identification of a Novel Myc-Regulated Gene Signature for Patients with Kidney Renal Clear Cell Carcinoma.
Prognostic Value of Drug Targets Predicted Using Deep Bioinformatic Analysis of m6A-Associated lncRNA-Based Pancreatic Cancer Model Characteristics and Its Tumour Microenvironment.
Correlation between chromatin epigenetic-related lncRNA signature (CELncSig) and prognosis, immune microenvironment, and immunotherapy in non-small cell lung cancer.
The predictive value of pyroptosis for the prognosis and immune escape of bladder cancer.
Comprehensive analysis of the association between tumor-infiltrating immune cells and the prognosis of lung adenocarcinoma.
Cancer-Associated Fibroblast Risk Model for Prediction of Colorectal Carcinoma Prognosis and Therapeutic Responses.
Development of a polyamine gene expression score for predicting prognosis and treatment response in clear cell renal cell carcinoma.
Novel exosome-related risk signature as prognostic biomarkers in glioblastoma.
P2RY14 downregulation in lung adenocarcinoma: a potential therapeutic target associated with immune infiltration.
A Model of Basement Membrane-Associated Gene Signature Predicts Liver Hepatocellular Carcinoma Response to Immune Checkpoint Inhibitors.
A Novel Defined Pyroptosis-Related Gene Signature for the Prognosis of Acute Myeloid Leukemia.
Metastasis Related Epithelial-Mesenchymal Transition Signature Predicts Prognosis and Response to Immunotherapy in Gastric Cancer.
A novel cuproptosis-related prognostic lncRNA signature for predicting immune and drug therapy response in hepatocellular carcinoma.
Lung adenocarcinoma-specific three-integrin signature contributes to poor outcomes by metastasis and immune escape pathways.
Identification of the collagen family as prognostic biomarkers in papillary thyroid carcinoma.
Macrophage-related molecular subtypes in lung adenocarcinoma identify novel tumor microenvironment with prognostic and therapeutic implications.
Identification of a cuproptosis-related lncRNA prognostic signature in lung adenocarcinoma.
Prognostic Gene Expression Signature in Patients With Distinct Glioma Grades.
Prognosis and Immunotherapy Significances of a Cancer-Associated Fibroblasts-Related Gene Signature in Gliomas.
Prognostic value of neutrophil extracellular trap signature in clear cell renal cell carcinoma.
Identification and Validation of an Immune-Related lncRNA Signature to Facilitate Survival Prediction in Gastric Cancer.

Exploration of Reduced Mitochondrial Content-Associated Gene Signature and Immunocyte Infiltration in Colon Adenocarcinoma by an Integrated Bioinformatic Analysis.
Up-regulated PIF1 predicts poor clinical outcomes and correlates with low immune infiltrates in clear cell renal cell carcinoma.
Bacterial lipopolysaccharide-related genes are involved in the invasion and recurrence of prostate cancer and are related to immune escape based on bioinformatics analysis.
Identification of a Hypoxia-Related Molecular Classification and Hypoxic Tumor Microenvironment Signature for Predicting the Prognosis of Patients with Triple-Negative Breast Cancer.
Development and Validation of Tumor Immunogenicity Based Gene Signature for Skin Cancer Risk Stratification.
Comprehensive exploration of tumor mutational burden and immune infiltration in diffuse glioma.
N6-Methyladenosine (m(6)A)-Related lncRNAs Are Potential Signatures for Predicting Prognosis and Immune Response in Lung Squamous Cell Carcinoma.
Prognostic and Predictive Value of Cadherin 11 for Patients with Gastric Cancer and Its Correlation with Tumor Microenvironment: Results from Microarray Analysis.
Pan-cancer analysis of GALNTs expression identifies a prognostic of GALNTs feature in low grade glioma.
Chordin-Like 2: A Possible Therapeutic Target for Gastric Cancer by Affecting Cell Cycle and Proliferation.
Comprehensive Analysis of Cellular Senescence-Related Genes in Prognosis, Molecular Characterization and Immunotherapy of Hepatocellular Carcinoma.
Multi-Omics Profiling Identifies Risk Hypoxia-Related Signatures for Ovarian Cancer Prognosis.
Low expression of the metabolism-related gene SLC25A21 predicts unfavourable prognosis in patients with acute myeloid leukaemia.
Identification and validation of a metabolism-related model and associated with tumor-infiltrating lymphocytes in p53 mutant lung adenocarcinoma patients.
A SERPINE1-Based Immune Gene Signature Predicts Prognosis and Immunotherapy Response in Gastric Cancer.
Analysis of Genome-Wide Alternative Splicing Profiling and Development of Potential Drugs in Lung Adenocarcinoma.
Identification of N7-methylguanosine related subtypes and construction of prognostic model in gastric cancer.
NF- $\kappa$ B-Related Metabolic Gene Signature Predicts the Prognosis and Immunotherapy Response in Gastric Cancer.
Establishment and validation of an immune-based prognostic score model in glioblastoma.
Identification of candidate biomarker EMP3 and its prognostic potential in clear cell renal cell carcinoma.
Comprehensive Prognostic Analysis of Immune Implication Value and Oxidative Stress Significance of NECAP2 in Low-Grade Glioma.
Necroptosis-Related Prognostic Model for Pancreatic Carcinoma Reveals Its Invasion and Metastasis Potential through Hybrid EMT and Immune Escape.
Tumor-associated macrophages related signature in glioma.
Leveraging Tumor Microenvironment Infiltration in Pancreatic Cancer to Identify Gene Signatures Related to Prognosis and Immunotherapy Response.
Pan-cancer analysis based on epigenetic modification explains the value of HJURP in the tumor microenvironment.

Comprehensive Analysis of N6-Methyladenosine-Related lncRNAs Prognostic Signature and the Associated Immune Infiltrates in Kidney Renal Clear Cell Carcinoma.
Potential prognostic biomarkers related to immunity in clear cell renal cell carcinoma using bioinformatic strategy.
Identification of Immune Cell Infiltration and Immune-Related Genes in the Tumor Microenvironment of Glioblastomas.
Immune-Related lncRNAs Affect the Prognosis of Osteosarcoma, Which Are Related to the Tumor Immune Microenvironment.
[Expression characteristics and functional analysis of ELK3 in gastric cancer].
Prognostic significance of HSF2BP in lung adenocarcinoma.
New Prognostic Gene Signature and Immune Escape Mechanisms of Bladder Cancer.
Ferroptosis-related lncRNAs signature to predict the survival and immune evasion for lung squamous cell carcinoma.
Identification of an Individualized Metabolism Prognostic Signature and Related Therapy Regimens in Early Stage Lung Adenocarcinoma.
Immune-related gene signature associates with immune landscape and predicts prognosis accurately in patients with Wilms tumour.
Machine learning-based on cytotoxic T lymphocyte evasion gene develops a novel signature to predict prognosis and immunotherapy responses for kidney renal clear cell carcinoma patients.
A signature based on anoikis-related genes for the evaluation of prognosis, immunoinfiltration, mutation, and therapeutic response in ovarian cancer.
Identification of INSRR as an immune-related gene in the tumor microenvironment of glioblastoma by integrated bioinformatics analysis.
Bioinformatic analysis of cancer-associated fibroblast related gene signature as a predictive model in clinical outcomes and immune characteristics of gastric cancer.
Autophagy-Dependent Ferroptosis-Related Signature is Closely Associated with the Prognosis and Tumor Immune Escape of Patients with Glioma.
Identification and validation of dichotomous immune subtypes based on intratumoral immune cells infiltration in clear cell renal cell carcinoma patients.
A novel T-cell proliferation-associated regulator signature pre-operatively predicted the prognostic of bladder cancer.
TAGLN2 Promotes the Proliferation, Migration, Invasion, and EMT of Clear Cell Renal Cell Carcinoma Through the PI3K/Akt Signaling Pathway.
Calbindin S100A16 Promotes Renal Cell Carcinoma Progression and Angiogenesis via the VEGF/VEGFR2 Signaling Pathway.
Homologous recombination repair gene-based risk model predicts prognosis and immune microenvironment for primary lung cancer after previous malignancies.
Could CTSK and COL4A2 be specific biomarkers of poor prognosis for patients with gastric cancer in Asia?-a microarray analysis based on regional population.
Establishment of lactate-metabolism-related signature to predict prognosis and immunotherapy response in patients with colon adenocarcinoma.
Genetic landscape of prognostic value in pancreatic ductal adenocarcinoma microenvironment.
Integrating Bulk and Single-Cell RNA Sequencing Reveals Heterogeneity, Tumor Microenvironment, and Immunotherapeutic Efficacy Based on Sialylation-Related Genes in Bladder Cancer.
Identification and validation of a novel cuproptosis-related lncRNA gene signature to predict prognosis and immune response in bladder cancer.
Prognostic model based on m6A-associated lncRNAs in esophageal cancer.



KLHDC8A Expression in Association with Macrophage Infiltration and Oxidative Stress Predicts Unfavorable Prognosis for Glioma.
Development and validation of a necroptosis-related gene prognostic score to predict prognosis and efficiency of immunotherapy in gastric cancer.
Comprehensive Profiling Reveals Distinct Microenvironment and Metabolism Characterization of Lung Adenocarcinoma.
An integrated bioinformatic investigation of mitochondrial energy metabolism genes in colon adenocarcinoma followed by preliminary validation of CPT2 in tumor immune infiltration.
Development and Validation of a Prognostic Autophagy-Related Gene Pair Index Related to Tumor-Infiltrating Lymphocytes in Early-Stage Lung Adenocarcinoma.
Fibroblast Growth Factor 11 Enables Tumor Cell Immune Escape by Promoting T Cell Exhaustion and Predicts Poor Prognosis in Patients with Lung Adenocarcinoma.
Development and Validation of a TNF Family-Based Signature for Predicting Prognosis, Tumor Immune Characteristics, and Immunotherapy Response in Colorectal Cancer Patients.
An Immune-Related Prognostic Signature for Predicting Clinical Outcomes and Immune Landscape in IDH-Mutant Lower-Grade Gliomas.
Prognostic values of the immune microenvironment-related non-coding RNA IGF2BP2-AS1 in bladder cancer.
Identification and validation of RB1 as an immune-related prognostic signature based on tumor mutation burdens in bladder cancer.
Prediction of hepatocellular carcinoma prognosis and immunotherapeutic effects based on tryptophan metabolism-related genes.
Schlafen family is a prognostic biomarker and corresponds with immune infiltration in gastric cancer.
Membrane tension sensing molecule-FNBP1 is a prognostic biomarker related to immune infiltration in BRCA, LUAD and STAD.
A novel 25-ferroptosis-related gene signature for the prognosis of gliomas.
Genomic and Immunological Characterization of Pyroptosis in Lung Adenocarcinoma.
Identifying tumour microenvironment-related signature that correlates with prognosis and immunotherapy response in breast cancer.
OLR1 Is a Pan-Cancer Prognostic and Immunotherapeutic Predictor Associated with EMT and Cuproptosis in HNSCC.
Prediction and identification of immune genes related to the prognosis of patients with colon adenocarcinoma and its mechanisms.
Epithelial-mesenchymal transition-associated secretory phenotype predicts survival in lung cancer patients.
Development and validation of cancer-associated fibroblasts-related gene landscape in prognosis and immune microenvironment of bladder cancer.
A prognostic risk model for patients with triple negative breast cancer based on stromal natural killer cells, tumor-associated macrophages and growth-arrest specific protein 6.
A systematic and comprehensive analysis of T cell exhaustion related to therapy in lung adenocarcinoma tumor microenvironment.
Gastric cancer peritoneal metastasis related signature predicts prognosis and sensitivity to immunotherapy in gastric cancer.
ARHGAP11A Is a Novel Prognostic and Predictive Biomarker Correlated with Immunosuppressive Microenvironment in Clear Cell Renal Cell Carcinoma.
High Expression of the Component 3a Receptor 1 (C3AR1) Gene in Stomach Adenocarcinomas Infers a Poor Prognosis and High Immune-Infiltration Levels.

Integrated pan-cancer analysis of CSMD2 as a potential prognostic, diagnostic, and immune biomarker.
Identification and validation of an immune-related gene prognostic signature for clear cell renal carcinoma.
Hexokinase 3 dysfunction promotes tumorigenesis and immune escape by upregulating monocyte/macrophage infiltration into the clear cell renal cell carcinoma microenvironment.
Identification of Prognostic Biomarkers Originating From the Tumor Stroma of Betel Quid-Associated Oral Cancer Tissues.
Integrative analysis of immune molecular subtypes and microenvironment characteristics of bladder cancer.
A Novel TAF-Related Signature Based on ECM Remodeling Genes Predicts Glioma Prognosis.
A damage-associated molecular patterns-related gene signature for the prediction of prognosis and immune microenvironment in children stage III acute lymphoblastic leukemia.
Integrated Analysis of RNA-Binding Proteins Associated With the Prognosis and Immunosuppression in Squamous Cell Carcinoma of Head and Neck.
Integrated Analysis of the Transcriptome Profile Reveals the Potential Roles Played by Long Noncoding RNAs in Immunotherapy for Sarcoma.
A novel immune prognostic model of non-M3 acute myeloid leukemia.
Comprehensive landscape of the miRNA-regulated prognostic marker LAYN with immune infiltration and stemness in pan-cancer.
Exploration of s new biomarker in osteosarcoma and association with clinical outcomes: (TOP2A+) cancer associated fibroblasts.
Immune Landscape Refines the Classification of Colorectal Cancer With Heterogeneous Prognosis, Tumor Microenvironment and Distinct Sensitivity to Frontline Therapies.
Comparative Analysis and in vitro Experiments of Signatures and Prognostic Value of Immune Checkpoint Genes in Colorectal Cancer.
PCK2 inhibits lung adenocarcinoma tumor cell immune escape through oxidative stress-induced senescence as a potential therapeutic target.
A novel ganglioside-related risk signature can reveal the distinct immune landscape of neuroblastoma and predict the immunotherapeutic response.
Machine learning-based identification of glycosyltransferase-related mRNAs for improving outcomes and the anti-tumor therapeutic response of gliomas.
Construction of an Immune Escape-Related Signature in Clear Cell Renal Cell Carcinoma and Identification of the Relationship between IFNAR1 and Immune Infiltration by Multiple Immunohistochemistry.
Weighted Gene Co-Expression Network Analysis (WGCNA) Reveals the Functions of Syndecan-1 to Regulate Immune Infiltration by Influenced T Cells in Glioma.
Integrated Analysis of Necroptosis-Related Genes for Prognosis, Immune Microenvironment Infiltration, and Drug Sensitivity in Colon Cancer.
CHSY3 can be a Poor Prognostic Biomarker and Mediates Immune Evasion in Stomach Adenocarcinoma.
A novel pyroptosis-related signature predicts prognosis and indicates immunotherapy in oral squamous cell carcinoma.
Deoxythymidylate kinase (DTYMK) participates in cell cycle arrest to promote pancreatic adenocarcinoma progression regulated by miR-491-5p through TP53 and is associated with tumor immune infiltration.

Upregulated Immunogenic Cell-Death-Associated Gene Signature Predicts Reduced Responsiveness to Immune-Checkpoint-Blockade Therapy and Poor Prognosis in High-Grade Gliomas.
Discovery of a Novel Immune Gene Signature with Profound Prognostic Value in Colorectal Cancer: A Model of Cooperativity Disorientation Created in the Process from Development to Cancer.
Tumor-intrinsic and -extrinsic (immune) gene signatures robustly predict overall survival and treatment response in high grade serous ovarian cancer patients.
Role of ICAM1 in tumor immunity and prognosis of triple-negative breast cancer.
Comprehensive Characterization of RNA-Binding Proteins in Colon Adenocarcinoma Identifies a Novel Prognostic Signature for Predicting Clinical Outcomes and Immunotherapy Responses Based on Machine Learning.
Bioinformatics Analysis of GFAP as a Potential Key Regulator in Different Immune Phenotypes of Prostate Cancer.
A Pan-Cancer Analysis of UBE2S in Tumorigenesis, Prognosis, Pathway, Immune Infiltration and Evasion, and Therapy Response from an Immune-Oncology Perspective.
Distinct signatures of lung cancer types: aberrant mucin O-glycosylation and compromised immune response.
Exploring the correlation analysis of immune microenvironment, mutation burden and prognosis of papillary thyroid carcinoma based on Estimate algorithm.
Pan-cancer analysis of chromothripsis-related gene expression patterns indicates an association with tumor immune and therapeutic agent responses.
Identification of fatty acid metabolism-related molecular subtype biomarkers and their correlation with immune checkpoints in cutaneous melanoma.
Expression of CYP1B1 and B7-H3 significantly correlates with poor prognosis in colorectal cancer patients.
Comprehensive Analysis of the Relationship Between Metabolic Reprogramming and Immune Function in Prostate Cancer.
CTLA4-Mediated Immunosuppression in Glioblastoma is Associated with the Infiltration of Macrophages in the Tumor Microenvironment.
Aggressive early-stage lung adenocarcinoma is characterized by epithelial cell plasticity with acquirement of stem-like traits and immune evasion phenotype.
A novel prognostic biomarker: GINS3 is correlated with methylation and immune escape in liver hepatocellular carcinoma.
Identification of a chromatin regulator signature for predicting prognosis of prostate cancer patient.
Diverse Macrophages Constituted the Glioma Microenvironment and Influenced by PTEN Status.
Prediction of the immunological and prognostic value of five signatures related to fatty acid metabolism in patients with cervical cancer.
KCNN4 may weaken anti-tumor immune response via raising Tregs and diminishing resting mast cells in clear cell renal cell carcinoma.
Multiple bioinformatics analysis identifies IGFBP1 as associated with the prognosis of stomach adenocarcinoma.
MMP25-AS1/hsa-miR-10a-5p/SERPINE1 axis as a novel prognostic biomarker associated with immune cell infiltration in KIRC.
Pan-cancer analysis of the prognostic and immunological role of ANLN: An onco-immunological biomarker.
Pan-Cancer Analysis Shows Enrichment of Macrophages, Overexpression of Checkpoint Molecules, Inhibitory Cytokines, and Immune Exhaustion Signatures in EMT-High Tumors.

Autophagy-related gene expression classification defines three molecular subtypes with distinct clinical and microenvironment cell infiltration characteristics in colon cancer.
INHBA is a novel mediator regulating cellular senescence and immune evasion in colorectal cancer.
Vemurafenib inhibits immune escape biomarker BCL2A1 by targeting PI3K/AKT signaling pathway to suppress breast cancer.
Expression, tumor immune infiltration, and prognostic impact of HMGs in gastric cancer.
An immunogenic cell death-related signature predicts prognosis and immunotherapy response in stomach adenocarcinoma.
Comprehensive Pan-Cancer Analysis of TRPM8 in Tumor Metabolism and Immune Escape.
Systematic integration of machine learning algorithms to develop immune escape-related signatures to improve clinical outcomes in lung adenocarcinoma patients.
Decoding Immune Heterogeneity of Melanoma and identifying immune-prognostic hub genes.
Immune Infiltrating Cells-Derived Risk Signature Based on Large-scale Analysis Defines Immune Landscape and Predicts Immunotherapy Responses in Glioma Tumor Microenvironment.
MET overexpression contributes to STAT4-PD-L1 signaling activation associated with tumor-associated, macrophages-mediated immunosuppression in primary glioblastomas.
KIF11 As a Potential Pan-Cancer Immunological Biomarker Encompassing the Disease Staging, Prognoses, Tumor Microenvironment, and Therapeutic Responses.
MXD3 as an onco-immunological biomarker encompassing the tumor microenvironment, disease staging, prognoses, and therapeutic responses in multiple cancer types.
Identifying immune subtypes of uterine corpus endometrial carcinoma and a four-paired-lncRNA signature with immune-related lncRNAs.
Intrinsic immune evasion patterns predict temozolomide sensitivity and immunotherapy response in lower-grade gliomas.
Identification of Endoplasmic Reticulum Stress-Related Subtypes, Infiltration Analysis of Tumor Microenvironment, and Construction of a Prognostic Model in Colorectal Cancer.
Single-cell profiling of the copy-number heterogeneity in colorectal cancer.
Identification and Validation of a Novel Multiomics Signature for Prognosis and Immunotherapy Response of Endometrial Carcinoma.
The CXCL Family Contributes to Immunosuppressive Microenvironment in Gliomas and Assists in Gliomas Chemotherapy.
STEAP3 can predict the prognosis and shape the tumor microenvironment of clear cell renal cell carcinoma.
Alteration in glycolytic/cholesterogenic gene expression is associated with bladder cancer prognosis and immune cell infiltration.
The Clinical Significance of DNA Damage Repair Signatures in Clear Cell Renal Cell Carcinoma.
miR-1251-5p Overexpression Inhibits Proliferation, Migration, and Immune Escape in Clear Cell Renal Cell Carcinoma by Targeting NPTX2.
ATP2C2 Has Potential to Define Tumor Microenvironment in Breast Cancer.
Immune cell infiltration signatures identified molecular subtypes and underlying mechanisms in gastric cancer.
A gene prognostic index from cellular senescence predicting metastasis and radioresistance for prostate cancer.
Molecular profiling of multiple human cancers defines an inflammatory cancer-associated molecular pattern and uncovers KPNA2 as a uniform poor prognostic cancer marker.
Ribonucleotide reductase small subunit M2 is a master driver of aggressive prostate cancer.

LINC00638/hsa-miR-29b-3p axis-mediated high expression of CDCA4 correlates with tumor immune infiltration and hepatocellular carcinoma progression.
Cuproptosis correlates with immunosuppressive tumor microenvironment based on pan-cancer multiomics and single-cell sequencing analysis.
Identification and verification of YBX3 and its regulatory gene HEIH as an oncogenic system: A multidimensional analysis in colon cancer.
Investigating the correlation between DNA methylation and immune-associated genes of lung adenocarcinoma based on a competing endogenous RNA network.
Identification of Cancer Hub Gene Signatures Associated with Immune-Suppressive Tumor Microenvironment and Ovatodiolide as a Potential Cancer Immunotherapeutic Agent.
SWATH-based proteomics reveals processes associated with immune evasion and metastasis in poor prognosis colorectal tumours.
PD-L1 Activity Is Associated with Partial EMT and Metabolic Reprogramming in Carcinomas.
Novel Cuproptosis-Related Gene Signature for Precise Identification of High-Risk Populations in Low-Grade Gliomas.
Integrative analysis of imaging and transcriptomic data of the immune landscape associated with tumor metabolism in lung adenocarcinoma: Clinical and prognostic implications.
ATG9B Is a Poor Prognostic Marker Associated With Immune Evasion in Colon Adenocarcinoma.
Poor Clinical Outcomes and Immuno-evasive Contexture in Intratumoral IL-10-Producing Macrophages Enriched Gastric Cancer Patients.
Epigenetically regulated lncRNAs dissect the intratumoural heterogeneity and facilitate immune evasion of glioblastomas.
Long non-coding RNAs involved in metastasis of gastric cancer.
Hyper-Methylated Hub Genes of T-Cell Receptor Signaling Predict a Poor Clinical Outcome in Lung Adenocarcinoma.
CD44 Is a Prognostic Biomarker Correlated With Immune Infiltrates and Metastasis in Clear Cell Renal Cell Carcinoma.
Construction of lncRNA prognostic model related to cuproptosis in esophageal carcinoma.
Novel Immune Infiltrating Cell Signature Based on Cell Pair Algorithm Is a Prognostic Marker in Cancer.
Comprehensive Molecular Analyses of a Novel Mutational Signature Classification System with Regard to Prognosis, Genomic Alterations, and Immune Landscape in Glioma.
Decreased m6A Modification of CD34/CD276(B7-H3) Leads to Immune Escape in Colon Cancer.
LCTL Is a Prognostic Biomarker and Correlates With Stromal and Immune Infiltration in Gliomas.
PDZ Binding Kinase/T-LAK Cell-Derived Protein Kinase Plays an Oncogenic Role and Promotes Immune Escape in Human Tumors.
Integration of single sample and population analysis for understanding immune evasion mechanisms of lung cancer.
A gene-based survival score for lung adenocarcinoma by multiple transcriptional datasets analysis.
Comprehensive analysis of TAMs marker genes in glioma for predicting prognosis and immunotherapy response.
lncRNA and its role in gastric cancer immunotherapy.
Pan-cancer analysis of clinical significance and associated molecular features of glycolysis.
CD248 as a bridge between angiogenesis and immunosuppression: a promising prognostic and therapeutic target for renal cell carcinoma.
Integrated DNA and RNA sequencing reveals early drivers involved in metastasis of gastric cancer.

Identification of a Glycosyltransferase Signature for Predicting Prognosis and Immune Microenvironment in Neuroblastoma.
Relationships Between Immune Landscapes, Genetic Subtypes and Responses to Immunotherapy in Colorectal Cancer.
Prognostic Value and Immunological Role of KIFC1 in Hepatocellular Carcinoma.
Signatures of immune cell infiltration for predicting immune escape and immunotherapy in cervical cancer.
Inhibition of NOS1 promotes the interferon response of melanoma cells.
Bioinformatic and Experimental Analyses Reveal That KIF4A Is a Biomarker of Therapeutic Sensitivity and Predicts Prognosis in Cervical Cancer Patients.
Immunological landscape of consensus clusters in colorectal cancer.
Comparative study of gene expression by cDNA microarray in human colorectal cancer tissues and normal mucosa.
Deep learning-based morphological feature analysis and the prognostic association study in colon adenocarcinoma histopathological images.
A Gene Prognostic Index Associated With Epithelial-Mesenchymal Transition Predicting Biochemical Recurrence and Tumor Chemoresistance for Prostate Cancer.
lncRNAs as Hallmarks for Individualized Treatment of Gastric Cancer.
Single-cell mapping of N6-methyladenosine in esophageal squamous cell carcinoma and exploration of the risk model for immune infiltration.
The immune phenotypes and different immune escape mechanisms in colorectal cancer.
Identification of lactylation related model to predict prognostic, tumor infiltrating immunocytes and response of immunotherapy in gastric cancer.
Integrated analysis of single cell and bulk RNA sequencing identifies CTHRC1(+) INHBA(+) CAF as drivers of colorectal cancer progression.
Expression Profile Analysis of m6A RNA Methylation Regulators Indicates They Are Immune Signature Associated and Can Predict Survival in Kidney Renal Cell Carcinoma.
A Ferroptosis-Related Prognostic Risk Score Model to Predict Clinical Significance and Immunogenic Characteristics in Glioblastoma Multiforme.
Reverse engineering a predictive signature characterized by proliferation, DNA damage, and immune escape from stage I lung adenocarcinoma recurrence.
CD2 Is a Novel Immune-Related Prognostic Biomarker of Invasive Breast Carcinoma That Modulates the Tumor Microenvironment.
LINE-1 promotes tumorigenicity and exacerbates tumor progression via stimulating metabolism reprogramming in non-small cell lung cancer.
Long Noncoding RNA: Shining Stars in the Immune Microenvironment of Gastric Cancer.
Large-scale bulk RNA-seq analysis defines immune evasion mechanism related to mast cell in gliomas.
Research Progress on Long Non-Coding RNA and Radiotherapy.
Immune Landscape in PTEN-Related Glioma Microenvironment: A Bioinformatic Analysis.
Long Non-Coding RNAs (lncRNAs) in Response and Resistance to Cancer Immunosurveillance and Immunotherapy.
Prognostic landscape of tumor-infiltrating immune cells and immune-related genes in the tumor microenvironment of gastric cancer.
SUMOylation patterns and signature characterize the tumor microenvironment and predict prognosis in lung adenocarcinoma.
Molecular Landscape of lncRNAs in Prostate Cancer: A focus on pathways and therapeutic targets for intervention.

Long non-coding RNAs: A view to kill ovarian cancer.
Prediction of prognosis, immune infiltration and immunotherapy response with N6-methyladenosine-related lncRNA clustering patterns in cervical cancer.
A Ferroptosis-Related Gene Prognostic Index Associated With Biochemical Recurrence and Radiation Resistance for Patients With Prostate Cancer Undergoing Radical Radiotherapy.
The Expression Pattern of Ferroptosis-Related Genes in Colon Adenocarcinoma: Highly Correlated to Tumor Microenvironment Characteristics.
DNA damage repair gene mutations and their association with tumor immune regulatory gene expression in muscle invasive bladder cancer subtypes.
Cuproptosis facilitates immune activation but promotes immune escape, and a machine learning-based cuproptosis-related signature is identified for predicting prognosis and immunotherapy response of gliomas.
A pan-cancer analysis of the oncogenic role of secreted phosphoprotein 1 (SPP1) in human cancers.
Identification of Four Immune Subtypes in Bladder Cancer Based on Immune Gene Sets.
Landscape of associations between long non-coding RNAs and infiltrating immune cells in liver hepatocellular carcinoma.
A PRRX1 Signature Identifies TIM-3 and VISTA as Potential Immune Checkpoint Targets in a Subgroup of Microsatellite Stable Colorectal Cancer Liver Metastases.
Association between orosomucoid types and cancer.
Transcriptional effects of copy number alterations in a large set of human cancers.
Integrative Molecular Characterization of Resistance to Neoadjuvant Chemoradiation in Rectal Cancer.
HMGB1/RAGE axis mediates the apoptosis, invasion, autophagy, and angiogenesis of the renal cell carcinoma.
An NF- $\kappa$ B- and Therapy-Related Regulatory Network in Glioma: A Potential Mechanism of Action for Natural Antiglioma Agents.
[Expression of MUC1 and distribution of tumor-infiltrating dendritic cells in human bladder transitional cell carcinoma].
Targeting bladder cancer: A sex sensitive perspective in mutations and outcomes.
PBK phosphorylates MSL1 to elicit epigenetic modulation of CD276 in nasopharyngeal carcinoma.
The coding microsatellite mutation profile of PMS2-deficient colorectal cancer.
Differential chemokine, chemokine receptor, cytokine and cytokine receptor expression in pulmonary adenocarcinoma: diffuse down-regulation is associated with immune evasion and brain metastasis.
An open-label, nonrandomized, phase Ib feasibility study of cusatuzumab in patients with nasopharyngeal carcinoma.
Better together: circulating tumor cell clustering in metastatic cancer.
Chemokine receptors in epithelial ovarian cancer.
Characterization of Macrophage Galactose-type Lectin (MGL) ligands in colorectal cancer cell lines.
Pentraxin 3: a novel biomarker for predicting progression from prostatic inflammation to prostate cancer.
Kidney tumor biomarkers revealed by simultaneous multiple matrix metabolomics analysis.
STAT5a-targeting miRNA enhances chemosensitivity to cisplatin and 5-fluorouracil in human colorectal cancer cells.
Genomic and transcriptomic characterization of heterogeneous immune subgroups of microsatellite instability-high colorectal cancers.
Checkpoint inhibitors: the new treatment paradigm for urothelial bladder cancer.

Canonical and truncated transglutaminase-2 regulate mucin-1 expression and androgen independency in prostate cancer cell lines.
Interleukin-10 gene promoter polymorphisms and the risk of nasopharyngeal carcinoma.
Long noncoding RNA amplified in lung cancer rewires cancer pathways.
Radiation-induced cancer: a modern view.
Metastasis of lung adenocarcinoma to the gingiva: a rare case report.
DUX4 is a common driver of immune evasion and immunotherapy failure in metastatic cancers.
Paths of Evolution of Progressive Anaplastic Meningiomas: A Clinical and Molecular Pathology Study.
Chronic inflammation in endometriosis and endometriosis-associated ovarian cancer: New roles for the "old" complement pathway.
HER-2-induced PI3K signaling pathway was involved in the pathogenesis of gastric cancer.
Characterization of the Roles of Suppressor of Cytokine Signaling-3 in Esophageal Carcinoma.
Association of interleukin-10 (IL10) promoter genotypes with nasopharyngeal carcinoma risk in Taiwan.
The role of B7 costimulation in benzene immunotoxicity and its potential association with cancer risk.
IL-12 regulates B7-H1 expression in ovarian cancer-associated macrophages by effects on NF- $\kappa$ B signalling.
[Pathogenesis and treatment of immune dysregulation associated with myelodysplastic syndromes].
Large-scale genomic analyses reveal alterations and mechanisms underlying clonal evolution and immune evasion in esophageal cancer.
N(6)-Methyladenosine-Modified CBX1 Regulates Nasopharyngeal Carcinoma Progression Through Heterochromatin Formation and STAT1 Activation.
CDH1 loss promotes diffuse-type gastric cancer tumorigenesis via epigenetic reprogramming and immune evasion.
Convergence of YAP/TAZ, TEAD and TP63 activity is associated with bronchial premalignant severity and progression.
The biology of micrometastases from uveal melanoma.
Epigenetic Suppression of the IL-7 Pathway in Progressive Glioblastoma.
The Changing Treatment Landscape for Metastatic Urothelial Carcinoma.
Significant association of Interleukin-10 genotypes and oral cancer susceptibility in Taiwan.
Evaluation of immune response according to the metastatic status in the regional lymph nodes in patients with gastric carcinoma.
Multomic analysis of homologous recombination-deficient end-stage high-grade serous ovarian cancer.
Intron retention-induced neoantigen load correlates with unfavorable prognosis in multiple myeloma.
Expression of the immune modulator secretory leukocyte protease inhibitor (SLPI) in colorectal cancer liver metastases and matched primary tumors is associated with a poorer prognosis.
Elevated Expression of the Immune Checkpoint Ligand CD276 (B7-H3) in Urothelial Carcinoma Cell Lines Correlates Negatively with the Cell Proliferation.
Th17 Cells and IL-17 As Novel Immune Targets in Ovarian Cancer Therapy.
Primary refractory plasmablastic lymphoma: A precision oncology approach.
Potential mechanisms underlying CDK5 related Osteosarcoma progression.
Adhesion of lymphocytes to bladder cancer cells: the role of the $\alpha$ (E) $\beta$ (7) integrin.
STAG2 loss-of-function mutation induces PD-L1 expression in U2OS cells.
Merkel Cell Carcinoma from Molecular Pathology to Novel Therapies.
Posttranscriptional regulation of cancer traits by HuR.



Progression of Metastasis through Lymphatic System.
GARP promotes the proliferation and therapeutic resistance of bone sarcoma cancer cells through the activation of TGF- $\beta$ .
Correction to: The lncRNA UCA1 promotes proliferation, migration, immune escape and inhibits apoptosis in gastric cancer by sponging anti-tumor miRNAs.
Replication stress defines distinct molecular subtypes across cancers.
Effects of miR-363 on the Biological Activities of Eutopic Endometrial Stromal Cells in Endometriosis.
Expression and distribution of S-100, CD83, and costimulatory molecules (CD80 and CD86) in tissues of thyroid papillary carcinoma.
Exploring the biological hallmarks of cancer of unknown primary: where do we stand today?
Pregnancy-Associated Plasma Protein-A (PAPP-A) in Ewing Sarcoma: Role in Tumor Growth and Immune Evasion.
Expression of EBAG9/RCA51 is associated with advanced disease in human epithelial ovarian cancer.
Association of Serum Immunoglobulin Levels with Solid Cancer: A Systematic Review and Meta-analysis.
Longitudinal analysis and prognostic effect of cancer-testis antigen expression in multiple myeloma.
IRGM promotes glioma M2 macrophage polarization through p62/TRAF6/NF- $\kappa$ B pathway mediated IL-8 production.
Serum and peritoneal fluid concentrations of soluble human leukocyte antigen, tumor necrosis factor alpha and interleukin 10 in patients with selected ovarian pathologies.
Resistance to CD95 (APO-1/Fas)-mediated apoptosis in human renal cell carcinomas: an important factor for evasion from negative growth control.
The m6A Reader YTHDF2 Promotes Bladder Cancer Progression by Suppressing RIG-I-Mediated Immune Response.
Helicobacter pylori and serum kynurenine-tryptophan ratio in patients with colorectal cancer.
Frontiers in the Pathology and Pathogenesis of Ovarian Cancer: Cancer Precursors and "Precursor Escape".
[Significance of the unbalanced expression of Th1/Th2 type cytokines in human glioma].
Genomic Alterations during the In Situ to Invasive Ductal Breast Carcinoma Transition Shaped by the Immune System.
$\beta$ -catenin regulates IRF3-mediated innate immune signalling in colorectal cancer.
Obesity alters the mouse endometrial transcriptome in a cell context-dependent manner.
Upregulation of Yin-Yang-1 Associates with Proliferation and Glutamine Metabolism in Esophageal Carcinoma.
Neuroblastoma cells negative for CD44 possess tumor-initiating properties.
Lytic induction and apoptosis of Epstein-Barr virus-associated gastric cancer cell line with epigenetic modifiers and ganciclovir.
Circulating tumor cell clusters: Insights into tumour dissemination and metastasis.
The yin-yang of immunity: Immune dysregulation in myelodysplastic syndrome with different risk stratification.
ADAM protein family - its role in tumorigenesis, mechanisms of chemoresistance and potential as diagnostic and prognostic factors.
DNA hypermethylation driven by DNMT1 and DNMT3A favors tumor immune escape contributing to the aggressiveness of adrenocortical carcinoma.
Spatially resolved, high-dimensional transcriptomics sorts out the evolution of biphasic malignant pleural mesothelioma: new paradigms for immunotherapy.

MYC Activity Inference Captures Diverse Mechanisms of Aberrant MYC Pathway Activation in Human Cancers.
Immune oppression array elucidating immune escape and survival mechanisms in uveal melanoma.
Y Chromosome Loss Drives Bladder Cancer Aggressiveness and Immune Evasion.
MUC1-C Oncoprotein Integrates a Program of EMT, Epigenetic Reprogramming and Immune Evasion in Human Carcinomas.
Renal cell carcinoma alters endothelial receptor expression responsible for leukocyte adhesion.
Isolation and molecular characterization of the 5'-upstream region of the human TRAIL gene.
High Immune Expression of Progesterone-Induced Blocking Factor in Epithelial Ovarian Cancer.
SNP array-based karyotyping: differences and similarities between aplastic anemia and hypocellular myelodysplastic syndromes.
Perspective: APOBEC mutagenesis in drug resistance and immune escape in HIV and cancer evolution.
Severe preeclampsia is characterized by increased placental expression of galectin-1.
[Expression of IgVH and B7-1 in proteome of the human colorectal carcinoma cell lines].
Morphological characteristics and co-stimulatory molecule (CD80, CD86, CD40) expression in tumor infiltrating dendritic cells in human endometrioid adenocarcinoma.
Recent advances in understanding clonal haematopoiesis in aplastic anaemia.
Anatomic and molecular pathology of intrahepatic cholangiocarcinoma.
Estrogen-regulated CD200 inhibits macrophage phagocytosis in endometriosis.
Retracted: LncRNA OIP5-AS1 Knockdown Facilitated the Ferroptosis and Immune Evasion by Modulating the GPX4 in Oesophageal Carcinoma.
Granulomas of intestinal tuberculosis and Crohn's disease can be differentiated by CD73 cell surface marker expression: a pilot study.
Management of nonviral mixed cryoglobulinemia vasculitis refractory to rituximab: Data from a European collaborative study and review of the literature.

Table A2-58, Cluster 57

Cluster 57 focuses on bacteria (especially <i>Neisseria meningitidis</i> ), emphasizing their pathogenesis and immune suppressive mechanisms (394)
Cellular and molecular biology of <i>Neisseria meningitidis</i> colonization and invasive disease.
Sialic Acid Protects Nontypeable <i>Haemophilus influenzae</i> from Natural IgM and Promotes Survival in Murine Respiratory Tract.
Mechanisms of avoidance of host immunity by <i>Neisseria meningitidis</i> and its effect on vaccine development.
Underlying Glycans Determine the Ability of Sialylated Lipooligosaccharide To Protect Nontypeable <i>Haemophilus influenzae</i> from Serum IgM and Complement.
Commensal <i>Neisseria</i> species share immune suppressive mechanisms with <i>Neisseria gonorrhoeae</i> .
Nontypeable <i>Haemophilus influenzae</i> : pathogenesis and prevention.
<i>Treponema pallidum</i> subsp. <i>pallidum</i> with an Artificially Impaired TprK Antigenic Variation System is Attenuated in the Rabbit Model of Syphilis.
<i>Neisseria meningitidis</i> : biology, microbiology, and epidemiology.
Phase variation of DNA methyltransferases and the regulation of virulence and immune evasion in the pathogenic <i>Neisseria</i> .
Examination of phase-variable haemoglobin-haptoglobin binding proteins in non-typeable <i>Haemophilus influenzae</i> reveals a diverse distribution of multiple variants.

Global population structure and evolution of <i>Bordetella pertussis</i> and their relationship with vaccination.
Longitudinal TprK profiling of in vivo and in vitro-propagated <i>Treponema pallidum</i> subsp. <i>pallidum</i> reveals accumulation of antigenic variants in absence of immune pressure.
Nucleotide Sequencing of Antigen Genes of <i>Neisseria meningitidis</i> .
Host-pathogen interactions of nontypeable <i>Haemophilus influenzae</i> : from commensal to pathogen.
Analysis of Pilin Antigenic Variation in <i>Neisseria meningitidis</i> by Next-Generation Sequencing.
Bacterium of one thousand and one variants: genetic diversity of <i>Neisseria gonorrhoeae</i> pathogenicity.
Nontypeable <i>Haemophilus influenzae</i> : the role of N-acetyl-5-neuraminic acid in biology.
<i>Neisseria gonorrhoeae</i> host adaptation and pathogenesis.
Complete genome sequence of <i>Neisseria meningitidis</i> serogroup B strain MC58.
Antigenic variation of pilin regulates adhesion of <i>Neisseria meningitidis</i> to human epithelial cells.
<i>Neisseria meningitidis</i> : an overview of the carriage state.
Vaccine development against <i>Neisseria meningitidis</i> .
Expression of the Gene for Autotransporter AutB of <i>Neisseria meningitidis</i> Affects Biofilm Formation and Epithelial Transmigration.
Evaluation of the Protective Ability of the <i>Treponema pallidum</i> subsp. <i>pallidum</i> Tp0126 OmpW Homolog in the Rabbit Model of Syphilis.
Selective Inhibition of Sialic Acid-Based Molecular Mimicry in <i>Haemophilus influenzae</i> Abrogates Serum Resistance.
High-Frequency Changes in Pilin Glycosylation Patterns during <i>Neisseria meningitidis</i> Serogroup a Meningitis Outbreaks in the African Meningitis Belt.
Insights into the population structure and pan-genome of <i>Haemophilus influenzae</i> .
Full-length TprK of <i>Treponema pallidum</i> subsp. <i>pallidum</i> in lipid nanodiscs is a monomeric porin.
Polypeptides of <i>Treponema pallidum</i> : progress toward understanding their structural, functional, and immunologic roles. <i>Treponema Pallidum</i> Polypeptide Research Group.
Naturally occurring bactericidal antibodies specific for <i>Haemophilus influenzae</i> lipooligosaccharide are present in healthy adult individuals.
Mobile DNA in the Pathogenic <i>Neisseria</i> .
<i>Neisseria meningitidis</i> B vaccines.
Biological Functions of the Secretome of <i>Neisseria meningitidis</i> .
Estimation of Full-Length TprK Diversity in <i>Treponema pallidum</i> subsp. <i>pallidum</i> .
Variable expression of class 1 outer membrane protein in <i>Neisseria meningitidis</i> is caused by variation in the spacing between the -10 and -35 regions of the promoter.
Macrophage- <i>Neisseria gonorrhoeae</i> Interactions: A Better Understanding of Pathogen Mechanisms of Immunomodulation.
Antibody Binding and Complement-Mediated Killing of Invasive <i>Haemophilus influenzae</i> Isolates from Spain, Portugal, and the Netherlands.
A <i>Haemophilus influenzae</i> IgA protease-like protein promotes intimate interaction with human epithelial cells.
Biological basis for syphilis.
Phase variation and microevolution at homopolymeric tracts in <i>Bordetella pertussis</i> .
Structural Modeling of the <i>Treponema pallidum</i> Outer Membrane Protein Repertoire: a Road Map for Deconvolution of Syphilis Pathogenesis and Development of a Syphilis Vaccine.
The adhesins of non-typeable <i>Haemophilus influenzae</i> .

Potential of Phase Variation in Multiple Outer-Membrane Proteins During Spread of the Hyperinvasive <i>Neisseria meningitidis</i> Serogroup W ST-11 Lineage.
Structure-Activity Relationship of Fluorinated Sialic Acid Inhibitors for Bacterial Sialylation.
Comparative investigation of the genomic regions involved in antigenic variation of the TprK antigen among treponemal species, subspecies, and strains.
Epigenetic Regulation of Virulence and Immuno-evasion by Phase-Variable Restriction-Modification Systems in Bacterial Pathogens.
New concepts in immunity to <i>Neisseria gonorrhoeae</i> : innate responses and suppression of adaptive immunity favor the pathogen, not the host.
Evolution of the exclusively human pathogen <i>Neisseria gonorrhoeae</i> : Human-specific engagement of immunoregulatory Siglecs.
Naturally occurring lipid A mutants in <i>Neisseria meningitidis</i> from patients with invasive meningococcal disease are associated with reduced coagulopathy.
Sequence diversity of <i>Treponema pallidum</i> subsp. <i>pallidum</i> tprK in human syphilis lesions and rabbit-propagated isolates.
Genomic basis of a polyagglutinating isolate of <i>Neisseria meningitidis</i> .
<i>Neisseria gonorrhoeae</i> suppresses dendritic cell-induced, antigen-dependent CD4 T cell proliferation.
Phase variation of PorA, a major outer membrane protein, mediates escape of bactericidal antibodies by <i>Neisseria meningitidis</i> .
The minor pilin PilV provides a conserved adhesion site throughout the antigenically variable meningococcal type IV pilus.
Characterization of terminal NeuNAc $\alpha$ 2-3Gal $\beta$ 1-4GlcNAc sequence in lipooligosaccharides of <i>Neisseria meningitidis</i> .
Serum resistance and phase variation of a nasopharyngeal non-typeable <i>Haemophilus influenzae</i> isolate.
The virulence factors of <i>Bordetella pertussis</i> : talented modulators of host immune response.
Mechanisms of host manipulation by <i>Neisseria gonorrhoeae</i> .
The role of <i>B. pertussis</i> vaccine antigen gene variants in pertussis resurgence and possible consequences for vaccine development.
<i>Neisseria gonorrhoeae</i> induces a tolerogenic phenotype in macrophages to modulate host immunity.
Structural and kinetic characterizations of the polysialic acid O-acetyltransferase OatWY from <i>Neisseria meningitidis</i> .
Proteome analysis of <i>Bordetella pertussis</i> isolated from human macrophages.
DNA Blocks the Lethal Effect of Human Beta-Defensin 2 Against <i>Neisseria meningitidis</i> .
Genetic, Functional, and Immunogenic Analyses of the O-Linked Protein Glycosylation System in <i>Neisseria meningitidis</i> Serogroup A ST-7 Isolates.
Phasome analysis of pathogenic and commensal <i>Neisseria</i> species expands the known repertoire of phase variable genes, and highlights common adaptive strategies.
<i>Bordetella pertussis</i> : new concepts in pathogenesis and treatment.
Diversion of the immune response to <i>Neisseria gonorrhoeae</i> from Th17 to Th1/Th2 by treatment with anti-transforming growth factor $\beta$ antibody generates immunological memory and protective immunity.
Slam is an outer membrane protein that is required for the surface display of lipidated virulence factors in <i>Neisseria</i> .
Molecular mimicry of host structures by lipooligosaccharides of <i>Neisseria meningitidis</i> : characterization of sialylated and nonsialylated lacto-N-neotetraose (Gal $\beta$ 1-4GlcNAc $\beta$ 1-3Gal $\beta$ 1-4Glc) structures in lipooligosaccharides using monoclonal antibodies and specific lectins.

Mycoplasma genitalium adhesin P110 binds sialic-acid human receptors.
ArcA-regulated glycosyltransferase lic2B promotes complement evasion and pathogenesis of nontypeable Haemophilus influenzae.
Neisseria gonorrhoeae infection protects human endocervical epithelial cells from apoptosis via expression of host antiapoptotic proteins.
Absence of mucosal immunity in the human upper respiratory tract to the commensal bacteria Neisseria lactamica but not pathogenic Neisseria meningitidis during the peak age of nasopharyngeal carriage.
Regulation of capsule in Neisseria meningitidis.
Identification and Functional Characterization of Peptides With Antimicrobial Activity From the Syphilis Spirochete, Treponema pallidum.
Characterization of a novel antisense RNA in the major pilin locus of Neisseria meningitidis influencing antigenic variation.
Haemophilus influenzae in chronic bronchitis.
Neisserial surface lipoproteins: structure, function and biogenesis.
Reverse vaccinology approaches to introduce promising immunogenic and drug targets against antibiotic-resistant Neisseria gonorrhoeae: Thinking outside the box in current prevention and treatment.
Gonococcal Defenses against Antimicrobial Activities of Neutrophils.
The 47-kDa major lipoprotein immunogen of Treponema pallidum is a penicillin-binding protein with carboxypeptidase activity.
Genome-scale analysis of the non-cultivable Treponema pallidum reveals extensive within-patient genetic variation.
[PERSISTENCE OF BORDETELLA PERTUSSIS BACTERIA AND A POSSIBLE MECHANISM OF ITS FORMATION].
Regulatory T cells are locally induced during intravaginal infection of mice with Neisseria gonorrhoeae.
Investigation of the immune escape mechanism of Treponema pallidum.
Gene conversion: a mechanism for generation of heterogeneity in the tprK gene of Treponema pallidum during infection.
The Pulmonary Extracellular Matrix Is a Bactericidal Barrier Against Haemophilus influenzae in Chronic Obstructive Pulmonary Disease (COPD): Implications for an in vivo Innate Host Defense Function of Collagen VI.
Vaccine research for gonococcal infections: where are we?
Evidence for capsule switching between carried and disease-causing Neisseria meningitidis strains.
Treponema pallidum, the stealth pathogen, changes, but how?
The Haemophilus influenzae Sap transporter mediates bacterium-epithelial cell homeostasis.
Competition, coinfection and strain replacement in models of Bordetella pertussis.
Antigenic variation in Treponema pallidum: TprK sequence diversity accumulates in response to immune pressure during experimental syphilis.
Questions about gonococcal pilus phase- and antigenic variation.
Identification and Characterization of msf, a Novel Virulence Factor in Haemophilus influenzae.
The Pilin N-terminal Domain Maintains Neisseria gonorrhoeae Transformation Competence during Pilus Phase Variation.
The pathogenesis of disease due to nontypeable Haemophilus influenzae.
Blockade of the Adenylate Cyclase Toxin Synergizes with Opsonizing Antibodies to Protect Mice against Bordetella pertussis.

Nontypeable <i>Haemophilus influenzae</i> Has Evolved Preferential Use of N-Acetylneuraminic Acid as a Host Adaptation.
Nontypeable <i>Haemophilus influenzae</i> exploits the interaction between protein-E and vitronectin for the adherence and invasion to bronchial epithelial cells.
Function and protective capacity of <i>Treponema pallidum</i> subsp. <i>pallidum</i> glycerophosphodiester phosphodiesterase.
Relevance in biology and mechanisms of immune and treatment evasion of <i>Treponema pallidum</i> .
Importance of (antibody-dependent) complement-mediated serum killing in protection against <i>Bordetella pertussis</i> .
Temperature triggers immune evasion by <i>Neisseria meningitidis</i> .
Host-pathogen interaction during bacterial vaccination.
Sialic acid transport in <i>Haemophilus influenzae</i> is essential for lipopolysaccharide sialylation and serum resistance and is dependent on a novel tripartite ATP-independent periplasmic transporter.
Comparative genomics of the transportome of Ten <i>Treponema</i> species.
<i>Moraxella catarrhalis</i> evades neutrophil oxidative stress responses providing a safer niche for nontypeable <i>Haemophilus influenzae</i> .
Zinc piracy as a mechanism of <i>Neisseria meningitidis</i> for evasion of nutritional immunity.
<i>Neisseria gonorrhoeae</i> selectively suppresses the development of Th1 and Th2 cells, and enhances Th17 cell responses, through TGF- $\beta$ -dependent mechanisms.
<i>Neisseria meningitidis</i> IgA1-specific serine protease exhibits novel cleavage activity against IgG3.
<i>Haemophilus influenzae</i> resides in tonsils and uses immunoglobulin D binding as an evasion strategy.
Insights on persistent airway infection by non-typeable <i>Haemophilus influenzae</i> in chronic obstructive pulmonary disease.
Complete posttranslational modification mapping of pathogenic <i>Neisseria meningitidis</i> pilins requires top-down mass spectrometry.
'Nothing is permanent but change'- antigenic variation in persistent bacterial pathogens.
<i>Neisseria gonorrhoeae</i> -Induced Inflammatory Pyroptosis in Human Macrophages is Dependent on Intracellular Gonococci and Lipooligosaccharide.
Exploring the Impact of Ketodeoxynonulosonic Acid in Host-Pathogen Interactions Using Uptake and Surface Display by Nontypeable <i>Haemophilus influenzae</i> .
Triple co-culture and perfusion bioreactor for studying the interaction between <i>Neisseria gonorrhoeae</i> and neutrophils: A novel 3D tissue model for bacterial infection and immunity.
Serotype and serovar distribution of <i>Neisseria gonorrhoeae</i> isolated from high-risk populations in Bangladesh.
Investigating pertussis toxin and its impact on vaccination.
<i>Bordetella pertussis</i> evolution in the (functional) genomics era.
Bacterial periplasmic sialic acid-binding proteins exhibit a conserved binding site.
Evasion of killing by human antibody and complement through multiple variations in the surface oligosaccharide of <i>Haemophilus influenzae</i> .
The impact of protein-conjugate polysaccharide vaccines: an endgame for meningitis?
Meningococcal vaccines.
Phase variation and host immunity against high molecular weight (HMW) adhesins shape population dynamics of nontypeable <i>Haemophilus influenzae</i> within human hosts.
Molecular determinants of the pathogenesis of disease due to non-typable <i>Haemophilus influenzae</i> .
Antibody to Rmp (outer membrane protein 3) increases susceptibility to gonococcal infection.
RecQ DNA helicase HRDC domains are critical determinants in <i>Neisseria gonorrhoeae</i> pilin antigenic variation and DNA repair.

Notes on syphilis vaccine development.
Pertussis: Identification, Prevention and Control.
The mimicry of human glycolipids and glycosphingolipids by the lipooligosaccharides of pathogenic neisseria and haemophilus.
Molecular analysis of the meningococcal LPS expression.
Two cases of type-a Haemophilus influenzae meningitis within the same week in the same hospital are phylogenetically unrelated but recently exchanged capsule genes.
Antibody responses elicited against the Treponema pallidum repeat proteins differ during infection with different isolates of Treponema pallidum subsp. pallidum.
Therapeutic CMP-Nonulosonates against Multidrug-Resistant Neisseria gonorrhoeae.
The obligate human pathogen, Neisseria gonorrhoeae, is polyploid.
Modeling Immune Evasion and Vaccine Limitations by Targeted Nasopharyngeal Bordetella pertussis Inoculation in Mice.
The minimal cellular genome of mycoplasma.
Global phylogeny of Treponema pallidum lineages reveals recent expansion and spread of contemporary syphilis.
Role of Evolutionary Selection Acting on Vaccine Antigens in the Re-Emergence of Bordetella Pertussis.
Bordetella pertussis entry into respiratory epithelial cells and intracellular survival.
Are lactoferrin receptors in Gram-negative bacteria viable vaccine targets?
Comparative genomics of mycoplasmas.
Neisseria gonorrhoeae arthritis in a patient with systemic lupus: resistance and virulence profiles.
Non-typeable Haemophilus influenzae airways infection: the next treatable trait in asthma?
Sialic acid acquisition in bacteria-one substrate, many transporters.
Genome-scale approaches to identify genes essential for Haemophilus influenzae pathogenesis.
Antimicrobial peptide resistance in Neisseria meningitidis.
Small-Molecule Inhibitors of Haemophilus influenzae IgA1 Protease.
Opa proteins and CEACAMs: pathways of immune engagement for pathogenic Neisseria.
Genetic Profiling of the Full-Length tprK Gene in Patients with Primary and Secondary Syphilis.
Characterization of a novel porin protein from Moraxella catarrhalis and identification of an immunodominant surface loop.
The role of peptide loops of the Bordetella pertussis protein P.69 pertactin in antibody recognition.
Emergence of a new epidemic Neisseria meningitidis serogroup A Clone in the African meningitis belt: high-resolution picture of genomic changes that mediate immune evasion.
The search for Brachyspira outer membrane proteins that interact with the host.
The tprK gene is heterogeneous among Treponema pallidum strains and has multiple alleles.
Antigenic diversity of Haemophilus somnus lipooligosaccharide: phase-variable accessibility of the phosphorylcholine epitope.
Insights into the genetic variation profile of tprK in Treponema pallidum during the development of natural human syphilis infection.
How bacteria hack the matrix and dodge the bullets of immunity.
Protection against syphilis correlates with specificity of antibodies to the variable regions of Treponema pallidum repeat protein K.
Neisseria meningitidis Type IV Pili Composed of Sequence Invariable Pilins Are Masked by Multisite Glycosylation.

Human milk lactoferrin inactivates two putative colonization factors expressed by <i>Haemophilus influenzae</i> .
Effect of FHA and Prn on <i>Bordetella pertussis</i> colonization of mice is dependent on vaccine type and anatomical site.
Transcription of a cis-acting, noncoding, small RNA is required for pilin antigenic variation in <i>Neisseria gonorrhoeae</i> .
<i>Treponema pallidum</i> , the syphilis spirochete: making a living as a stealth pathogen.
Colonization of dermal arterioles by <i>Neisseria meningitidis</i> provides a safe haven from neutrophils.
Structure of the N-terminal domain of human CEACAM1: binding target of the opacity proteins during invasion of <i>Neisseria meningitidis</i> and <i>N. gonorrhoeae</i> .
Structural Basis for Recombinatorial Permissiveness in the Generation of <i>Anaplasma marginale</i> Msp2 Antigenic Variants.
Colonization of healthy children by <i>Moraxella catarrhalis</i> is characterized by genotype heterogeneity, virulence gene diversity and co-colonization with <i>Haemophilus influenzae</i> .
Sequence variation and immunogenicity of the <i>Mycoplasma genitalium</i> MgpB and MgpC adherence proteins during persistent infection of men with non-gonococcal urethritis.
Pertussis in Poland.
Lower airway colonization and inflammatory response in COPD: a focus on <i>Haemophilus influenzae</i> .
Localized frameshift mutation generates selective, high-frequency phase variation of a surface lipoprotein encoded by a mycoplasma ABC transporter operon.
Non-typeable <i>Haemophilus influenzae</i> chronic colonization in chronic obstructive pulmonary disease (COPD).
Emergence of pertactin-deficient pertussis strains in Australia can be explained by models of vaccine escape.
[Immunological mechanisms of <i>Neisseria gonorrhoeae</i> infection: An update].
<i>Mycoplasma genitalium</i> Nonadherent Phase Variants Arise by Multiple Mechanisms and Escape Antibody-Dependent Growth Inhibition.
Influence of serogroup B meningococcal vaccine antigens on growth and survival of the meningococcus in vitro and in ex vivo and in vivo models of infection.
PRELP Enhances Host Innate Immunity against the Respiratory Tract Pathogen <i>Moraxella catarrhalis</i> .
Molecular biology and pathogenicity of mycoplasmas.
Role of Vpma phase variation in <i>Mycoplasma agalactiae</i> pathogenesis.
The Pathobiology of <i>Neisseria gonorrhoeae</i> Lower Female Genital Tract Infection.
Circumvention of herd immunity during an outbreak of meningococcal disease could be correlated to escape mutation in the <i>porA</i> gene of <i>Neisseria meningitidis</i> .
Endotoxin, capsule, and bacterial attachment contribute to <i>Neisseria meningitidis</i> resistance to the human antimicrobial peptide LL-37.
<i>Bordetella pertussis</i> autotransporter Vag8 binds human C1 esterase inhibitor and confers serum resistance.
Molecular variation among type IV pilin ( <i>bfpA</i> ) genes from diverse enteropathogenic <i>Escherichia coli</i> strains.
Upregulation of ATF3 inhibits expression of the pro-inflammatory cytokine IL-6 during <i>Neisseria gonorrhoeae</i> infection.
Bacterial Sialic Acid Catabolism at the Host-Microbe Interface.
<i>Haemophilus influenzae</i> protein F mediates binding to laminin and human pulmonary epithelial cells.
<i>Bordetella bronchiseptica</i> and <i>Bordetella pertussis</i> : Similarities and Differences in Infection, Immuno-Modulation, and Vaccine Considerations.



The OmpA family of proteins: roles in bacterial pathogenesis and immunity.
Genetics and molecular specificity of sialylation of <i>Haemophilus somni</i> lipooligosaccharide (LOS) and the effect of LOS sialylation on Toll-like receptor-4 signaling.
Biology of the <i>Gonococcus</i> : Disease and Pathogenesis.
Beware of <i>Mycoplasma</i> Anti-immunoglobulin Strategies.
Bioinformatic analysis of meningococcal Msf and Opc to inform vaccine antigen design.
Structure and mechanism of a molecular rheostat, an RNA thermometer that modulates immune evasion by <i>Neisseria meningitidis</i> .
<i>Haemophilus influenzae</i> Protein D antibody suppression in a multi-component vaccine formulation.
Antigenic variation of <i>Anaplasma marginale</i> by expression of MSP2 mosaics.
Identification and characterization of important residues in the catalytic mechanism of CMP-Neu5Ac synthetase from <i>Neisseria meningitidis</i> .
Virulence Associated Gene 8 of <i>Bordetella pertussis</i> Enhances Contact System Activity by Inhibiting the Regulatory Function of Complement Regulator C1 Inhibitor.
Identification of a Large Family of Slam-Dependent Surface Lipoproteins in Gram-Negative Bacteria.
Inhibition of <i>Neisseria meningitidis</i> sialic acid synthase by a tetrahedral intermediate analogue.
Biological significance of IgA1 proteases in bacterial colonization and pathogenesis: critical evaluation of experimental evidence.
Meningococcal interactions with the host.
Heterogeneity in non-epitope loop sequence and outer membrane protein complexes alters antibody binding to the major porin protein PorB in serogroup B <i>Neisseria meningitidis</i> .
Molecular Basis of Ligand-Dependent Regulation of NadR, the Transcriptional Repressor of Meningococcal Virulence Factor NadA.
Impaired alveolar macrophage response to <i>Haemophilus</i> antigens in chronic obstructive lung disease.
Map-based comparative genomic analysis of virulent <i>Haemophilus parasuis</i> serovars 4 and 5.
Simultaneous variation of the immunodominant outer membrane proteins, MSP2 and MSP3, during <i>Anaplasma marginale</i> persistence in vivo.
<i>Haemophilus influenzae</i> acquires vitronectin via the ubiquitous Protein F to subvert host innate immunity.
In Silico Designed Multi-Epitope Immunogen "Tpme-VAC/LGCM-2022" May Induce Both Cellular and Humoral Immunity against <i>Treponema pallidum</i> Infection.
Age-related genotypic and phenotypic differences in <i>Moraxella catarrhalis</i> isolates from children and adults presenting with respiratory disease in 2001-2002.
Phenotypic variation of <i>Mycoplasma iowae</i> surface antigen.
<i>Haemophilus parasuis</i> $\alpha$ -2,3-sialyltransferase-mediated lipooligosaccharide sialylation contributes to bacterial pathogenicity.
The Unique Microbiology and Molecular Pathogenesis of <i>Mycoplasma genitalium</i> .
<i>Moraxella catarrhalis</i> Evades Host Innate Immunity via Targeting Cartilage Oligomeric Matrix Protein.
Phosphorylcholine allows for evasion of bactericidal antibody by <i>Haemophilus influenzae</i> .
Triggering Closure of a Sialic Acid TRAP Transporter Substrate Binding Protein through Binding of Natural or Artificial Substrates.
Antigenic variation of TprK facilitates development of secondary syphilis.
Development of Sialic Acid-coated Nanoparticles for Targeting Cancer and Efficient Evasion of the Immune System.
Heme utilization by nontypeable <i>Haemophilus influenzae</i> is essential and dependent on Sap transporter function.

RecA-binding pilE G4 sequence essential for pilin antigenic variation forms monomeric and 5' end-stacked dimeric parallel G-quadruplexes.
mgpB and mgpC sequence diversity in Mycoplasma genitalium is generated by segmental reciprocal recombination with repetitive chromosomal sequences.
Genomic analysis of the meningococcal ST-4821 complex-Western clade, potential sexual transmission and predicted antibiotic susceptibility and vaccine coverage.
CEACAM1 inhibits Toll-like receptor 2-triggered antibacterial responses of human pulmonary epithelial cells.
Capsule switching of Neisseria meningitidis sequence type 7 serogroup A to serogroup X.
The Capricious Nature of Bacterial Pathogens: Phasevarions and Vaccine Development.
A biphasic epigenetic switch controls immunoevasion, virulence and niche adaptation in non-typeable Haemophilus influenzae.
Identification of a phage-encoded Ig-binding protein from invasive Neisseria meningitidis.
Exploring the Ability of Meningococcal Vaccines to Elicit Mucosal Immunity: Insights from Humans and Mice.
Pertactin-negative Bordetella pertussis strains: evidence for a possible selective advantage.
Frequent capsule switching in 'ultra-virulent' meningococci - Are we ready for a serogroup B ST-11 complex outbreak?
Antigenic and/or phase variation of PorA protein in non-subtypable Neisseria meningitidis strains isolated in Spain.
Host cell invasion by pathogenic Neisseriae.
The immunization-induced antibody response to the Anaplasma marginale major surface protein 2 and its association with protective immunity.
Loss of both Holliday junction processing pathways is synthetically lethal in the presence of gonococcal pilin antigenic variation.
Phase-variable bacterial loci: how bacteria gamble to maximise fitness in changing environments.
Phasevarions of bacterial pathogens - phase-variable epigenetic regulators evolving from restriction-modification systems.
Genetic diversity of the 28-kilodalton outer membrane protein gene in human isolates of Ehrlichia chaffeensis.
Physiologic cold shock of Moraxella catarrhalis affects the expression of genes involved in the iron acquisition, serum resistance and immune evasion.
Substrate-bound outward-open structure of a Na(+)-coupled sialic acid symporter reveals a new Na(+) site.
Persistence, immune response, and antigenic variation of Mycoplasma genitalium in an experimentally infected pig-tailed macaque (Macaca nemestrina).
A multi-enzyme machine polymerizes the Haemophilus influenzae type b capsule.
Proteins of Bartonella bacilliformis: Candidates for Vaccine Development.
Suppression of Alternative Lipooligosaccharide Glycosyltransferase Activity by UDP-Galactose Epimerase Enhances Murine Lung Infection and Evasion of Serum IgM.
Emergence and genomic diversification of a virulent serogroup W:ST-2881(CC175) Neisseria meningitidis clone in the African meningitis belt.
Primary Structural Variation in Anaplasma marginale Msp2 Efficiently Generates Immune Escape Variants.
Structural and Biosynthetic Diversity of Nonulosonic Acids (NulOs) That Decorate Surface Structures in Bacteria.

MBOVPG45_0375 Encodes an IgG-Binding Protein and MBOVPG45_0376 Encodes an IgG-Cleaving Protein in <i>Mycoplasma bovis</i> .
Ultrahigh resolution and full-length pilin structures with insights for filament assembly, pathogenic functions, and vaccine potential.
Update on bacterial pathogenesis in BRD.
MIB-MIP is a mycoplasma system that captures and cleaves immunoglobulin G.
Multiple gonococcal pilin antigenic variants are produced during experimental human infections.
Complete Genome Sequencing of <i>Leptospira interrogans</i> Isolates from Malaysia Reveals Massive Genome Rearrangement but High Conservation of Virulence-Associated Genes.
Expression of sialic acids and other nonulosonic acids in <i>Leptospira</i> .
Crystal structure of the retaining galactosyltransferase LgtC from <i>Neisseria meningitidis</i> in complex with donor and acceptor sugar analogs.
The Sialoglycan Binding Adhesins of <i>Mycoplasma genitalium</i> and <i>Mycoplasma pneumoniae</i> .
The Role of Lipoproteins in Mycoplasma-Mediated Immunomodulation.
Structural insight into the dual function of LbpB in mediating <i>Neisserial</i> pathogenesis.
Role of <i>Bordetella</i> O antigen in respiratory tract infection.
Segmental Variation in a Duplicated msp2 Pseudogene Generates <i>Anaplasma marginale</i> Antigenic Variants.
Characterization of <i>Treponema denticola</i> Major Surface Protein (Msp) by Deletion Analysis and Advanced Molecular Modeling.
Antigenic variation of <i>Anaplasma marginale</i> msp2 occurs by combinatorial gene conversion.
Long-Read Sequencing Reveals Genetic Adaptation of <i>Bartonella</i> Adhesin A Among Different <i>Bartonella henselae</i> Isolates.
A plate-based high-throughput activity assay for polysialyltransferase from <i>Neisseria meningitidis</i> .
Generation of antigenic variants via gene conversion: Evidence for recombination fitness selection at the locus level in <i>Anaplasma marginale</i> .
Experimental Infection of Pig-Tailed Macaques ( <i>Macaca nemestrina</i> ) with <i>Mycoplasma genitalium</i> .
Intrastrain heterogeneity of the mgpB gene in <i>Mycoplasma genitalium</i> is extensive in vitro and in vivo and suggests that variation is generated via recombination with repetitive chromosomal sequences.
Crystallization and preliminary X-ray diffraction analysis of antigen-binding fragments which are specific for antigenic conformations of sialic acid homopolymers.
The Use of the Antigenically Variable Major Surface Protein 2 in the Establishment of Superinfection during Natural Tick Transmission of <i>Anaplasma marginale</i> in Southern Ghana.
Expression patterns of <i>Anaplasma marginale</i> Msp2 variants change in response to growth in cattle, and tick cells versus mammalian cells.
Phase variations of the <i>Mycoplasma penetrans</i> main surface lipoprotein increase antigenic diversity.
A novel approach to decrease sialic acid expression in cells by a C-3-modified N-acetylmannosamine.
Phenotypic and genotypic analyses of <i>Neisseria gonorrhoeae</i> isolates that express frequently recovered PorB PIA variable region types suggest that certain P1a porin sequences confer a selective advantage for urogenital tract infection.
Erythroid Suppressor Cells Compromise Neonatal Immune Response against <i>Bordetella pertussis</i> .
Protocols to Interrogate the Interactions Between <i>Neisseria gonorrhoeae</i> and Primary Human Neutrophils.
In vitro resistance mechanisms of <i>Neisseria meningitidis</i> against neutrophil extracellular traps.
Epitope clusters in the major outer membrane protein of <i>Chlamydia trachomatis</i> .
Ascending Reproductive Tract Infection in Pig-Tailed Macaques Inoculated with <i>Mycoplasma genitalium</i> .

In silico structural homology modeling and functional characterization of <i>Mycoplasma gallisepticum</i> variable lipoprotein hemagglutinin proteins.
Selection for functional diversity drives accumulation of point mutations in Dr adhesins of <i>Escherichia coli</i> .
Adhesive activity of the <i>Haemophilus</i> cryptic genospecies <i>cha</i> autotransporter is modulated by variation in tandem Peptide repeats.
Predominant Single Stable VpmaV Expression in Strain GM139 and Major Differences with <i>Mycoplasma agalactiae</i> Type Strain PG2.
Molecular aspects of <i>Moraxella catarrhalis</i> pathogenesis.
Variability of trinucleotide tandem repeats in the MgPa operon and its repetitive chromosomal elements in <i>Mycoplasma genitalium</i> .
Distinctly different msp2 pseudogene repertoires in <i>Anaplasma marginale</i> strains that are capable of superinfection.
Kinetics of Genetic Variation of the <i>Mycoplasma genitalium</i> MG192 Gene in Experimentally Infected Chimpanzees.
Extensive variation and rapid shift of the MG192 sequence in <i>Mycoplasma genitalium</i> strains from patients with chronic infection.
Bacterial Outer Membrane Polysaccharide Export (OPX) Proteins Occupy Three Structural Classes with Selective $\beta$ -Barrel Porin Requirements for Polymer Secretion.
Probing the determinants of phosphorylated sugar-substrate binding for human sialic acid synthase.
A novel non-homologous recombination-mediated mechanism for <i>Escherichia coli</i> unilateral flagellar phase variation.
Conformational comparisons of <i>Pasteurella multocida</i> types B and E and structurally related capsular polysaccharides.
<i>Moraxella catarrhalis</i> : from interactions with the host immune system to vaccine development.
<i>Neisseria gonorrhoeae</i> elicits extracellular traps in primary neutrophil culture while suppressing the oxidative burst.
Impact of sequence diversity in the <i>Moraxella catarrhalis</i> UspA2/UspA2H head domain on vitronectin binding and antigenic variation.
Host cell interactions of novel antigenic membrane proteins of <i>Mycoplasma agalactiae</i> .
Antigenic Variation in <i>Neisseria gonorrhoeae</i> Occurs Independently of RecQ-Mediated Unwinding of the pilE G Quadruplex.
The lipooligosaccharide-modifying enzyme LptA enhances gonococcal defence against human neutrophils.
Versatile substrates and probes for IgA1 protease activity.
Contribution of a Novel Pertussis Toxin-Like Factor in Mediating Persistent Otitis Media.
Molecular Basis for <i>Bordetella pertussis</i> Interference with Complement, Coagulation, Fibrinolytic, and Contact Activation Systems: the Cryo-EM Structure of the Vag8-C1 Inhibitor Complex.
Recombination in the ompA gene but not the omcB gene of <i>Chlamydia</i> contributes to serovar-specific differences in tissue tropism, immune surveillance, and persistence of the organism.
Reconstitution of surface lipoprotein translocation through the Slam translocon.
Analysis of potential virulence genes and competence to transformation in <i>Haemophilus influenzae</i> biotype <i>aegyptius</i> associated with Brazilian Purpuric Fever.
Variability in the 28-kDa surface antigen protein multigene locus of isolates of the emerging disease agent <i>Ehrlichia chaffeensis</i> suggests that it plays a role in immune evasion.
Global Changes in <i>Mycoplasma gallisepticum</i> Phase-Variable Lipoprotein Gene vlhA Expression during In Vivo Infection of the Natural Chicken Host.

In silico analysis of Pap31 from <i>Bartonella bacilliformis</i> and other <i>Bartonella</i> spp.
Processing-independent CRISPR RNAs limit natural transformation in <i>Neisseria meningitidis</i> .
A Phase-Variable Surface Layer from the Gut Symbiont <i>Bacteroides thetaiotaomicron</i> .
The structure of an archaeal pilus.
Crystal Structure of a Complex of Surfactant Protein D (SP-D) and <i>Haemophilus influenzae</i> Lipopolysaccharide Reveals Shielding of Core Structures in SP-D-Resistant Strains.
Structure and function relationships in IgA.
Identification and characterization of a <i>Mycoplasma hyopneumoniae</i> adhesin.
Ureaplasma antigenic variation beyond MBA phase variation: DNA inversions generating chimeric structures and switching in expression of the MBA N-terminal paralogue UU172.
Efficacy of Antigonococcal CMP-Nonulosonate Therapeutics Require Cathelicidins.
Bacterial lyso-form lipoproteins are synthesized via an intramolecular acyl chain migration.
Defining the remarkable structural malleability of a bacterial surface protein Rib domain implicated in infection.
<i>Histophilus somni</i> : Antigenic and Genomic Changes Relevant to Bovine Respiratory Disease.
Phase variation of the gonococcal siderophore receptor FetA.
NeuNAc Oxime: A Slow-Binding and Effectively Irreversible Inhibitor of the Sialic Acid Synthase NeuB.
Variable Lipoprotein Hemagglutinin A Gene (vlhA) Expression in Variant <i>Mycoplasma gallisepticum</i> Strains In Vivo.
Comparative Analysis of <i>Mycoplasma gallisepticum</i> vlhA Promoters.
The <i>Mycoplasma hominis</i> P120 membrane protein contains a 216 amino acid hypervariable domain that is recognized by the human humoral immune response.
<i>Glaesserella parasuis</i> autotransporters EspP1 and EspP2 are novel IgA-specific proteases.
A novel group of <i>Moraxella catarrhalis</i> UspA proteins mediates cellular adhesion via CEACAMs and vitronectin.
Recombinational switching of the <i>Clostridium difficile</i> S-layer and a novel glycosylation gene cluster revealed by large-scale whole-genome sequencing.
Differential expression and glycosylation of anaplasma phagocytophilum major surface protein 2 paralogs during cultivation in sialyl Lewis x-deficient host cells.
The influence of variable-heavy chain families on IgG(2), (3), (4), FcγRs and B-cell superantigens protein G and L binding using biolayer interferometry.
Structural insights into the molecular organization of the S-layer from <i>Clostridium difficile</i> .
Identification of the <i>Bartonella</i> autotransporter CFA as a protective antigen and hypervariable target of neutralizing antibodies in mice.
<i>Bacillus anthracis</i> Diversity and Geographic Potential across Nigeria, Cameroon and Chad: Further Support of a Novel West African Lineage.
Autophagy Is Associated with Pathogenesis of <i>Haemophilus parasuis</i> .
UV light-induced spatial loss of sialic acid capping using a photoactivatable sialyltransferase inhibitor.
Restricted changes in major surface protein-2 (msp2) transcription after prolonged in vitro passage of <i>Anaplasma phagocytophilum</i> .
Nucleotide and phylogenetic analyses of the <i>Chlamydia trachomatis</i> ompA gene indicates it is a hotspot for mutation.
Role of sialic acid-containing glycans of matrix metalloproteinase-9 (MMP-9) in the interaction between MMP-9 and staphylococcal superantigen-like protein 5.
Molecular characterization of Msp2/P44 of <i>Anaplasma phagocytophilum</i> isolated from infected patients and <i>Haemaphysalis longicornis</i> in Laizhou Bay, Shandong Province, China.

Neurosyphilis with ocular involvement and normal magnetic resonance imaging results affirmed by metagenomic next-generation sequencing.
The central region of the msp gene of <i>Treponema denticola</i> has sequence heterogeneity among clinical samples, obtained from patients with periodontitis.
Siglecs family used by pathogens for immune escape may engaged in immune tolerance in pregnancy.
Fowl Adenovirus (FAdV) Recombination with Intertypic Crossovers in Genomes of FAdV-D and FAdV-E, Displaying Hybrid Serological Phenotypes.
Reflectron MALDI TOF and MALDI TOF/TOF mass spectrometry reveal novel structural details of native lipooligosaccharides.
Lateral opening and exit pore formation are required for BamA function.
<i>Haemophilus parasuis</i> infection in 3D4/21 cells induces autophagy through the AMPK pathway.
Xer1-independent mechanisms of Vpma phase variation in <i>Mycoplasma agalactiae</i> are triggered by Vpma-specific antibodies.
Designing of novel chimeric PvpA-pMGA protein of <i>Mycoplasma gallisepticum</i> , applicable for indirect ELISA.
IgA: Structure, Function, and Developability.
A Phylogeographic Analysis of Porcine Parvovirus 1 in Africa.
Nucleotide polymorphism assay for the identification of west African group <i>Bacillus anthracis</i> : a lineage lacking anthrose.
A comprehensive screen for chicken proteins that interact with proteins unique to virulent strains of Marek's disease virus.
Host Immune Response to <i>Histophilus somni</i> .
Phylogenetic analysis of three orf virus strains isolated from different districts in Shandong Province, East China.
<i>Bartonella</i> spp. in human and animal populations in Gauteng, South Africa, from 2007 to 2009.
Molecular Mechanisms of <i>Bartonella</i> and Mammalian Erythrocyte Interactions: A Review.
Binding of bovine IgG2a and IgG2b allotypes to protein A, protein G, and <i>Haemophilus somnus</i> IgBPs.
Evidence of possible evasion of protective immunity by NAD-independent isolates of <i>Haemophilus paragallinarum</i> in poultry.
Serum antibody titers to <i>Bacteroides forsythus</i> in elderly subjects with gingivitis or periodontitis.
Structural insights into substrate and inhibitor binding sites in human indoleamine 2,3-dioxygenase 1.
Target cell range of <i>Haemophilus ducreyi</i> hemolysin and its involvement in invasion of human epithelial cells.
Generating tertiary protein structures via interpretable graph variational autoencoders.
Similar proportions of immunoglobulin A1 (IgA1) protease-producing streptococci in initial dental plaque of selectively IgA-deficient and normal individuals.
Proteomic analysis of proteins released by <i>Tritrichomonas foetus</i> : Identification of potential targets for the development of new diagnostic methods.
Emergence of a novel immune-evasion strategy from an ancestral protein fold in bacteriophage Mu.
Homology modeling, virtual screening and dynamics study of proteins involved in Pebrine - Serine protease inhibitor 106 and spore wall protein 26.
Optimization of protein-protein docking for predicting Fc-protein interactions.
The critical role of residues 43R and 44Q of carcinoembryonic antigen cell adhesion molecules-1 in the protection from killing by human NK cells.
Staphylococcal superantigen-like protein 5 inhibits matrix metalloproteinase 9 from human neutrophils.

Identification and partial characterization of an immunoglobulin A protease associated with human sperm.
Protease activity of <i>Blastocystis hominis</i> subtype3 in symptomatic and asymptomatic patients.
The LMW surface-layer proteins of <i>Clostridium difficile</i> PCR ribotypes 027 and 001 share common immunogenic properties.
Measurement of Protein-Protein Interactions through Microscale Thermophoresis (MST).
Functional analysis of the CXXC motif using phage antibodies that cross-react with protein disulphide-isomerase family proteins.
<i>Necator americanus</i> Ancylostoma Secreted Protein-2 (Na-ASP-2) Binds an Ascaroside (ascr#3) in Its Fatty Acid Binding Site.
Synthesis of lactosamine-based building blocks on a practical scale and investigations of their assembly for the preparation of (19)F-labelled LacNAc oligomers.
Conformation of the branched O-specific polysaccharide of <i>Shigella dysenteriae</i> type 2: molecular mechanics calculations show a compact helical structure exposing an epitope which potentially mimics galabiose.
Structural basis for Fc gammaRIIIa recognition of human IgG and formation of inflammatory signaling complexes.
Subtilase cytotoxin cleaves newly synthesized BiP and blocks antibody secretion in B lymphocytes.
Complete assignment of Ala, Ile, Leu(ProS), Met and Val(ProS) methyl groups of the protruding domain from human norovirus GII.4 Saga.
The Role of IgG4 in the Fine Tuning of Tolerance in IgE-Mediated Allergy and Cancer.
Bulged and Canonical G-Quadruplex Conformations Determine NDPK Binding Specificity.
Characterization of PEGylated Asparaginase: New Opportunities from NMR Analysis of Large PEGylated Therapeutics.
[Specific coalescent peptide of CD59 screened by phage library].
Enhanced biocompatibility of CD47-functionalized vascular stents.
Purification and characterization of a 33 kDa serine protease from <i>Acanthamoeba lugdunensis</i> KA/E2 isolated from a Korean keratitis patient.
Prostate secretory protein 94 inhibits sterol binding and export by the mammalian CAP protein CRISP2 in a calcium-sensitive manner.
A procedure for producing an anti-AXL nanobody in <i>E. coli</i> .
A Sensitive and Reversible Labeling Strategy Enables Global Mapping of the Core-Fucosylated Glycoproteome on Cell Surfaces.
Desmoplakin and KIF20B as target antigens in patients with paroxysmal nocturnal haemoglobinuria.
Crystal structures of HER3 extracellular domain 4 in complex with the designed ankyrin-repeat protein D5.
Mirolysin structures open a window on gum disease.
Mutations for decreasing the immunogenicity and maintaining the function of core streptavidin.
Structural Basis for CD96 Immune Receptor Recognition of Nectin-like Protein-5, CD155.
Small ruminant lentivirus infection influences expression of acute phase proteins and cathelicidin genes in milk somatic cells and peripheral blood leukocytes of dairy goats.

Table A2-59, Cluster 58

Cluster 58 focuses on T cells (especially CD8+), emphasizing the role of T cell response suppression (in the tumor microenvironment) in immune system evasion (1478)
--

Transfer of Allogeneic CD4+ T Cells Rescues CD8+ T Cells in Anti-PD-L1-Resistant Tumors Leading to Tumor Eradication.
PD1-CD28 Fusion Protein Enables CD4+ T Cell Help for Adoptive T Cell Therapy in Models of Pancreatic Cancer and Non-hodgkin Lymphoma.
[Cytotoxic T lymphocytes: role in immunosurveillance and in immunotherapy].
Viral immune evasion due to persistence of activated T cells without effector function.
CD8(+) cytotoxic T lymphocytes in cancer immunotherapy: A review.
Visualizing the course of antigen-specific CD8 and CD4 T cell responses to a growing tumor.
PD-1 does not mark tumor-infiltrating CD8+ T cell dysfunction in human gastric cancer.
Why Do CD8+ T Cells become Indifferent to Tumors: A Dynamic Modeling Approach.
A Human Trypanosome Suppresses CD8+ T Cell Priming by Dendritic Cells through the Induction of Immune Regulatory CD4+ Foxp3+ T Cells.
Intra-tumoral production of IL18, but not IL12, by TCR-engineered T cells is non-toxic and counteracts immune evasion of solid tumors.
Mechanism of T cell tolerance induced by myeloid-derived suppressor cells.
Dual control of antitumor CD8 T cells through the programmed death-1/programmed death-ligand 1 pathway and immunosuppressive CD4 T cells: regulation and counterregulation.
Up-regulation of PD-L1, IDO, and T(regs) in the melanoma tumor microenvironment is driven by CD8(+) T cells.
Dendritic Cells and CD8 T Cell Immunity in Tumor Microenvironment.
CD8(+) T-cell-Mediated Immunoediting Influences Genomic Evolution and Immune Evasion in Murine Gliomas.
Tumor progression despite massive influx of activated CD8(+) T cells in a patient with malignant melanoma ascites.
Suppression of T cell responses in the tumor microenvironment.
Tumor-derived chemokine CCL5 enhances TGF- $\beta$ -mediated killing of CD8(+) T cells in colon cancer by T-regulatory cells.
PLA2G2A(+) cancer-associated fibroblasts mediate pancreatic cancer immune escape via impeding antitumor immune response of CD8(+) cytotoxic T cells.
Chimeric NKG2D expressing T cells eliminate immunosuppression and activate immunity within the ovarian tumor microenvironment.
Increased PD-1 expression on CD4+ and CD8+ T cells is involved in immune evasion in gastric cancer.
An intra-tumoral niche maintains and differentiates stem-like CD8 T cells.
Basophils Promote Tumor Rejection via Chemotaxis and Infiltration of CD8+ T Cells.
Turning on/off tumor-specific CTL response during progressive tumor growth.
Metabolic Regulation of CD8(+) T Cells: From Mechanism to Therapy.
Enhancement of adoptive T cell transfer with single low dose pretreatment of doxorubicin or paclitaxel in mice.
The elevated expression of LAG-3 on CD8+T cells correlates with disease severity of pulmonary TB.
HMGB2 regulates the differentiation and stemness of exhausted CD8(+) T cells during chronic viral infection and cancer.
SMAR1 promotes immune escape of Tri-negative Breast Cancer through a mechanism involving T-bet/PD-1 Axis.
Intratumoral CXCL13(+)CD8(+)T cell infiltration determines poor clinical outcomes and immunoevasive contexture in patients with clear cell renal cell carcinoma.
Trypanosoma cruzi-specific CD8(+) T cells and other immunological hallmarks in chronic Chagas cardiomyopathy: Two decades of research.



Selection of Tumor-Specific Cytotoxic T Lymphocytes in Acute Myeloid Leukemia Patients Through the Identification of T-Cells Capable to Establish Stable Interactions With the Leukemic Cells: "Doublet Technology".
CD8+ T cell immunity against a tumor/self-antigen is augmented by CD4+ T helper cells and hindered by naturally occurring T regulatory cells.
Generation of CD8(+) T cells expressing two additional T-cell receptors (TETARs) for personalised melanoma therapy.
Selectively impaired CD8+ but not CD4+ T cell cycle arrest during priming as a consequence of dendritic cell interaction with plasmodium-infected red cells.
Reinforcement of cancer immunotherapy by adoptive transfer of cblb-deficient CD8+ T cells combined with a DC vaccine.
Blocking Tumor Necrosis Factor $\alpha$ Enhances CD8 T-cell-Dependent Immunity in Experimental Melanoma.
Mixed lineage kinase 3 inhibition induces T cell activation and cytotoxicity.
Long-term staphylococcal enterotoxin C1 exposure induces soluble factor-mediated immunosuppression by bovine CD4+ and CD8+ T cells.
Transcription factor Batf3 is important for development of CD8+ T-cell response against a phagosomal bacterium regardless of the location of antigen.
PD-1 blockade restores helper activity of tumor-infiltrating, exhausted PD-1hiCD39+ CD4 T cells.
Platycodon grandiflorum Triggers Antitumor Immunity by Restricting PD-1 Expression of CD8(+) T Cells in Local Tumor Microenvironment.
Cytotoxicity of tumor antigen specific human T cells is unimpaired by arginine depletion.
LSECtin expressed on melanoma cells promotes tumor progression by inhibiting antitumor T-cell responses.
Distinct mechanisms of CD4+ and CD8+ T-cell activation and bystander apoptosis induced by human immunodeficiency virus type 1 virions.
T cells in multiple myeloma display features of exhaustion and senescence at the tumor site.
Inhibition of host Ogr1 enhances effector CD8(+) T-cell function by modulating acidic microenvironment.
Re-adapting T cells for cancer therapy: from mouse models to clinical trials.
Galectins and their ligands: negative regulators of anti-tumor immunity.
Combination of T-Cell Bispecific Antibodies With PD-L1 Checkpoint Inhibition Elicits Superior Anti-Tumor Activity.
GCN2 is essential for CD8(+) T cell survival and function in murine models of malignant glioma.
Adoptive tumor therapy with T lymphocytes enriched through an IFN-gamma capture assay.
T-cell immune function in tumor, skin, and peripheral blood of advanced stage melanoma patients: implications for immunotherapy.
Distinct role for CD8 T cells toward cutaneous tumors and visceral metastases.
Targeting costimulatory pathways for tumor immunotherapy.
Insufficient CD100 shedding contributes to suppression of CD8(+) T-cell activity in non-small cell lung cancer.
Induction of contact-dependent CD8(+) regulatory T cells through stimulation with staphylococcal and streptococcal superantigens.
Cellular immune responses to hepatocellular carcinoma: lessons for immunotherapy.
[Not Available].
Defining the mechanisms of CD8 T-cell tumor tolerance.
Monitoring Patient Response to Pembrolizumab With Peripheral Blood Exhaustion Marker Profiles.

CD4(+) T cell exhaustion leads to adoptive transfer therapy failure which can be prevented by immune checkpoint blockade.
T cell anergy, exhaustion, senescence, and stemness in the tumor microenvironment.
LAG-3xPD-L1 bispecific antibody potentiates antitumor responses of T cells through dendritic cell activation.
Transgenic Expression of IL15 Retains CD123-Redirected T Cells in a Less Differentiated State Resulting in Improved Anti-AML Activity in Autologous AML PDX Models.
Curcumin reverses T cell-mediated adaptive immune dysfunctions in tumor-bearing hosts.
Latent HIV reservoirs exhibit inherent resistance to elimination by CD8+ T cells.
The CD4+ T Cell Response to Human Cytomegalovirus in Healthy and Immunocompromised People.
Clinical experience with gene therapy and bispecific antibodies for T cell-based therapy of cancer.
CD8+ T cell evasion mandates CD4+ T cell control of chronic gamma-herpesvirus infection.
Liver sinusoidal endothelial cells contribute to CD8 T cell tolerance toward circulating carcinoembryonic antigen in mice.
CD8(+) T cells specific for tumor antigens can be rendered dysfunctional by the tumor microenvironment through upregulation of the inhibitory receptors BTLA and PD-1.
Modulation of antigenic location converts chronic into acute infection by forcing CD8+ T cell recognition.
Cross-presentation of a spread-defective MCMV is sufficient to prime the majority of virus-specific CD8+ T cells.
The immune checkpoint regulator PD-L1 is a specific target for naturally occurring CD4(+) T cells.
Reduced tumor-antigen density leads to PD-1/PD-L1-mediated impairment of partially exhausted CD8+ T cells.
Peculiar Phenotypic and Cytotoxic Features of Pulmonary Mucosal CD8 T Cells in People Living with HIV Receiving Long-Term Antiretroviral Therapy.
SMAD4, activated by the TCR-triggered MEK/ERK signaling pathway, critically regulates CD8(+) T cell cytotoxic function.
Neuroendocrine Regulation of Stress-Induced T Cell Dysfunction during Lung Cancer Immunosurveillance via the Kisspeptin/GPR54 Signaling Pathway.
Genome-wide fitness gene identification reveals Roquin as a potent suppressor of CD8 T cell expansion and anti-tumor immunity.
Co-inhibition of TIGIT, PD1, and Tim3 reverses dysfunction of Wilms tumor protein-1 (WT1)-specific CD8+ T lymphocytes after dendritic cell vaccination in gastric cancer.
Durable adoptive immunotherapy for leukemia produced by manipulation of multiple regulatory pathways of CD8+ T-cell tolerance.
Antibodies Against Immune Checkpoint Molecules Restore Functions of Tumor-Infiltrating T Cells in Hepatocellular Carcinomas.
Colorectal cancer-derived Foxp3(+) IL-17(+) T cells suppress tumour-specific CD8+ T cells.
TRIB3 reduces CD8(+) T cell infiltration and induces immune evasion by repressing the STAT1-CXCL10 axis in colorectal cancer.
Revisiting T Cell Tolerance as a Checkpoint Target for Cancer Immunotherapy.
COPS6 promotes tumor progression and reduces CD8(+) T cell infiltration by repressing IL-6 production to facilitate tumor immune evasion in breast cancer.
An increased number of PD-1+ and Tim-3+ CD8+ T cells is involved in immune evasion in gastric cancer.
Adoptive Immunotherapy with Antigen-Specific T Cells Expressing a Native TCR.

CXCR5(+)CD8(+) T cells present elevated capacity in mediating cytotoxicity toward autologous tumor cells through interleukin 10 in diffuse large B-cell lymphoma.
Tumor-infiltrating regulatory T cells delineated by upregulation of PD-1 and inhibitory receptors.
Antigen presentation under the influence of 'immune evasion' proteins and its modulation by interferon-gamma: implications for immunotherapy of cytomegalovirus infection with antiviral CD8 T cells.
Immune-related genes in tumor-specific CD4(+) and CD8(+) T cells in colon cancer.
CD40 ligation reverses T cell tolerance in acute myeloid leukemia.
A Novel Immunomodulator Delivery Platform Based on Bacterial Biomimetic Vesicles for Enhanced Antitumor Immunity.
CD8(+) T Cell-Based Molecular Classification With Heterogeneous Immunogenomic Landscapes and Clinical Significance of Clear Cell Renal Cell Carcinoma.
IL10 and PD-1 Cooperate to Limit the Activity of Tumor-Specific CD8+ T Cells.
Bone marrow endothelial cells sustain a tumor-specific CD8(+) T cell subset with suppressive function in myeloma patients.
Clonal Abundance of Tumor-Specific CD4(+) T Cells Potentiates Efficacy and Alters Susceptibility to Exhaustion.
Batf3 transcription factor-dependent DC subsets in murine CMV infection: differential impact on T-cell priming and memory inflation.
Increased Coexpression of PD-1, TIGIT, and KLRG-1 on Tumor-Reactive CD8(+) T Cells During Relapse after Allogeneic Stem Cell Transplantation.
Tumor antigen-specific CD8(+) T cells are negatively regulated by PD-1 and Tim-3 in human gastric cancer.
A TIGIT-based chimeric co-stimulatory switch receptor improves T-cell anti-tumor function.
T cell sensitivity and the outcome of viral infection.
CD4 and CD8 T lymphocyte interplay in controlling tumor growth.
Silencing EGFR-upregulated expression of CD55 and CD59 activates the complement system and sensitizes lung cancer to checkpoint blockade.
New Strategies in Engineering T-cell Receptor Gene-Modified T cells to More Effectively Target Malignancies.
Tgf- $\beta$ 1 produced by activated CD4(+) T Cells Antagonizes T Cell Surveillance of Tumor Development.
Engineering adoptive T cell therapy to co-opt Fas ligand-mediated death signaling in ovarian cancer enhances therapeutic efficacy.
The role of CD8 T lymphocytes in rickettsial infections.
Antigen spreading-induced CD8+T cells confer protection against the lethal challenge of wild-type malignant mesothelioma by eliminating myeloid-derived suppressor cells.
Bortezomib augments lymphocyte stimulatory cytokine signaling in the tumor microenvironment to sustain CD8+T cell antitumor function.
Engineering T Cells to Express Tumoricidal MDA-7/IL24 Enhances Cancer Immunotherapy.
Effect of pembrolizumab on CD4(+) CD25(+) , CD4(+) LAP(+) and CD4(+) TIM-3(+) T cell subsets.
IL-10 elicits IFN $\gamma$ -dependent tumor immune surveillance.
PD-1/PD-L1 interactions contribute to functional T-cell impairment in patients who relapse with cancer after allogeneic stem cell transplantation.
PD-L1 Overexpression, SWI/SNF Complex Deregulation, and Profound Transcriptomic Changes Characterize Cancer-Dependent Exhaustion of Persistently Activated CD4(+) T Cells.
Pretreatment Innate Cell Populations and CD4 T Cells in Blood Are Associated With Response to Immune Checkpoint Blockade in Melanoma Patients.

Tipping the scales: Immunotherapeutic strategies that disrupt immunosuppression and promote immune activation.
Premature terminal exhaustion of Friend virus-specific effector CD8+ T cells by rapid induction of multiple inhibitory receptors.
Human Pancreatic Carcinoma-Associated Fibroblasts Promote Expression of Co-inhibitory Markers on CD4(+) and CD8(+) T-Cells.
Simulated microgravity-mediated reversion of murine lymphoma immune evasion.
Adoptive transfer of cytotoxic T lymphocytes targeting two different antigens limits antigen loss and tumor escape.
HIV-1 Nef equips dendritic cells to reduce survival and function of CD8+ T cells: a mechanism of immune evasion.
[Elevated Fas expression is related to increased apoptosis of circulating CD8(+)T cell in patients with hepatocellular carcinoma].
Adenovirus Improves the Efficacy of Adoptive T-cell Therapy by Recruiting Immune Cells to and Promoting Their Activity at the Tumor.
Reciprocal changes in tumor antigenicity and antigen-specific T cell function during tumor progression.
Combination of Sunitinib and PD-L1 Blockade Enhances Anticancer Efficacy of TLR7/8 Agonist-Based Nanovaccine.
Infiltration of T cells promotes the metastasis of ovarian cancer cells via the modulation of metastasis-related genes and PD-L1 expression.
Contrasting Roles of the PD-1 Signaling Pathway in Dendritic Cell-Mediated Induction and Regulation of HIV-1-Specific Effector T Cell Functions.
CD8(+) T cells mediate the antitumor activity of frankincense and myrrh in hepatocellular carcinoma.
Enhancing adoptive CD8 T cell therapy by systemic delivery of tumor associated antigens.
Optimal activation of tumor-reactive T cells by selected antigenic peptide analogues.
Diacylglycerol Kinase $\zeta$ Limits Cytokine-dependent Expansion of CD8(+) T Cells with Broad Antitumor Capacity.
Tumor antigen-specific CD8 T cells infiltrating the tumor express high levels of PD-1 and are functionally impaired.
A new hope in immunotherapy for malignant gliomas: adoptive T cell transfer therapy.
Tumor resistance to CD8+ T cell-based therapeutic vaccination.
Regulation of virus-specific CD4+ T cell function by multiple costimulatory receptors during chronic HIV infection.
Tumor-Residing Batf3 Dendritic Cells Are Required for Effector T Cell Trafficking and Adoptive T Cell Therapy.
Indoleamine 2,3-dioxygenase regulates anti-tumor immunity in lung cancer by metabolic reprogramming of immune cells in the tumor microenvironment.
The Establishment and Experimental Verification of an lncRNA-Derived CD8+ T Cell Infiltration ceRNA Network in Colorectal Cancer.
Crucial role of CD69 in anti-tumor immunity through regulating the exhaustion of tumor-infiltrating T cells.
Clonality, Antigen Recognition, and Suppression of CD8(+) T Cells Differentially Affect Prognosis of Breast Cancer Subtypes.
Unbiased chemokine receptor screening reveals similar efficacy of lymph node- and tumor-targeted T cell immunotherapy.
PSGL-1 Is an Immune Checkpoint Regulator that Promotes T Cell Exhaustion.

Squamous cell carcinomas escape immune surveillance via inducing chronic activation and exhaustion of CD8+ T Cells co-expressing PD-1 and LAG-3 inhibitory receptors.
Neoantigen landscape dynamics during human melanoma-T cell interactions.
Successful elimination of large established tumors and avoidance of antigen-loss variants by aggressive adoptive T cell immunotherapy.
Tumor evasion from T cell surveillance.
Neoadjuvant PD-1 blockade induces T cell and cDC1 activation but fails to overcome the immunosuppressive tumor associated macrophages in recurrent glioblastoma.
Ex vivo priming of CD4 T cells converts immunological tolerance into effective antitumor immunity in a murine model of acute lymphoblastic leukemia.
Urothelial bladder cancer may suppress perforin expression in CD8+ T cells by an ICAM-1/TGFβ2 mediated pathway.
T cell surveillance of oncogene-induced prostate cancer is impeded by T cell-derived TGF-β1 cytokine.
Carboxyamidotriazole combined with IDO1-Kyn-AhR pathway inhibitors profoundly enhances cancer immunotherapy.
Resisting fatal attraction: a glioma oncometabolite prevents CD8+ T cell recruitment.
[Expression and Significance of PD-1, TIM-3 and VISTA on T Cell of Acute Myeloid Leukemia Patients].
Soluble monomeric human programmed cell death-ligand 1 inhibits the functions of activated T cells.
A(2A) Adenosine Receptor Gene Deletion or Synthetic A(2A) Antagonist Liberate Tumor-Reactive CD8(+) T Cells from Tumor-Induced Immunosuppression.
DNA fusion vaccines enter the clinic.
T-cell immunometabolism against cancer.
Engineered Adoptive T-cell Therapy Prolongs Survival in a Preclinical Model of Advanced-Stage Ovarian Cancer.
23 assessment of T-cell immune dysfunction in patients with renal cell carcinoma.
Mechanisms of T-Cell Exhaustion in Pancreatic Cancer.
Altered peptide ligands narrow the repertoire of cellular immune responses by interfering with T-cell priming.
Modulation of microenvironment acidity reverses anergy in human and murine tumor-infiltrating T lymphocytes.
Immune suppression in the tumor microenvironment.
Anti-CTLA-4 antibody-functionalized dendritic cell-derived exosomes targeting tumor-draining lymph nodes for effective induction of antitumor T-cell responses.
Determination of poor prognostic immune features of tumour microenvironment in non-smoking patients with lung adenocarcinoma.
Natural T cell immunity against cancer.
Spontaneous apoptosis of tumor-specific tetramer+ CD8+ T lymphocytes in the peripheral circulation of patients with head and neck cancer.
Differences in PD-1 expression on CD8+ T-cells in chronic myeloid leukemia patients according to disease phase and TKI medication.
CD147 regulates antitumor CD8(+) T-cell responses to facilitate tumor-immune escape.
Tumors Escape CD4+ T-cell-Mediated Immunosurveillance by Impairing the Ability of Infiltrating Macrophages to Indirectly Present Tumor Antigens.
Improving T cell therapy for cancer.
miR-149-3p reverses CD8(+) T-cell exhaustion by reducing inhibitory receptors and promoting cytokine secretion in breast cancer cells.

The cytomegalovirus UL146 gene product vCXCL1 promotes the resistance of hepatic cells to CD8(+) T cells through up-regulation of PD-L1.
TIGIT in cancer immunotherapy.
Staphylococcus aureus alpha-toxin inhibits CD8(+) T cell-mediated killing of cancer cells in cutaneous T-cell lymphoma.
The role of dendritic cells in cancer.
PHA eludes macrophage suppression to activate CD8(+) T cells.
T-cell agonists in cancer immunotherapy.
Tumor microenvironment and immune escape.
Integrating T cell metabolism in cancer immunotherapy.
A model for CD8+ CTL tumor immunosurveillance and regulation of tumor escape by CD4 T cells through an effect on quality of CTL.
PD-1 is a regulator of NY-ESO-1-specific CD8+ T cell expansion in melanoma patients.
Targeting the immunoglobulin IGSF9 enhances anti-tumor T cell activity and sensitivity to anti-PD-1 immunotherapy.
Overexpression of PD-1 and CD39 in tumor-infiltrating lymphocytes compared with peripheral blood lymphocytes in triple-negative breast cancer.
Tumor-induced CD11b+Gr-1+ myeloid cells suppress T cell sensitization in tumor-draining lymph nodes.
Antigen loss and tumor-mediated immunosuppression facilitate tumor recurrence.
VPS9D1-AS1 overexpression amplifies intratumoral TGF- $\beta$ signaling and promotes tumor cell escape from CD8(+) T cell killing in colorectal cancer.
Newly emerged immunogenic neoantigens in established tumors enable hosts to regain immunosurveillance in a T-cell-dependent manner.
IFN- $\gamma$ R $\alpha$ is a key determinant of CD8+ T cell-mediated tumor elimination or tumor escape and relapse in FVB mouse.
Tumor progression can occur despite the induction of very high levels of self/tumor antigen-specific CD8+ T cells in patients with melanoma.
A novel approach to characterize clonality and differentiation of human melanoma-specific T cell responses: spontaneous priming and efficient boosting by vaccination.
Identifying and overcoming immune resistance mechanisms in the melanoma tumor microenvironment.
Active secretion of CXCL10 and CCL5 from colorectal cancer microenvironments associates with GranzymeB+ CD8+ T-cell infiltration.
Blockade of transforming growth factor- $\beta$ signaling in tumor-reactive CD8(+) T cells activates the antitumor immune response cycle.
Memory CD8+ T cells require CD28 costimulation.
A Well-Controlled Experimental System to Study Interactions of Cytotoxic T Lymphocytes with Tumor Cells.
In Vivo Priming of Peritoneal Tumor-Reactive Lymphocytes With a Potent Oncolytic Virus for Adoptive Cell Therapy.
PRC2 Plays Red Light, Green Light with MHC-I and CD8(+) T Cells.
Regulation of suppressive function of myeloid-derived suppressor cells by CD4+ T cells.
Enhanced therapeutic potential of adoptive immunotherapy by in vitro CD28/4-1BB costimulation of tumor-reactive T cells against a poorly immunogenic, major histocompatibility complex class I-negative A9P melanoma.

Expression of PD1 and BTLA on the CD8(+) T Cell and $\gamma\delta$ T Cell Subsets in Peripheral Blood of Non-Small Cell Lung Cancer Patients.
Generation of CD8+ T cell-mediated immunity against idiotype-negative lymphoma escapees.
Hepatitis D Virus-Specific CD8(+) T Cells Have a Memory-Like Phenotype Associated With Viral Immune Escape in Patients With Chronic Hepatitis D Virus Infection.
Eomes-Dependent Loss of the Co-activating Receptor CD226 Restrains CD8(+) T Cell Anti-tumor Functions and Limits the Efficacy of Cancer Immunotherapy.
A superagonist variant of peptide MART1/Melan A27-35 elicits anti-melanoma CD8+ T cells with enhanced functional characteristics: implication for more effective immunotherapy.
Nanoliposome C6-Ceramide Increases the Anti-tumor Immune Response and Slows Growth of Liver Tumors in Mice.
NFAT1 supports tumor-induced anergy of CD4(+) T cells.
Downregulation of MARC2 Promotes Immune Escape and Is Associated With Immunosuppression of Hepatocellular Carcinoma.
Distinct changes of in BTLA, ICOS, PD-1, and TIGIT expression on peripheral blood and decidual CD8+ T cells in women with unexplained recurrent spontaneous abortion <sup>†</sup> .
IRF1 Inhibits Antitumor Immunity through the Upregulation of PD-L1 in the Tumor Cell.
Metabolic regulation of T cells in the tumor microenvironment by nutrient availability and diet.
Immunoregulatory T cells in tumor immunity.
PD1 is transcriptionally regulated by LEF1 in mature T cells.
Cyclin-dependent kinase 9 expression and its association with CD8(+) T cell infiltration in microsatellite-stable colorectal cancer.
Preclinical models for prediction of immunotherapy outcomes and immune evasion mechanisms in genetically heterogeneous multiple myeloma.
T helper responses are maintained by basal-like breast cancer cells and confer to immune modulation via upregulation of PD-1 ligands.
Shifting the equilibrium in cancer immunoediting: from tumor tolerance to eradication.
Adoptive T cell therapy promotes the emergence of genomically altered tumor escape variants.
TIGIT signaling and its influence on T cell metabolism and immune cell function in the tumor microenvironment.
Differential Role of PD-1 Expressed by Various Immune and Tumor Cells in the Tumor Immune Microenvironment: Expression, Function, Therapeutic Efficacy, and Resistance to Cancer Immunotherapy.
Multivalent Forms of the Notch Ligand DLL-1 Enhance Antitumor T-cell Immunity in Lung Cancer and Improve Efficacy of EGFR-Targeted Therapy.
Sialic Acid Blockade Suppresses Tumor Growth by Enhancing T-cell-Mediated Tumor Immunity.
IL-6-mediated environmental conditioning of defective Th1 differentiation dampens antitumour immune responses in old age.
Tumor-infiltrating CD39(+)CD8(+) T cells determine poor prognosis and immune evasion in clear cell renal cell carcinoma patients.
CD8+ T cell-Dependent Remodeling of the Tumor Microenvironment Overcomes Chemoresistance.
A high proportion of bone marrow T cells with regulatory phenotype (CD4+CD25hiFoxP3+) in Ewing sarcoma patients is associated with metastatic disease.
Mycobacterium tuberculosis Membrane Vesicles Inhibit T Cell Activation.
Combination of PD-1 Inhibitor and OX40 Agonist Induces Tumor Rejection and Immune Memory in Mouse Models of Pancreatic Cancer.
T cell immunity to cytomegalovirus infection.

Rescue of tumor-infiltrating lymphocytes from activation-induced cell death enhances the antitumor CTL response in CD5-deficient mice.
The Progress of T Cell Immunity Related to Prognosis in Gastric Cancer.
PD-1 Expression on Circulating CD8(+) T-Cells as a Prognostic Marker for Patients With Gastric Cancer.
Numbers of CD8+PD-1+ and CD4+PD-1+ Cells in Peripheral Blood of Patients with Chronic Lymphocytic Leukemia Are Independent of Binet Stage and Are Significantly Higher Compared to Healthy Volunteers.
CXCR6 and CCR5 localize T lymphocyte subsets in nasopharyngeal carcinoma.
Functional and Clinical Characterization of Tumor-Infiltrating T Cell Subpopulations in Hepatocellular Carcinoma.
PD-1 coinhibitory signals: the link between pathogenesis and protection.
Tim-3 is highly expressed in T cells in acute myeloid leukemia and associated with clinicopathological prognostic stratification.
Tissue homing and persistence of defined antigen-specific CD8+ tumor-reactive T-cell clones in long-term melanoma survivors.
Less yin, more yang: confronting the barriers to cancer immunotherapy.
Tumor Endothelial Cell-Mediated Antigen-Specific T-cell Suppression via the PD-1/PD-L1 Pathway.
Characteristics and critical function of CD8+ T cells in the Toxoplasma-infected brain.
Follicular CD8(+) T Cells: Origin, Function and Importance during HIV Infection.
Costimulation tunes tumor-specific activation of redirected T cells in adoptive immunotherapy.
Cellular and cytokine-dependent immunosuppressive mechanisms of grm1-transgenic murine melanoma.
CD8+ T cells stimulated by exosomes derived from RenCa cells mediate specific immune responses through the FasL/Fas signaling pathway and, combined with GM-CSF and IL-12, enhance the anti-renal cortical adenocarcinoma effect.
Using signaling pathways to overcome immune tolerance to tumors.
Therapeutic cancer vaccination against mutant calreticulin in myeloproliferative neoplasms induces expansion of specific T cells in the periphery but specific T cells fail to enrich in the bone marrow.
Cytotoxic T-cells mediate exercise-induced reductions in tumor growth.
Immune Checkpoint Blockade for Breast Cancer.
Lymphatic dysfunction attenuates tumor immunity through impaired antigen presentation.
NKILA lncRNA promotes tumor immune evasion by sensitizing T cells to activation-induced cell death.
Long-term surviving influenza infected cells evade CD8+ T cell mediated clearance.
Targeted inhibition of galectin-1 gene expression in tumor cells results in heightened T cell-mediated rejection; A potential mechanism of tumor-immune privilege.
Indoleamine 2,3-dioxygenase regulates T cell activity through Vav1/Rac pathway.
Slow progression of pediatric HIV associates with early CD8+ T cell PD-1 expression and a stem-like phenotype.
Analysis of cancer cell-intrinsic immune regulation in response to CD8+ T cell attack.
Exosome-derived circCCAR1 promotes CD8 + T-cell dysfunction and anti-PD1 resistance in hepatocellular carcinoma.
Infiltration of tumor-reactive transforming growth factor-beta insensitive CD8+ T cells into the tumor parenchyma is associated with apoptosis and rejection of tumor cells.
To Remember or to Forget: The Role of Good and Bad Memories in Adoptive T Cell Therapy for Tumors.
DCOne as an Allogeneic Cell-based Vaccine for Multiple Myeloma.



Overcoming current challenges to T-cell receptor therapy via metabolic targeting to increase antitumor efficacy, durability, and tolerability.
Increased regulatory T-cell fraction amidst a diminished CD4 compartment explains cellular immune defects in patients with malignant glioma.
Induction of anergic or regulatory tumor-specific CD4(+) T cells in the tumor-draining lymph node.
Gastric Carcinomas with Stromal B7-H3 Expression Have Lower Intratumoural CD8+ T Cell Density.
Role of FABP5 in T Cell Lipid Metabolism and Function in the Tumor Microenvironment.
A universal anti-cancer vaccine: Chimeric invariant chain potentiates the inhibition of melanoma progression and the improvement of survival.
Melanoma-intrinsic $\beta$ -catenin signalling prevents anti-tumour immunity.
An Engineered AAV6-Based Vaccine Induces High Cytolytic Anti-Tumor Activity by Directly Targeting DCs and Improves Ag Presentation.
Immunogenomics of Colorectal Cancer Response to Checkpoint Blockade: Analysis of the KEYNOTE 177 Trial and Validation Cohorts.
CD8+ T cell tolerance following antigen recognition on hepatocytes.
B7-DC-Ig enhances vaccine effect by a novel mechanism dependent on PD-1 expression level on T cell subsets.
IFN- $\gamma$ down-regulates the PD-1 expression and assist nivolumab in PD-1-blockade effect on CD8+ T-lymphocytes in pancreatic cancer.
Improving T cell therapy for cancer.
Immune Escape After Adoptive T-cell Therapy for Malignant Gliomas.
In situ T cells in melanoma.
Single-cell analysis of localized prostate cancer patients links high Gleason score with an immunosuppressive profile.
B lymphocytes can be activated to act as antigen presenting cells to promote anti-tumor responses.
Tumors attenuating the mitochondrial activity in T cells escape from PD-1 blockade therapy.
IL4I1: Key immunoregulator at a crossroads of divergent T-cell functions.
The use of endogenous T cells for adoptive transfer.
Immune suppressive landscape in the human esophageal squamous cell carcinoma microenvironment.
Intratumoral TIGIT(+) CD8(+) T-cell infiltration determines poor prognosis and immune evasion in patients with muscle-invasive bladder cancer.
Lipid Metabolism in Tumor-Infiltrating T Cells.
Eradication of medullary multiple myeloma by CD4+ cytotoxic human T lymphocytes directed at a single minor histocompatibility antigen.
Malaria drives T cells to exhaustion.
Cross-presentation of antigens from apoptotic tumor cells by liver sinusoidal endothelial cells leads to tumor-specific CD8+ T cell tolerance.
Role of LAP(+)CD4(+) T cells in the tumor microenvironment of colorectal cancer.
Adenosine mediates functional and metabolic suppression of peripheral and tumor-infiltrating CD8(+) T cells.
Emerging strategies in regulatory T-cell immunotherapies.
Directing Traffic: How to Effectively Drive T Cells into Tumors.
Molecular Profile of Tumor-Specific CD8+ T Cell Hypofunction in a Transplantable Murine Cancer Model.
Active evasion of CTL mediated killing and low quality responding CD8+ T cells contribute to persistence of brucellosis.

Combination Approaches with Immune-Checkpoint Blockade in Cancer Therapy.
Survival of the fittest: Cancer challenges T cell metabolism.
ZEB1 transcription factor promotes immune escape in melanoma.
Immune-escape markers in relation to clinical outcome of advanced melanoma patients following immunotherapy.
Novel strategies for cancer therapy: the potential of genetically modified T lymphocytes.
HBZ is an immunogenic protein, but not a target antigen for human T-cell leukemia virus type 1-specific cytotoxic T lymphocytes.
A Model for Apoptotic-Cell-Mediated Adaptive Immune Evasion via CD80-CTLA-4 Signaling.
Analysis of expression of the melanoma-associated antigens MART-1 and gp100 in metastatic melanoma cell lines and in situ lesions.
NFATc2 is an intrinsic regulator of melanoma dedifferentiation.
Dissection of transcriptome dysregulation and immune characterization in women with germline BRCA1 mutation at single-cell resolution.
The role of the immune response in merkel cell carcinoma.
NPM1 and DNMT3A mutations are associated with distinct blast immunophenotype in acute myeloid leukemia.
N-3-(oxododecanoyl)-L-homoserine lactone suppresses dendritic cell maturation by upregulating the long noncoding RNA NRIR.
Surfaceome Profiling of Rhabdomyosarcoma Reveals B7-H3 as a Mediator of Immune Evasion.
Epigenetic Therapy Promotes the Ratio of Th1/Th17 Lineage to Reverse Immune Evasion and Treat Leukemia Relapse Post-allogeneic Stem Cell Transplantation in Non-APL AML Patients.
Dysregulated Expression of CD28 and CTLA-4 Molecules in Patients with Acute Myeloid Leukemia and Possible Association with Development of Graft versus Host Disease after Hematopoietic Stem Cell Transplantation.
PSMD2 promotes the progression of bladder cancer and is correlated with immune infiltration.
MYC-driven synthesis of Siglec ligands is a glycoimmune checkpoint.
Investigation of indoleamine 2,3-dioxygenase 1 expression in uveal melanoma.
A fiber-modified adenoviral vector interacts with immunoevasion molecules of the B7 family at the surface of murine leukemia cells derived from dormant tumors.
Development of Multidrug Resistance in Acute Myeloid Leukemia Is Associated with Alterations of the LPHN1/GAL-9/TIM-3 Signaling Pathway.
Cutaneous T cell lymphoma expresses immunosuppressive CD80 (B7-1) cell surface protein in a STAT5-dependent manner.
Lymphocyte apoptosis induced by CD95 (APO-1/Fas) ligand-expressing tumor cells--a mechanism of immune evasion?
[The expression of zeta-chain of the T cell receptor as prognostic marker for patients with head and neck cancer].
ATM-dependent spontaneous regression of early Eμ-myc-induced murine B-cell leukemia depends on natural killer and T cells.
Integrated bioinformatic analysis and cell line experiments reveal the significant role of the novel immune checkpoint TIGIT in kidney renal clear cell carcinoma.
Tim-3 inhibits macrophage control of Listeria monocytogenes by inhibiting Nrf2.
Interferon-stimulated gene 15 promotes progression of endometrial carcinoma and weakens antitumor immune response.
Humoral immunodeficiency in chronic lymphocytic leukemia: role of CD95/CD95L in tumoral damage and escape.

Local immunomodulation combined to radiofrequency ablation results in a complete cure of local and distant colorectal carcinoma.
Anthrax lethal toxin blocks MAPK kinase-dependent IL-2 production in CD4+ T cells.
Uveal melanoma expression of indoleamine 2,3-deoxygenase: establishment of an immune privileged environment by tryptophan depletion.
Expression of the immune checkpoint modulator OX40 indicates poor survival in acute myeloid leukemia.
Soluble decoy receptor 3 is expressed by malignant gliomas and suppresses CD95 ligand-induced apoptosis and chemotaxis.
Tim-3 regulates pro- and anti-inflammatory cytokine expression in human CD14+ monocytes.
Cytokine Networks Dysregulation during HTLV-1 Infection and Associated Diseases.
Absent or low expression of T-cell receptor zeta-chain in T cells infiltrating human pathological skin conditions.
The Role of HBZ in HTLV-1-Induced Oncogenesis.
All infiltrating T-lymphocytes in Hodgkin's disease express immunohistochemically detectable T-cell receptor zeta-chains in situ.
Escape from IFN- $\gamma$ -dependent immunosurveillance in tumorigenesis.
CXCR3 blockade protects against <i>Listeria monocytogenes</i> infection-induced fetal wastage.
Expression of costimulatory molecules, B7-1 and B7-2 on human gastric carcinoma.
The relation of blood cell division control protein 42 level with disease risk, comorbidity, tumor features/markers, and prognosis in colorectal cancer patients.
Long noncoding RNA MIAT regulates TP53 ubiquitination and expedites prostate adenocarcinoma progression by recruiting TBL1X.
hMan2c1 transgene promotes tumor progress in mice.
Potential role of CCL27 and CCR10 expression in melanoma progression and immune escape.
Characterisation of the immune-related transcriptome in resected biliary tract cancers.
Targeting of IL-2 to cytotoxic lymphocytes as an improved method of cytokine-driven immunotherapy.
MTAP deficiency contributes to immune landscape remodelling and tumour evasion.
TNF Superfamily Networks: bidirectional and interference pathways of the herpesvirus entry mediator (TNFSF14).
NOX4 activation is involved in ROS-dependent Jurkat T-cell death induced by <i>Entamoeba histolytica</i> .
Application of monoclonal antibodies in a sandwich enzyme-linked immunosorbent assay for identification and detection of soluble human B and T lymphocyte attenuator.
Immunocyte density in parathyroid carcinoma is correlated with disease relapse.
Cholangiocarcinoma with respect to IgG4 Reaction.
B7-H4 expression in bladder urothelial carcinoma and immune escape mechanisms.
Molecular mechanisms of death ligand-mediated immune modulation: a gene therapy model to prolong islet survival in type 1 diabetes.
<i>Fusobacterium nucleatum</i> CbpF Mediates Inhibition of T Cell Function Through CEACAM1 Activation.
Methionine-enkephalin secreted by human colorectal cancer cells suppresses T lymphocytes.
TGF $\beta$ Promotes Immune Evasion to Limit the Efficacy of Anti-PD-1/PD-L1.
Engineered type 1 regulatory T cells designed for clinical use kill primary pediatric acute myeloid leukemia cells.
Cellular distribution of CD200 receptor in rats and its interaction with cytomegalovirus e127 protein.
Protective roles of epithelial cells in the survival of adult T-cell leukemia/lymphoma cells.

Clinical impact of CD200 expression in patients with acute myeloid leukemia and correlation with other molecular prognostic factors.
Altered phenotypic and functional characteristics of CD3+CD56+ NKT-like cells in human gastric cancer.
Effects of <i>Cysticercus cellulosae</i> Excretory-Secretory Antigens on the TGF- $\beta$ Signaling Pathway and Th17 Cell Differentiation in Piglets, a Proteomic Analysis.
Sub-clonal analysis of the murine C1498 acute myeloid leukaemia cell line reveals genomic and immunogenic diversity.
Evidence of immune elimination, immuno-editing and immune escape in patients with hematological cancer.
Uveal and cutaneous melanoma: shared expression characteristics of melanoma-associated antigens.
HTLV-1 Tax protects against CD95-mediated apoptosis by induction of the cellular FLICE-inhibitory protein (c-FLIP).
Selection and expression of CD40 single chain variable fragment by phage display and evaluation of tumor specific immune activation.
Immune-dysregulation in subacute sclerosing panencephalitis: An exploratory case-control study.
Zap70 controls the interaction of talin with integrin to regulate the chemotactic directionality of T-cell migration.
Engineering human stem cell-derived islets to evade immune rejection and promote localized immune tolerance.
Transgenic expression of decoy receptor 3 protects islets from spontaneous and chemical-induced autoimmune destruction in nonobese diabetic mice.
ATRX loss promotes immunosuppressive mechanisms in IDH1 mutant glioma.
Multiple Mechanisms of NOTCH1 Activation in Chronic Lymphocytic Leukemia: NOTCH1 Mutations and Beyond.
Intratumoral CCR5(+) neutrophils identify immunogenic subtype muscle-invasive bladder cancer with favorable prognosis and therapeutic responses.
Loss of the Immune Checkpoint CD85j/LILRB1 on Malignant Plasma Cells Contributes to Immune Escape in Multiple Myeloma.
Correction to: $\alpha$ 2,6-Sialylation promotes immune escape in hepatocarcinoma cells by regulating T cell functions and CD147/MMP signaling.
Autologous bone marrow Th cells can support multiple myeloma cell proliferation in vitro and in xenografted mice.
The binding of LDN193189 to CD133 C-terminus suppresses the tumorigenesis and immune escape of liver tumor-initiating cells.
Epigenetic modulation of MAGE-A3 antigen expression in multiple myeloma following treatment with the demethylation agent 5-azacitidine and the histone deacetylase inhibitor MGCD0103.
Parenchymal regression in chronic pancreatitis spares islets reprogrammed for the expression of NFkappaB and IAPs.
A "stealth effect": adenocarcinoma cells engineered to express TRAIL elude tumor-specific and allogeneic T cell reactions.
IFN-Independent STING Signaling: Friend or Foe?
Pan-Cancer Analysis Reveals Disrupted Circadian Clock Associates With T Cell Exhaustion.
CD123/CD33 dual-antibody modified liposomes effectively target acute myeloid leukemia cells and reduce antigen-negative escape.
Interactions between Ibrutinib and Anti-CD20 Antibodies: Competing Effects on the Outcome of Combination Therapy.

Th22 cells as well as Th17 cells expand differentially in patients with early-stage and late-stage myelodysplastic syndrome.
microRNA-155, induced by interleukin-1 $\beta$ , represses the expression of microphthalmia-associated transcription factor (MITF-M) in melanoma cells.
Tasmanian devil CD28 and CTLA4 capture CD80 and CD86 from adjacent cells.
Ectopic CD137 expression by rhabdomyosarcoma provides selection advantages but allows immunotherapeutic targeting.
Hoechst 33342 Staining Identifies the Progenitor Side Population in NOD.Cg-PrkdcscidIL2rgtmWjl/Sz Mice Harboring Pediatric Leukemias.
Human peripheral basophils extended phenotype shows a high expression of CD244 immuno-regulatory receptor.
[Effect of bone marrow stromal cells transfected with interleukin 18 on growth of intracranial glioma in rats].
[Parenchymal regression in chronic pancreatitis spares islets reprogrammed for expression of NFkappaB and IAPs].
Somatic mutations and clonal expansions in paroxysmal nocturnal hemoglobinuria.
Mixed hematopoietic molecular chimerism results in permanent transgene expression from retrovirally transduced hepatocytes in mice.
B7 family molecules: novel immunomodulators at the maternal-fetal interface.
[B7-H4-mediated immunoresistance is suppressed by PI3K/Akt/mTOR pathway inhibitors].
Canine mammary tumor cells transfected with B7-1 or B7-2 stimulate proliferation of peripheral blood mononuclear cells.
TGF $\beta$ activating integrins $\beta$ 6 and $\beta$ 8 are dysregulated in inflammatory skin disease and cutaneous melanoma.
IL-18 enhances SCF production of melanoma cells by regulating ROI and p38 MAPK activity.
Development of hemolytic paroxysmal nocturnal hemoglobinuria without graft loss following hematopoietic stem cell transplantation for acquired aplastic anemia.
Relationship between bone marrow failure syndromes and the presence of glycoposphatidyl inositol-anchored protein-deficient clones.

Table A2-60, Cluster 59

Cluster 59 focuses on cancer immunotherapy, emphasizing the targeting of cancer immune evasion (1368)
Cancer Vaccines: Toward the Next Breakthrough in Cancer Immunotherapy.
Immunotherapeutic and their immunological aspects: Current treatment strategies and agents.
Combination of cancer immunotherapy with clinically available drugs that can block immunosuppressive cells.
Immunotherapy of cancer in 2012.
Cancer immunotherapy: Strategies for personalization and combinatorial approaches.
The Evolving Role of Immune Checkpoint Inhibitors in Cancer Treatment.
Clinical Combinatorial Treatments Based on Cancer Vaccines: Combination with Checkpoint Inhibitors and Beyond.
Primary, Adaptive, and Acquired Resistance to Cancer Immunotherapy.
Cancer Vaccine Therapeutics: Limitations and Effectiveness-A Literature Review.
[Development of novel immunotherapy targeting cancer immune evasion].

Targeting the immune system in cancer.
Immune based therapies in cancer.
Future approaches in immunotherapy.
Immune escape mechanisms as a guide for cancer immunotherapy.
Mechanisms of Immunotherapy Resistance in Cutaneous Melanoma: Recognizing a Shapeshifter.
Towards curative cancer immunotherapy: overcoming posttherapy tumor escape.
Tumor immunology and cancer immunotherapy: summary of the 2013 SITC primer.
Perspectives in Melanoma: meeting report from the Melanoma Bridge (December 2nd - 4th, 2021, Italy).
Current perspectives on immunotherapy.
Combinatorial Approach to Improve Cancer Immunotherapy: Rational Drug Design Strategy to Simultaneously Hit Multiple Targets to Kill Tumor Cells and to Activate the Immune System.
Immunotherapy as a Promising Treatment for Prostate Cancer: A Systematic Review.
The Reciprocity between Radiotherapy and Cancer Immunotherapy.
Biomarkers of related driver genes predict anti-tumor efficacy of immune checkpoint inhibitors.
Lung cancer-associated tumor antigens and the present status of immunotherapy against non-small-cell lung cancer.
The Evasion Mechanisms of Cancer Immunity and Drug Intervention in the Tumor Microenvironment.
Cancer immunotherapies: advances and bottlenecks.
Therapeutic targets and biomarkers of tumor immunotherapy: response versus non-response.
From immune checkpoints to vaccines: The past, present and future of cancer immunotherapy.
Multi-target combinatory strategy to overcome tumor immune escape.
Developments in cancer immunotherapy.
Mechanisms of immune evasion in bladder cancer.
Top 10 Challenges in Cancer Immunotherapy.
Engineered Nanoparticles for Cancer Vaccination and Immunotherapy.
Immunotherapy in gastric cancer.
Immunomodulatory effects of current cancer treatment and the consequences for follow-up immunotherapeutics.
Unraveling the crosstalk between melanoma and immune cells in the tumor microenvironment.
Progress and prospects of immune checkpoint inhibitors in advanced gastric cancer.
Therapeutic Cancer Vaccine and Combinations With Antiangiogenic Therapies and Immune Checkpoint Blockade.
Associating resistance to immune checkpoint inhibitors with immunological escape in colorectal cancer.
Cancer immunotherapy: A comprehensive appraisal of its modes of application.
A Highlight of the Mechanisms of Immune Checkpoint Blocker Resistance.
Active immunization against cancer cells: impediments and advances.
SITC cancer immunotherapy resource document: a compass in the land of biomarker discovery.
Frontiers in cancer immunotherapy-a symposium report.
Altering Landscape of Cancer Vaccines: Unique Platforms, Research on Therapeutic Applications and Recent Patents.
[Immunotherapy of cancer: promise and reality].
Acquired resistance to cancer immunotherapy.
Tackling Resistance to Cancer Immunotherapy: What Do We Know?

The history and advances in cancer immunotherapy: understanding the characteristics of tumor-infiltrating immune cells and their therapeutic implications.
Harnessing the immune system by targeting immune checkpoints: Providing new hope for Oncotherapy.
Development of novel immune interventions for prostate cancer.
Immunotherapy for lung cancers.
Reverting Immune Suppression to Enhance Cancer Immunotherapy.
Overcoming malignant cell-based mechanisms of resistance to immune checkpoint blockade antibodies.
Intratumoral Immunotherapy: From Trial Design to Clinical Practice.
Rational Combinations of Targeted Therapy and Immune Checkpoint Inhibitors in Head and Neck Cancers.
HDAC inhibitors as epigenetic regulators for cancer immunotherapy.
Engineering Nanoparticles for Targeted Remodeling of the Tumor Microenvironment to Improve Cancer Immunotherapy.
The Role of Somatic Mutations on the Immune Response of the Tumor Microenvironment in Prostate Cancer.
Immunotherapeutic strategies employing RNA interference technology for the control of cancers.
Bispecific PSMA antibodies and CAR-T in metastatic castration-resistant prostate cancer.
Enhancing Therapeutic Approaches for Melanoma Patients Targeting Epigenetic Modifiers.
Immunotherapy in hematologic malignancies: achievements, challenges and future prospects.
Emerging Prospects for Nanoparticle-Enabled Cancer Immunotherapy.
The Immunotherapy and Immunosuppressive Signaling in Therapy-Resistant Prostate Cancer.
Cancer-induced heterogeneous immunosuppressive tumor microenvironments and their personalized modulation.
Recent updates on cancer immunotherapy.
Immune tumor board: integral part in the multidisciplinary management of cancer patients treated with cancer immunotherapy.
Cancer immunotherapy: accomplishments to date and future promise.
Interleukin-34 and immune checkpoint inhibitors: Unified weapons against cancer.
siRNA and cancer immunotherapy.
Immunotherapy for colorectal cancer.
Immunotherapies catering to the unmet medical need of cold colorectal cancer.
Mechanisms and therapeutic potentials of cancer immunotherapy in combination with radiotherapy and/or chemotherapy.
Tumor-induced escape mechanisms and their association with resistance to checkpoint inhibitor therapy.
Primary and Acquired Resistance to Immune Checkpoint Inhibitors in Metastatic Melanoma.
Turning the corner on therapeutic cancer vaccines.
Vaccinations for Colorectal Cancer: Progress, Strategies, and Novel Adjuvants.
Immunotherapy for melanoma.
Understanding of Immune Escape Mechanisms and Advances in Cancer Immunotherapy.
Advantages of targeting the tumor immune microenvironment over blocking immune checkpoint in cancer immunotherapy.
Cancer Immunotherapy: The Checkpoint between Chronic Colitis and Colorectal Cancer.
The interplay of immunotherapy and chemotherapy: harnessing potential synergies.

Immune Escape in Glioblastoma Multiforme and the Adaptation of Immunotherapies for Treatment.
DNA vaccines to attack cancer: Strategies for improving immunogenicity and efficacy.
Combination Immunotherapy: Taking Cancer Vaccines to the Next Level.
Vaccine Therapies for Cancer: Then and Now.
Identification and targeting of tumor escape mechanisms: a new hope for cancer therapy?
CHALLENGES IN MANIPULATING IMMUNE SYSTEM TO TREAT PROSTATE CANCER.
Understanding Tumor-Antigen Presentation in the New Era of Cancer Immunotherapy.
Perspectives in Immunotherapy: meeting report from Immunotherapy Bridge (Naples, November 30th-December 1st, 2022).
Advances in the Lung Cancer Immunotherapy Approaches.
Combining Radiotherapy and Immunotherapy in Lung Cancer: Can We Expect Limitations Due to Altered Normal Tissue Toxicity?
Transient cell-in-cell formation underlies tumor relapse and resistance to immunotherapy.
Immunotherapy Resistance in Glioblastoma.
Immunotherapy of gastric cancer: Past, future perspective and challenges.
Prostate cancer immunotherapy.
Non-coding RNAs: Emerging roles in the characterization of immune microenvironment and immunotherapy of prostate cancer.
Inhibiting the inhibitors: evaluating agents targeting cancer immunosuppression.
The relationship between autophagy and PD-L1 and their role in antitumor therapy.
Various Uses of PD1/PD-L1 Inhibitor in Oncology: Opportunities and Challenges.
Current state and future of co-inhibitory immune checkpoints for the treatment of glioblastoma.
The role of immunotherapy in prostate cancer: an overview of current approaches in development.
Developments in immunotherapy for gastrointestinal cancer.
Cancer immunotherapy: are we there yet?
Therapeutic Strategies to Enhance Tumor Antigenicity: Making the Tumor Detectable by the Immune System.
Nanoparticle mediated cancer immunotherapy.
Immunomodulatory MicroRNAs in cancer: targeting immune checkpoints and the tumor microenvironment.
Immune Reactivation by Cell-Free Fetal DNA in Healthy Pregnancies Re-Purposed to Target Tumors: Novel Checkpoint Inhibition in Cancer Therapeutics.
Immune system and melanoma biology: a balance between immunosurveillance and immune escape.
Advances of immune checkpoints in colorectal cancer treatment.
Future Challenges in Cancer Resistance to Immunotherapy.
Oncolytic Viruses and Immune Checkpoint Inhibition: The Best of Both Worlds.
Immunotherapy of melanoma.
Prognostic Biomarkers for Melanoma Immunotherapy.
Trends and advances in tumor immunology and lung cancer immunotherapy.
Review of the Immune Checkpoint Inhibitors in the Context of Cancer Treatment.
The emerging role of miRNA in the perturbation of tumor immune microenvironment in chemoresistance: Therapeutic implications.
Prostate cancer vaccines: current status and future potential.
Immunotherapy for colorectal cancer: where are we heading?
Molecular Biomarkers of Primary and Acquired Resistance to T-Cell-Mediated Immunotherapy in Cancer: Landscape, Clinical Implications, and Future Directions.



The Changing Landscape of Therapeutic Cancer Vaccines- Novel Platforms and Neoantigen Identification.
Current immunotherapy for solid tumors.
Immuno-oncology-101: overview of major concepts and translational perspectives.
Perspectives in immunotherapy: meeting report from the "Immunotherapy Bridge", Napoli, November 30th 2016.
[At the crossroads of cancer].
Vaccine therapy for melanoma: current status and future directions.
[The Pedigree of Immuno-Oncology - History and Current Status of Immunotherapy against Malignancy].
Dendritic-cell vaccination for prostate cancer.
Guiding immunotherapy combinations: Who gets what?
Functional genomics: paving the way for more successful cancer immunotherapy.
Cancer Immunotherapy: Diverse Approaches and Obstacles.
Immunotherapy Approaches in Cancer Treatment.
Next generation of immunotherapy for melanoma.
Evolving Role of Immunotherapy in Metastatic Castration Refractory Prostate Cancer.
Bladder cancer, a unique model to understand cancer immunity and develop immunotherapy approaches.
Anti-cancer vaccine therapy for hematologic malignancies: An evolving era.
The Angiopoietin-2 and TIE Pathway as a Therapeutic Target for Enhancing Antiangiogenic Therapy and Immunotherapy in Patients with Advanced Cancer.
Generation of more effective cancer vaccines.
Recent progress in therapeutic antibodies for cancer immunotherapy.
Pancreatic Cancer and Immunotherapy: A Clinical Overview.
OMIC signatures to understand cancer immunosurveillance and immunoediting: Melanoma and immune cells interplay in immunotherapy.
Resistance mechanisms in melanoma to immuneoncologic therapy with checkpoint inhibitors.
Normalization Cancer Immunotherapy for Melanoma.
An Update on the Role of Immunotherapy and Vaccine Strategies for Primary Brain Tumors.
[Immune-checkpoints: the new anti-cancer immunotherapies].
Immunosuppressive mechanisms in human tumors: why we still cannot cure cancer.
Theranostics Applications of Nanoparticles in Cancer Immunotherapy.
Do novel treatment strategies enhance T cell-mediated Immunity: Opportunities and challenges in pancreatic cancer immunotherapy.
Therapeutic Cancer Vaccines for Nonmelanoma Skin Cancer.
Challenges and prospects of immunotherapy as cancer treatment.
Enhancing immune responses to tumor-associated antigens.
Fundamentals of Cancer Immunology and Their Application to Cancer Vaccines.
In situ vaccination: Harvesting low hanging fruit on the cancer immunotherapy tree.
Combining immunotherapy and radiotherapy in head and neck squamous cell cancers: which perspectives?
Prophylactic vaccine approach for colon and pancreatic cancers: present and future.
From Melanoma Development to RNA-Modified Dendritic Cell Vaccines: Highlighting the Lessons From the Past.
Cancer vaccines: harnessing the potential of anti-tumor immunity.

The Treatment of Advanced Melanoma: Therapeutic Update.
Ultrasound-Targeted Microbubble Destruction: Modulation in the Tumor Microenvironment and Application in Tumor Immunotherapy.
[Not Available].
Immunology of melanoma.
Vaccines for melanoma and renal cell carcinoma.
The Tumor Microenvironment: A Milieu Hindering and Obstructing Antitumor Immune Responses.
Prognostic significance of tumor immune microenvironment and immunotherapy: Novel insights and future perspectives in gastric cancer.
Immunotherapy for metastatic colorectal cancer: present status and new options.
Targeting cancer-related inflammation in the era of immunotherapy.
Review Article: Immune Landscape and Immunotherapy Options in Cervical Carcinoma.
Cancer Immunotherapy and Personalized Medicine: Emerging Technologies and Biomarker-Based Approaches.
Mechanistic rationales for combining immunotherapy with radiotherapy.
Advances in immunotherapy for prostate cancer.
Research advance of natural products in tumor immunotherapy.
Comparing immunotherapies to other frequently used treatments of gastric cancer.
Checkpoint Inhibitors in Cancer Therapy: Clinical Benefits for Head and Neck Cancers.
Targeting CTCFL/BORIS for the immunotherapy of cancer.
The importance of immune checkpoints in immune monitoring: A future paradigm shift in the treatment of cancer.
Immunoscore and Immunoprofiling in cancer: an update from the melanoma and immunotherapy bridge 2015.
Cancer immune escape: the role of antigen presentation machinery.
Cancer-cell-intrinsic mechanisms shaping the immunosuppressive landscape of prostate cancer.
Immune checkpoint molecules in neuroblastoma: A clinical perspective.
Perspectives in immunotherapy: meeting report from the immunotherapy bridge (December 2nd-3rd, 2020, Italy).
Targeted therapies to improve tumor immunotherapy.
Victory and defeat in the induction of a therapeutic response through vaccine therapy for human and canine brain tumors: a review of the state of the art.
Perspectives on immunotherapy via oncolytic viruses.
Tumor immune microenvironment and nano-immunotherapeutics in colorectal cancer.
Perspectives in melanoma: meeting report from the "Melanoma Bridge" (December 5th-7th, 2019, Naples, Italy).
Intestinal Microbiota: The Driving Force behind Advances in Cancer Immunotherapy.
Vaccines in urologic malignancies.
Cancer immune exclusion: breaking the barricade for a successful immunotherapy.
Immunotherapy for advanced gastric cancer.
Next-generation immunotherapy for pancreatic ductal adenocarcinoma: navigating pathways of immune resistance.
Immunotherapy of malignant diseases. Challenges and strategies.
Targeted Tumor Therapy Remixed-An Update on the Use of Small-Molecule Drugs in Combination Therapies.
TREM2: Keeping Pace With Immune Checkpoint Inhibitors in Cancer Immunotherapy.

Immunotherapy for glioma: from illusion to realistic prospects?
Transcriptional determinants of cancer immunotherapy response and resistance.
Potentiating prostate cancer immunotherapy with oncolytic viruses.
Immune checkpoint inhibitors: The linchpins of modern immunotherapy.
Immunotherapy for gastrointestinal malignancies.
HDAC Inhibition to Prime Immune Checkpoint Inhibitors.
Conference scene: immune effector mechanisms in tumor immunity.
Current Advances in Immunotherapy for Glioblastoma.
Immune Therapy Resistance and Immune Escape of Tumors.
Basic Overview of Current Immunotherapy Approaches in Cancer.
Antigen-specific vaccines for cancer treatment.
Overcoming resistance to targeted therapy with immunotherapy and combination therapy for metastatic melanoma.
Application of nanotechnology in circumventing immunotolerance.
Next generation immune-checkpoints for cancer therapy.
Targeting phagocytosis to enhance antitumor immunity.
Emerging nanotechnological strategies to reshape tumor microenvironment for enhanced therapeutic outcomes of cancer immunotherapy.
Immunotherapeutic Approaches to Biliary Cancer.
Prostate cancer, tumor immunity and a renewed sense of optimism in immunotherapy.
The Role of Antigen Processing and Presentation in Cancer and the Efficacy of Immune Checkpoint Inhibitor Immunotherapy.
Immunological-based approaches for cancer therapy.
The role of CD47-SIRP $\alpha$ immune checkpoint in tumor immune evasion and innate immunotherapy.
Immunovirotherapy: The role of antibody based therapeutics combination with oncolytic viruses.
Immunotherapy against antigenic tumors: a game with a lot of players.
Role of Tumor-Mediated Dendritic Cell Tolerization in Immune Evasion.
Immune Checkpoint Regulators: A New Era Toward Promising Cancer Therapy.
Activated melanoma vessels: A sticky point for successful immunotherapy.
PD-1 and PD-L1 Checkpoint Signaling Inhibition for Cancer Immunotherapy: Mechanism, Combinations, and Clinical Outcome.
Molecular Immunotherapy: Promising Approach to Treat Metastatic Colorectal Cancer by Targeting Resistant Cancer Cells or Cancer Stem Cells.
Overcoming Immune Evasion in Melanoma.
Radiation and Immune Checkpoint Blockade: From Bench to Clinic.
Society for Immunotherapy of Cancer (SITC) consensus definitions for resistance to combinations of immune checkpoint inhibitors with targeted therapies.
Immunotherapy for nasopharyngeal cancer-a review.
Immune Checkpoint Inhibitors and Prostate Cancer: A New Frontier?
The PD-1 pathway as a therapeutic target to overcome immune escape mechanisms in cancer.
Screening and identification of molecular targets for cancer therapy.
Immunotherapy for gastrointestinal cancer: current status and strategies for improving efficacy.
Immunoediting and antigen loss: overcoming the achilles heel of immunotherapy with antigen non-specific therapies.
Can Natural Products be Used to Overcome the Limitations of Colorectal Cancer Immunotherapy?
[Melanoma immunotherapy: dendritic cell vaccines].

[Review on immunotherapies for lung cancer].
Immunoprecipitation therapy, a potential multi-epitope cancer vaccine.
Can calcium signaling be harnessed for cancer immunotherapy?
Lung cancer: a classic example of tumor escape and progression while providing opportunities for immunological intervention.
Emerging Targeted and Immune-Based Therapies in Sarcoma.
[Immune tolerance and active suppression in oncology. Immunological principles and therapeutic options].
Exploring the food-gut axis in immunotherapy response of cancer patients.
Immunoengineering through cancer vaccines - A personalized and multi-step vaccine approach towards precise cancer immunity.
[Neoantigens Are Critical Targets in Naturally and Therapeutically Induced Immune Responses to Cancer].
A novel combination immunotherapy for cancer by IL-13R $\alpha$ 2-targeted DNA vaccine and immunotoxin in murine tumor models.
Strategies for cancer vaccine development.
Author's view: radiation and immunotherapy as systemic therapy for solid tumors.
Epigenetics Regulates Antitumor Immunity in Melanoma.
Current challenges for cancer vaccine adjuvant development.
[Tumor/cytotoxic effector cross-talk in the control of tumor susceptibility to lysis].
A bibliometric and visualized research on global trends of immune checkpoint inhibitors related complications in melanoma, 2011-2021.
Immune Escape in Prostate Cancer: Known and Predicted Mechanisms and Targets.
[Immunotherapy: Activation of a system not a pathway].
Emerging monoclonal antibodies for the treatment of renal cell carcinoma (RCC).
Radiation therapy and immunotherapy: what is the optimal timing or sequencing?
[Epitope peptide vaccine with oncoantigen for cancer and its biomarker].
Preclinical imaging for targeting cancer immune evasion.
Immunotherapy of advanced or metastatic melanoma.
The direct display of costimulatory proteins on tumor cells as a means of vaccination for cancer immunotherapy.
Transcriptional Regulation of Cancer Immune Checkpoints: Emerging Strategies for Immunotherapy.
Interaction of molecular alterations with immune response in melanoma.
Targeted Nanophotoimmunotherapy Potentiates Cancer Treatment by Enhancing Tumor Immunogenicity and Improving the Immunosuppressive Tumor Microenvironment.
Evolving role of tumor antigens for future melanoma therapies.
Improving cancer immunotherapy by targeting tumor-induced immune suppression.
Advances in gene therapy and immunotherapy for brain tumors.
Research progress of nanovaccine in anti-tumor immunotherapy.
Immune checkpoint of B7-H3 in cancer: from immunology to clinical immunotherapy.
Immunotherapy: New Strategies for the Treatment of Gynecologic Malignancies.
[Immune response and cancer].
Translational Research in Cutaneous Melanoma: New Therapeutic Perspectives.
Immune checkpoint inhibitors and combinations with other agents in cholangiocarcinoma.
The Five Immune Forces Impacting DNA-Based Cancer Immunotherapeutic Strategy.

Key Players of the Immunosuppressive Tumor Microenvironment and Emerging Therapeutic Strategies.
RhoC a new target for therapeutic vaccination against metastatic cancer.
4th Summer School in Immuno-Oncology, July 1st-3rd, 2021, Athens, Greece.
Active specific immunotherapy for colorectal cancer.
Bispecific Antibodies: A Novel Approach for the Treatment of Solid Tumors.
Immunotherapy for Hepatocellular Carcinoma: Current Advances and Future Expectations.
Proteomics to study cancer immunity and improve treatment.
Twelve immunotherapy drugs that could cure cancers.
Inflammatory Markers in Cancer Immunotherapy.
The rationale for including immune checkpoint inhibition into multimodal primary treatment concepts of head and neck cancer.
Impact of Diets on Response to Immune Checkpoint Inhibitors (ICIs) Therapy against Tumors.
The need of radiotherapy optimization for glioblastomas considering immune responses.
Genomics and emerging biomarkers for immunotherapy of colorectal cancer.
Combined therapy for gastrointestinal carcinomas: exploiting synergies between gene therapy and classical chemo-radiotherapy.
Mechanism of action of immunotherapy.
Mathematical Prostate Cancer Evolution: Effect of Immunotherapy Based on Controlled Vaccination Strategy.
Interplay between the DNA Damage Response and Immunotherapy Response in Cancer.
Targeting inhibitory pathways in cancer immunotherapy.
Immunotherapy in Colorectal Cancer.
Focus on TILs: Prognostic significance of tumor infiltrating lymphocytes in human bladder cancer.
Immunotherapy for gastric cancers: emerging role and future perspectives.
A comparison of cancer vaccine adjuvants in clinical trials.
Developments in cancer vaccination.
Regulation of cell death in cancer-possible implications for immunotherapy.
The mechanism and clinical application of DNA damage repair inhibitors combined with immune checkpoint inhibitors in the treatment of urologic cancer.
Advances in personalized neoantigen vaccines for cancer immunotherapy.
Targets of immunotherapy for hepatocellular carcinoma: An update.
Society for Immunotherapy of Cancer (SITC) consensus definitions for resistance to combinations of immune checkpoint inhibitors.
Modern Approaches to Cancer Therapy: Immunotherapy and Targeted Drug Delivery.
Developing recombinant and synthetic vaccines for the treatment of melanoma.
Implications of corticotropin releasing factor in targeted anticancer therapy.
A MD Simulation Prediction for Regulation of N-Terminal Modification on Binding of CD47 to CD172a in a Force-Dependent Manner.
Immune evasion of microsatellite unstable colorectal cancers.
Romiplostim for the treatment of primary immune thrombocytopenia.
The confluence of stereotactic ablative radiotherapy and tumor immunology.
Identification of novel survivin-derived CTL epitopes.
Targeting telomerase reverse transcriptase with the covalent inhibitor NU-1 confers immunogenic radiation sensitization.

miR-2909 regulates ISGylation system via STAT1 signalling through negative regulation of SOCS3 in prostate cancer.
Androgen receptor agonist and antagonist reduce response of cytokine-induced killer cells on prostate cancer cells.
Management of higher risk myelodysplastic syndromes after hypomethylating agents failure: are we about to exit the black hole?
Thrombin contributes to cancer immune evasion via proteolysis of platelet-bound GARP to activate LTGF- $\beta$ .
Meeting report: The 13th Annual Meeting of the Translational Research Cancer Centers Consortium (TrC3); Immune Suppression and the Tumor Microenvironment, Columbus, Ohio; March 1-2, 2010.
Gene and cell therapies for diabetes mellitus: strategies and clinical potential.
All is not lost: learning from 9p21 loss in cancer.
Treating Hepatobiliary Cancer: The Immunologic Approach.
Ubiquitin Ligases in Cancer Immunotherapy - Balancing Antitumor and Autoimmunity.
Leukoplakia and Immunology: New Chemoprevention Landscapes?
Oxidative Stress and Immune Response in Melanoma: Ion Channels as Targets of Therapy.
The emerging roles of HDACs and their therapeutic implications in cancer.
Selective FGFR/FGF pathway inhibitors: inhibition strategies, clinical activities, resistance mutations, and future directions.
Human chorionic gonadotropin as a target for cancer vaccines.
Epigenetic escape of immunosurveillance by malignant cell precursors.
The roles of E3 ubiquitin ligases in cancer progression and targeted therapy.
Synergistic Anticancer Effect of a Combination of Berbamine and Arcyriaflavin A against Glioblastoma Stem-like Cells.
Models, mechanisms and clinical evidence for cancer dormancy.
Immunological off-target effects of imatinib.
Advances in Hypofractionated Irradiation-Induced Immunosuppression of Tumor Microenvironment.
Optimal organization of a polypeptide-based candidate cancer vaccine composed of cryptic tumor peptides with enhanced immunogenicity.
High-content CRISPR screening in tumor immunology.
Anti-leukemic activity and tolerability of anti-human CD47 monoclonal antibodies.
Chromosome instability and aneuploidy as context-dependent activators or inhibitors of antitumor immunity.
Heterotypic cell-in-cell structures between cancer and NK cells are associated with enhanced anticancer drug resistance.
Genetic and pharmacological modulation of DNA mismatch repair heterogeneous tumors promotes immune surveillance.
Evolutionary dynamics of neoantigens in growing tumors.
Restoring antigen presentation: the first step on the road to complete immune restoration.
High-Throughput CRISPR Screening in Hematological Neoplasms.
The Double Engines and Single Checkpoint Theory of Endometriosis.
A comparative genomics approach to understanding transmissible cancer in Tasmanian devils.
Two is better than one: Complementing oncolytic virotherapy with gemcitabine to potentiate antitumor immune responses.
Cyclin-Dependent Kinase 8: A New Hope in Targeted Cancer Therapy?
Tumor immunology meets oncology (TIMO) XVI, July 07-09 2022 in Halle/Saale, Germany.

The imitation game: How glioblastoma outmaneuvers immune attack.
Targeting Heat-Shock Protein 90 in Cancer: An Update on Combination Therapy.
Cancer as an evolutionary and ecological process.
Liver antigen-presenting cells.
Her-2/neu as a paradigm of a tumor-specific target for therapy.
Escape from nonsense-mediated decay associates with anti-tumor immunogenicity.
Glutathione depletion and carbon ion radiation potentiate clustered DNA lesions, cell death and prevent chromosomal changes in cancer cells progeny.
The KDM5B and KDM1A lysine demethylases cooperate in regulating androgen receptor expression and signalling in prostate cancer.
Clonal evolution and immune evasion in posttransplantation relapses.
Tumor antigen CA125 suppresses antibody-dependent cellular cytotoxicity (ADCC) via direct antibody binding and suppressed Fc-γ receptor engagement.
Trick and treat: Ambushing recurrent tumors before they get out of the door.
Novel therapies for malignant pleural effusion: Anti-angiogenic therapy and immunotherapy (Review).
From immunotoxicity to carcinogenicity: the effects of carbamate pesticides on the immune system.
The transcription factor NRF2 enhances melanoma malignancy by blocking differentiation and inducing COX2 expression.
Innovative Approaches in the Battle Against Cancer Recurrence: Novel Strategies to Combat Dormant Disseminated Tumor Cells.
Emerging Roles of ALK in Immunity and Insights for Immunotherapy.
The Diversification of Cell Death and Immunity: Memento Mori.
The JAK2/STAT3 inhibitor pacritinib effectively inhibits patient-derived GBM brain tumor initiating cells in vitro and when used in combination with temozolomide increases survival in an orthotopic xenograft model.
LSD1 inhibition induces differentiation and cell death in Merkel cell carcinoma.
Barriers in Heart Failure Gene Therapy and Approaches to Overcome Them.
"Tumor immunology meets oncology (TIMO) XV", April 25th-27th 2019, Halle/Saale, Germany.
Modelling the immunosuppressive effect of liver SBRT by simulating the dose to circulating lymphocytes: an in-silico planning study.
Placenta immune infiltrates and perinatal outcomes.
Immunobiology of Merkel cell carcinoma: implications for immunotherapy of a polyomavirus-associated cancer.
Melanogenesis inhibits respiration in B16-F10 melanoma cells whereas enhances mitochondrial cell content.
Passive immunotherapeutic strategies for the treatment of malignant gliomas.
The use of doxorubicine at low doses for elevation of LAK-activity toward explants and cells of MC-rhabdomyosarcoma and B16 melanoma resistant to doxorubicin.
Targeting MRP4 expression by anti-androgen treatment reverses MRP4-mediated docetaxel resistance in castration-resistant prostate cancer.
Endoplasmic reticulum Ca(2+) content decrease by PKA-dependent hyperphosphorylation of type 1 IP3 receptor contributes to prostate cancer cell resistance to androgen deprivation.
Oncogene abnormalities in a series of primary melanomas of the sinonasal tract: NRAS mutations and cyclin D1 amplification are more frequent than KIT or BRAF mutations.
Single-Cell Ssequencing in Cancer: Recent Applications to Immunogenomics and Multi-omics Tools.
The unfolded protein response and the biology of uveal melanoma.

Targeting Glioma Stem Cells.
Regenerative medicine: prospects for the treatment of inflammatory bowel disease.
Perioperative Enteral Immunonutrition Modulates Systemic and Mucosal Immunity and the Inflammatory Response in Patients With Periapillary Cancer Scheduled for Pancreaticoduodenectomy: A Randomized Clinical Trial.
Corrigendum: The TNF Paradox in Cancer Progression and Immunotherapy.
HO-1 Limits the Efficacy of Vemurafenib/PLX4032 in BRAF(V600E) Mutated Melanoma Cells Adapted to Physiological Normoxia or Hypoxia.
Recent advances in small molecule and peptide inhibitors of glucose-regulated protein 78 for cancer therapy.
Revisiting the tolerogenicity of epidermal Langerhans cells.
CD200 expression in patients with Multiple Myeloma: another piece of the puzzle.
Discovery of a new class of reversible TEA domain transcription factor inhibitors with a novel binding mode.
Immunologically silent cancer clone transmission from mother to offspring.
Towards single cell encapsulation for precision biology and medicine.
The immunoregulatory function of polyphenols: implications in cancer immunity.
Clonal evolution in aplastic anemia: failed tumor surveillance or maladaptive recovery?
Tissue Engineering for Mimics and Modulations of Immune Functions.
YAP/TAZ Signaling and Resistance to Cancer Therapy.
[Immunological effects of silica/asbestos].
Pediatric osteosarcoma - treatment of the primary tumor with intravenous cis-diamminedichloroplatinum-ii (cdp) - comparison of the results with the reported efficacy of intraarterial cdp.
The concurrence of multiple sclerosis and glioblastoma.
Xenotransplantation of Microencapsulated Parathyroid Cells as a Potential Treatment for Autoimmune-Related Hypoparathyroidism.
Timed GDNF gene therapy using an immune-evasive gene switch promotes long distance axon regeneration.
Novel insights into the synergistic interaction of Bortezomib and TRAIL: tBid provides the link.
Devil Facial Tumor Disease.
A 'devil' of a problem.
Immunological aspects of blood transfusions.
Microencapsulated parathyroid allotransplantation in the omental tissue.
Partial Decellularization for Segmental Tracheal Scaffold Tissue Engineering: A Preliminary Study in Rabbits.
Immunology of normal pregnancy.

Table A2-61, Cluster 60

Cluster 60 focuses on viruses, emphasizing viral strategies for immune system evasion (1244)
Innate antiviral immune signaling, viral evasion and modulation by HIV-1.
[How the bovine viral diarrhea virus outwits the immune system].
An Overview of Immune Evasion Strategies of DNA and RNA Viruses.
High-Content Screening and Computational Prediction Reveal Viral Genes That Suppress the Innate Immune Response.



How do viruses control mitochondria-mediated apoptosis?
Diverse mechanisms evolved by DNA viruses to inhibit early host defenses.
Evasion of influenza A viruses from innate and adaptive immune responses.
Non-cytotoxic antiviral activities of granzymes in the context of the immune antiviral state.
Cytotoxic T-lymphocyte escape viral variants: how important are they in viral evasion of immune clearance in vivo?
Innate Immune Evasion by Human Respiratory RNA Viruses.
Cell cycle regulation during viral infection.
Subversion and piracy: DNA viruses and immune evasion.
Replication of many human viruses is refractory to inhibition by endogenous cellular microRNAs.
Lactate dehydrogenase-elevating virus: an ideal persistent virus?
Innate Sensing of DNA Virus Genomes.
Tricks and threats of RNA viruses - towards understanding the fate of viral RNA.
HIV Exploits Antiviral Host Innate GCN2-ATF4 Signaling for Establishing Viral Replication Early in Infection.
Influenza A virus-induced degradation of eukaryotic translation initiation factor 4B contributes to viral replication by suppressing IFITM3 protein expression.
RNA viruses and microRNAs: challenging discoveries for the 21st century.
Modulation of innate immune signalling pathways by viral proteins.
NS1 Protein Mutation I64T Affects Interferon Responses and Virulence of Circulating H3N2 Human Influenza A Viruses.
Viruses, microRNAs and cancer.
Comparative interactomics for virus-human protein-protein interactions: DNA viruses versus RNA viruses.
Molecular Mechanisms of Innate Immune Inhibition by Non-Segmented Negative-Sense RNA Viruses.
Acute Infection of Viral Pathogens and Their Innate Immune Escape.
RNA-virus proteases counteracting host innate immunity.
The viral manipulation of the host cellular and immune environments to enhance propagation and survival: a focus on RNA viruses.
Immunomodulation by viruses: the myxoma virus story.
Host-virus interaction and viral evasion.
To sense or not to sense viral RNA--essentials of coronavirus innate immune evasion.
[Viral infections].
Induction of innate immunity and its perturbation by influenza viruses.
Induction and Evasion of Type-I Interferon Responses during Influenza A Virus Infection.
Viral infection: an evolving insight into the signal transduction pathways responsible for the innate immune response.
Regulation of antiviral innate immune signaling and viral evasion following viral genome sensing.
Cytosolic and nuclear recognition of virus and viral evasion.
Host and viral determinants of influenza A virus species specificity.
Viral MicroRNAs: Interfering the Interferon Signaling.
Immune regulation and evasion of Mammalian host cell immunity during viral infection.
The role of autophagy in viral infections.
When Poly(A) Binding Proteins Meet Viral Infections, Including SARS-CoV-2.
Viral pathogenesis, modulation of immune receptor signaling and treatment.
The Diverse Roles of microRNAs at the Host-Virus Interface.

Viral Infection of Human Natural Killer Cells.
Viral Innate Immune Evasion and the Pathogenesis of Emerging RNA Virus Infections.
Tiny Regulators in Viral Infection: Carving SARS-CoV-2 by miRNAs.
Vaccinia virus immune evasion: mechanisms, virulence and immunogenicity.
Viral CD229 (Ly9) homologs as new manipulators of host immunity.
Virus-Encoded Complement Regulators: Current Status.
The Importance of Glycans of Viral and Host Proteins in Enveloped Virus Infection.
Evasion and disruption of innate immune signalling by hepatitis C and West Nile viruses.
First Impressions-the Potential of Altering Initial Host-Virus Interactions for Rational Design of Herpesvirus Vaccine Vectors.
Amino acid substitutions in norovirus VP1 dictate cell tropism via an attachment process dependent on membrane mobility.
Viral deubiquitinases and innate antiviral immune response in livestock and poultry.
Recent insights into the role of Toll-like receptors in viral infection.
Control of human viral infections by natural killer cells.
Regulation of Antiviral Immune Response by N (6)-Methyladenosine of mRNA.
Viral strategies to subvert the mammalian translation machinery.
Immune evasion during foot-and-mouth disease virus infection of swine.
Recurrent evolution of high virulence in isolated populations of a DNA virus.
[Virokines and viroceptors--viral immunomodulators with clinical and therapeutic implications].
Pellino Proteins in Viral Immunity and Pathogenesis.
The modulation of apoptosis by oncogenic viruses.
The SCHOOL of nature: IV. Learning from viruses.
Intracellular sensing of viral genomes and viral evasion.
The Role of the Host Ubiquitin System in Promoting Replication of Emergent Viruses.
Common Nodes of Virus-Host Interaction Revealed Through an Integrated Network Analysis.
Does enterovirus 71 urge for effective vaccine control strategies? Challenges and current opinion.
Viral Evasion of the Complement System and Its Importance for Vaccines and Therapeutics.
Discovery and characterization of a novel alphavirus-like RNA virus from the red firebug <i>Pyrrhocoris apterus</i> L. (Heteroptera).
Incoming influenza A virus evades early host recognition, while influenza B virus induces interferon expression directly upon entry.
Modulation of Innate Immune Responses by the Influenza A NS1 and PA-X Proteins.
Cellular versus viral microRNAs in host-virus interaction.
A host susceptibility gene, DR1, facilitates influenza A virus replication by suppressing host innate immunity and enhancing viral RNA replication.
Monocytes and Macrophages as Viral Targets and Reservoirs.
Viral evasion mechanisms of early antiviral responses involving regulation of ubiquitin pathways.
Evolution of transient RNA structure-RNA polymerase interactions in respiratory RNA virus genomes.
Innate immune evasion strategies of influenza viruses.
Systems biology unravels interferon responses to respiratory virus infections.
Subcellular trafficking in rhabdovirus infection and immune evasion: a novel target for therapeutics.
Evasion of the Innate Immune Type I Interferon System by Monkeypox Virus.
Induction and Suppression of NF- $\kappa$ B Signalling by a DNA Virus of <i>Drosophila</i> .
Host Defenses to Viruses: Lessons from Inborn Errors of Immunity.
Viral miRNAs and immune evasion.

Innate barriers to viral infection.
Viral persistence: HIV's strategies of immune system evasion.
The Evasion of Antiviral Innate Immunity by Chicken DNA Viruses.
Host detection and the stealthy phenotype in influenza virus infection.
Mechanisms underlying the inhibition of interferon signaling by viruses.
Epigenetic modulation of host: new insights into immune evasion by viruses.
Genetic 'budget' of viruses and the cost to the infected host: a theory on the relationship between the genetic capacity of viruses, immune evasion, persistence and disease.
Elucidation of the Ebola virus VP24 cellular interactome and disruption of virus biology through targeted inhibition of host-cell protein function.
Viral mimicry of cytokines, chemokines and their receptors.
Safeguard against DNA sensing: the role of TREX1 in HIV-1 infection and autoimmune diseases.
Special Issue: "Innate Immune Sensing of Viruses and Viral Evasion".
Pathogenic Roles of m6A Modification in Viral Infection and Virus-driven Carcinogenesis.
Immunotherapy of viral infections.
Viral proteins targeting host protein kinase R to evade an innate immune response: a mini review.
Influenza virus directly infects human natural killer cells and induces cell apoptosis.
Mechanisms of viral clearance and persistence.
Post-translational Modification-Based Regulation of HIV Replication.
The Interplay Between Viral-Derived miRNAs and Host Immunity During Infection.
The Wnt pathway: a key network in cell signalling dysregulated by viruses.
Extracellular Vesicles in Viral Spread and Antiviral Response.
Vaccinia virus immune evasion.
When Dendritic Cells Go Viral: The Role of Siglec-1 in Host Defense and Dissemination of Enveloped Viruses.
The Promise of Proteomics in the Study of Oncogenic Viruses.
Modulation of natural killer cell activity by viruses.
Evasion of innate and adaptive immune responses by influenza A virus.
microRNA-21: a key modulator in oncogenic viral infections.
The domain landscape of virus-host interactomes.
Host Innate Immune Response and Viral Immune Evasion During Alphaherpesvirus Infection.
A Comprehensive Insight into the Role of Exosomes in Viral Infection: Dual Faces Bearing Different Functions.
Dicer-2 processes diverse viral RNA species.
Respiratory viral infections in infants: causes, clinical symptoms, virology, and immunology.
[Ebola hemorrhagic fever: Properties of the pathogen and development of vaccines and chemotherapeutic agents].
Exploitation of glycosylation in enveloped virus pathobiology.
Abscisic Acid Connects Phytohormone Signaling with RNA Metabolic Pathways and Promotes an Antiviral Response that Is Evaded by a Self-Controlled RNA Virus.
Innate immune sensing of coronavirus and viral evasion strategies.
HIV-1 Envelope Glycoprotein at the Interface of Host Restriction and Virus Evasion.
Host Immune Response to Influenza A Virus Infection.
Viral control of mitochondrial apoptosis.
Genome guardian p53 and viral infections.
Influenza Virus Infections and Cellular Kinases.

Sensor Sensibility-HIV-1 and the Innate Immune Response.
Host Defence RNases as Antiviral Agents against Enveloped Single Stranded RNA Viruses.
Modulation of chemokine activity by viruses.
Conserved charged amino acids within Sendai virus C protein play multiple roles in the evasion of innate immune responses.
[Viral interactions with the host's immune system].
Apoptosis prevention as a mechanism of immune evasion.
Clearance or Hijack: Universal Interplay Mechanisms Between Viruses and Host Autophagy From Plants to Animals.
Tumour viruses and innate immunity.
Coinfection by influenza A virus and respiratory syncytial virus produces hybrid virus particles.
Viruses and miRNAs: More Friends than Foes.
Evasion of short interfering RNA-directed antiviral silencing in <i>Musa acuminata</i> persistently infected with six distinct banana streak pararetroviruses.
Influenza Virus: A Master Tactician in Innate Immune Evasion and Novel Therapeutic Interventions.
MicroRNA and Nonsense Transcripts as Putative Viral Evasion Mechanisms.
[Modulation of the host toll-like receptor signaling pathways by virus infection].
An Overview of the Infectious Cycle of Bunyaviruses.
Synthesizing within-host and population-level selective pressures on viral populations: the impact of adaptive immunity on viral immune escape.
Viral miRNA regulation of host gene expression.
The Role of Viruses in Carcinogenesis and Molecular Targeting: From Infection to Being a Component of the Tumor Microenvironment.
Virus-cell interactions: impact on cytokine production, immune evasion and tumor growth.
Caspase cleavage of viral proteins, another way for viruses to make the best of apoptosis.
Viral deubiquitinases: role in evasion of anti-viral innate immunity.
Quantitative analysis of cellular proteome alterations in human influenza A virus-infected mammalian cell lines.
Antisense approaches for elucidating ranavirus gene function in an infected fish cell line.
A proteomics perspective on viral DNA sensors in host defense and viral immune evasion mechanisms.
Host Innate Immunity Against Hepatitis Viruses and Viral Immune Evasion.
Contemplating the Importance of Toll-like Receptors I and II Regarding Human Viral Pathogenesis.
Filovirus pathogenesis and immune evasion: insights from Ebola virus and Marburg virus.
Autoimmune diseases initiated by pathogen infection: Mathematical modeling.
Phosphorylation of JIP4 at S730 Presents Antiviral Properties against Influenza A Virus Infection.
Virus infections: escape, resistance, and counterattack.
The RNA helicase DDX5 promotes viral infection via regulating N6-methyladenosine levels on the DHX58 and NFkB transcripts to dampen antiviral innate immunity.
Identification of Myricetin as an Ebola Virus VP35-Double-Stranded RNA Interaction Inhibitor through a Novel Fluorescence-Based Assay.
How vaccinia virus has evolved to subvert the host immune response.
A Map of the Arenavirus Nucleoprotein-Host Protein Interactome Reveals that Junín Virus Selectively Impairs the Antiviral Activity of Double-Stranded RNA-Activated Protein Kinase (PKR).
Influenza virus polymerase: Functions on host range, inhibition of cellular response to infection and pathogenicity.

Exosomes in Viral Disease.
Mouse adaptation of the H9N2 avian influenza virus causes the downregulation of genes related to innate immune responses and ubiquitin-mediated proteolysis in mice.
Virus-specific host miRNAs: antiviral defenses or promoters of persistent infection?
Respiratory syncytial virus infection of airway cells: Role of microRNAs.
RNA Viruses and RNAi: Quasispecies Implications for Viral Escape.
Defective viral genomes are key drivers of the virus-host interaction.
Dual role of glycans and binding receptors in pathogenesis of enveloped viruses (by mainly focusing on two recent pandemics).
The Dynamic Interface of Viruses with STATs.
Vaccinia virus immune evasion.
Viral appropriation of apoptotic and NF-kappaB signaling pathways.
Novel viral translation strategies.
Model of influenza A virus infection: dynamics of viral antagonism and innate immune response.
SAFA facilitates chromatin opening of immune genes through interacting with anti-viral host RNAs.
Innate immune evasion strategies by human immunodeficiency virus type 1.
Human immunodeficiency virus-1 (HIV-1)-mediated apoptosis: new therapeutic targets.
[Exosomes: another arena for the game between viruses and hosts].
Mechanisms of innate immune evasion in re-emerging RNA viruses.
New concepts in measles virus replication: getting in and out in vivo and modulating the host cell environment.
Infections persistantes à entérovirus et pathologies humaines.
Immune evasion in ebolavirus infections.
SV40 miR-S1 and Cellular miR-1266 Sequester Each Other from Their Targets, Enhancing Telomerase Activity and Viral Replication.
Skin Viral Infections: Host Antiviral Innate Immunity and Viral Immune Evasion.
Picornavirus infection leading to immunosuppression.
Immunity and Viral Infections: Modulating Antiviral Response via CRISPR-Cas Systems.
Viral immune evasion strategies and the underlying cell biology.
MicroRNAs as mediators of viral evasion of the immune system.
Structure and Function of Viral Deubiquitinating Enzymes.
Vesicular Stomatitis Virus Phosphoprotein Dimerization Domain Is Dispensable for Virus Growth.
Assessing the Structure and Function of Vaccinia Virus Gene Products by Transient Complementation.
Infection Defects of RNA and DNA Viruses Induced by Antiviral RNA Interference.
Innate immune sensing of retroviral infection via Toll-like receptor 7 occurs upon viral entry.
Evasion of innate immunity by vaccinia virus.
Interplay between the virus and the ubiquitin-proteasome system: molecular mechanism of viral pathogenesis.
Plant miRNAome and antiviral resistance: a retrospective view and prospective challenges.
Subversion of cytokine networks by virally encoded decoy receptors.
Evasion of host immune responses by tumours and viruses.
RNA viruses can hijack vertebrate microRNAs to suppress innate immunity.
Viral mechanisms of immune evasion.
Pestiviruses: how to outmaneuver your hosts.
Emerging Roles of Cyclophilin A in Regulating Viral Cloaking.
The genome of swinepox virus.

Interactions between Autophagy and DNA Viruses.
Exploiting the therapeutic potential of microRNAs in viral diseases: expectations and limitations.
Plagiarism of the host immune system: lessons about chemokine immunology from viruses.
The biology of influenza viruses.
Ribosome Profiling Reveals Translational Upregulation of Cellular Oxidative Phosphorylation mRNAs during Vaccinia Virus-Induced Host Shutoff.
How Human Papillomavirus Replication and Immune Evasion Strategies Take Advantage of the Host DNA Damage Repair Machinery.
Virus complement evasion strategies.
Viral and cellular oncogenes promote immune evasion.
Short-Sighted Virus Evolution and a Germline Hypothesis for Chronic Viral Infections.
Specific phenotypic restoration of an attenuated virus by knockout of a host resistance gene.
Virus encoded cytokines and cytokine receptors.
Modulation of host immune responses by clinically relevant human DNA and RNA viruses.
Viruses and human cancers: a long road of discovery of molecular paradigms.
The imitation game: a viral strategy to subvert the complement system.
Rational design of a flavivirus vaccine by abolishing viral RNA 2'-O methylation.
Poxviral Strategies to Overcome Host Cell Apoptosis.
Enhanced Replication of Mouse Adenovirus Type 1 following Virus-Induced Degradation of Protein Kinase R (PKR).
ARHGAP1 Transported with Influenza Viral Genome Ensures Integrity of Viral Particle Surface through Efficient Budding Formation.
Characterization of host immune responses in Ebola virus infections.
RNA Virus-Encoded miRNAs: Current Insights and Future Challenges.
ADAR2 Is Involved in Self and Nonself Recognition of Borna Disease Virus Genomic RNA in the Nucleus.
Global Foot-and-Mouth Disease Research Update and Gap Analysis: 7 - Pathogenesis and Molecular Biology.
Towards an understanding of the poliovirus replication complex: the solution structure of the soluble domain of the poliovirus 3A protein.
Immunology 101 at poxvirus U: immune evasion genes.
A skeptical look at viral immune evasion.
Viral infection: Moving through complex and dynamic cell-membrane structures.
Identification of Poxvirus Genome Uncoating and DNA Replication Factors with Mutually Redundant Roles.
Roles of nuclear trafficking in infection by cytoplasmic negative-strand RNA viruses: paramyxoviruses and beyond.
Inhibition of human natural killer cell activity by influenza virions and hemagglutinin.
Proteomics approaches for the identification of protease substrates during virus infection.
Human adenovirus infections: update and consideration of mechanisms of viral persistence.
Unravelling the Immunomodulatory Effects of Viral Ion Channels, towards the Treatment of Disease.
Immunological Perspective for Ebola Virus Infection and Various Treatment Measures Taken to Fight the Disease.
Physical and Functional Analysis of Viral RNA Genomes by SHAPE.
Optimal control strategy for abnormal innate immune response.
Synthesis and quality control of viral membrane proteins.

Emerging Infections of CNS: Avian Influenza A Virus, Rift Valley Fever Virus and Human Parechovirus.
Killer Archaea: Virus-Mediated Antagonism to CRISPR-Immune Populations Results in Emergent Virus-Host Mutualism.
Cellular sensing of viral DNA and viral evasion mechanisms.
Dissecting innate immune signaling in viral evasion of cytokine production.
Manipulation of viral infection by deubiquitinating enzymes: new players in host-virus interactions.
Human Adenoviruses, Cholesterol Trafficking, and NF- $\kappa$ B Signaling.
Infection of dendritic cells by lymphocytic choriomeningitis virus.
The role of cholesterol 25-hydroxylase in viral infections: Mechanisms and implications.
Influenza A virus attenuation by codon deoptimization of the NS gene for vaccine development.
Antiviral defense in salmonids - Mission made possible?
Subversion of host defense mechanisms by adenoviruses.
Implications of Nef: host cell interactions in viral persistence and progression to AIDS.
Modulation of mitochondria by viral proteins.
Current and future developments in the treatment of virus-induced hypercytokinemia.
Viral Interactions with Adaptor-Protein Complexes: A Ubiquitous Trait among Viral Species.
On revealing the gene targets of Ebola virus microRNAs involved in the human skin microbiome.
Common Mechanisms of Viral Injury to the Kidney.
Mechanisms of RNA interference in the HIV-1-host cell interplay.
The Dual Role of Exosomes in Hepatitis A and C Virus Transmission and Viral Immune Activation.
A Next-Generation Sequencing Approach Uncovers Viral Transcripts Incorporated in Poxvirus Virions.
Viral hijacking of the host ubiquitin system to evade interferon responses.
Mechanism of herpesvirus protein kinase UL13 in immune escape and viral replication.
Innate immunity against HIV-1 infection.
Immune regulation of viral infection and vice versa.
Role of Virally-Encoded Deubiquitinating Enzymes in Regulation of the Virus Life Cycle.
Induction and Subversion of Human Protective Immunity: Contrasting Influenza and Respiratory Syncytial Virus.
Intercellular communication for innate immunity.
Gestational Viral Infections: Role of Host Immune System.
Virus usurps alternative splicing to clear the decks for infection.
Global aspects of viral glycosylation.
Exchange of Genetic Sequences Between Viruses and Hosts.
Host Factors Involved in Ebola Virus Replication.
Role of cytokines in poxvirus host tropism and adaptation.
Translational control during poxvirus infection.
The essential role of mitochondrial dynamics in antiviral immunity.
Viral Nucleases from Herpesviruses and Coronavirus in Recombination and Proofreading: Potential Targets for Antiviral Drug Discovery.
Arbovirus Infections As Screening Tools for the Identification of Viral Immunomodulators and Host Antiviral Factors.
Immune evasion by adenoviruses.
Nef performance in macrophages: the master orchestrator of viral persistence and spread.
VExD: A curated resource for human gene expression alterations following viral infection.
Activation of the Antiviral Kinase PKR and Viral Countermeasures.
Modulation of influenza A virus replication by microRNA-9 through targeting MCP1P1.

Innate immune viral recognition: relevance to CNS infections.
IL-10 encoded by viruses: a remarkable example of independent acquisition of a cellular gene by viruses and its subsequent evolution in the viral genome.
Tumor necrosis factor inhibitors from poxviruses with an emphasis on tanapoxvirus-2L protein.
Race between virus and inflammasomes: inhibition or escape, intervention and therapy.
Interferon Control of Human Coronavirus Infection and Viral Evasion: Mechanistic Insights and Implications for Antiviral Drug and Vaccine Development.
Sindbis Virus Can Exploit a Host Antiviral Protein To Evade Immune Surveillance.
Drawing on disorder: How viruses use histone mimicry to their advantage.
Micro-Players of Great Significance-Host microRNA Signature in Viral Infections in Humans and Animals.
Dendritic cells harbor infectious porcine circovirus type 2 in the absence of apparent cell modulation or replication of the virus.
A FACT-ETS-1 Antiviral Response Pathway Restricts Viral Replication and is Countered by Poxvirus A51R Proteins.
Visualizing Viral Infection In Vivo by Multi-Photon Intravital Microscopy.
Review of Mammarenavirus Biology and Replication.
Myeloid C-Type Lectin Receptors in Viral Recognition and Antiviral Immunity.
viral silencing suppressors: Tools forged to fine-tune host-pathogen coexistence.
Natural killer cell dysfunction during acute infection with foot-and-mouth disease virus.
MicroRNAs, HIV and HCV: a complex relation towards pathology.
Recent progress on gene silencing/suppression by virus-derived small interfering RNAs in rice viruses especially Rice grassy stunt virus.
RNA Polymerase Mutations Selected during Experimental Evolution Enhance Replication of a Hybrid Vaccinia Virus with an Intermediate Transcription Factor Subunit Replaced by the Myxoma Virus Ortholog.
Cellular Targets of HIV-1 Protease: Just the Tip of the Iceberg?
Battle Royale: Innate Recognition of Poxviruses and Viral Immune Evasion.
Modulation of Macrophage Polarization by Viruses: Turning Off/On Host Antiviral Responses.
Avian influenza virus directly infects human natural killer cells and inhibits cell activity.
ZBP1: Innate Sensor Regulating Cell Death and Inflammation.
Expression Profile of Glossina pallidipes MicroRNAs During Symptomatic and Asymptomatic Infection With Glossina pallidipes Salivary Gland Hypertrophy Virus (Hytrosavirus).
Marburg virus VP35 can both fully coat the backbone and cap the ends of dsRNA for interferon antagonism.
Current research for a vaccine against Lassa hemorrhagic fever virus.
2'-O-Methylation of the second transcribed nucleotide within the mRNA 5' cap impacts the protein production level in a cell-specific manner and contributes to RNA immune evasion.
The role of Rel/NF-kappa B proteins in viral oncogenesis and the regulation of viral transcription.
Viral interference with functions of the cellular receptor tyrosine phosphatase CD45.
A cryptic pocket in Ebola VP35 allosterically controls RNA binding.
A unique secreted adenovirus E3 protein binds to the leukocyte common antigen CD45 and modulates leukocyte functions.
Evasion of antiviral innate immunity by Theiler's virus L* protein through direct inhibition of RNase L.
Ubiquitin and Not Only Unfolded Domains Drives Toscana Virus Non-Structural NSs Protein Degradation.



Poxvirus-encoded TNF decoy receptors inhibit the biological activity of transmembrane TNF.
Heparan sulfate binding by natural eastern equine encephalitis viruses promotes neurovirulence.
Increased bovine Tim-3 and its ligand expressions during bovine leukemia virus infection.
Transcriptome analysis of rainbow trout in response to non-virion (NV) protein of viral haemorrhagic septicaemia virus (VHSV).
Genome characterization of cetaceanpox virus from a managed Indo-Pacific bottlenose dolphin ( <i>Tursiops aduncus</i> ).
Codon Usage Bias Analysis of Bluetongue Virus Causing Livestock Infection.
Immune escape phenomenon in molluscum contagiosum and the induction of apoptosis.
IFN-Lambda: The Key to Norovirus's Secret Hideaway.
Modified-Live Feline Calicivirus Vaccination Reduces Viral RNA Loads, Duration of RNAemia, and the Severity of Clinical Signs after Heterologous Feline Calicivirus Challenge.
COVID-19 - Considerations for the paediatric rheumatologist.
Cell-to-Cell Spread of Retroviruses.
Comparative proteomic analysis of PK15 swine kidney cells infected with a pseudorabies pathogenic variant and the Bartha-K/61 vaccine strain.
Conserved Fever Pathways across Vertebrates: A Herpesvirus Expressed Decoy TNF- $\alpha$ Receptor Delays Behavioral Fever in Fish.
Release of Immunomodulatory Ebola Virus Glycoprotein-Containing Microvesicles Is Suppressed by Tetherin in a Species-Specific Manner.
The role of sialic acid-binding immunoglobulin-like-lectin-1 (siglec-1) in immunology and infectious disease.
Genome-Wide Transcriptional Analysis Reveals Novel AhR Targets That Regulate Dendritic Cell Function during Influenza A Virus Infection.
Analyses of HTLV-1 sequences suggest interaction between ORF-I mutations and HAM/TSP outcome.
Intrinsic and extrinsic negative regulators of nuclear protein transport processes.
Glycoprotein G from pseudorabies virus binds to chemokines with high affinity and inhibits their function.
Hantavirus immunology of rodent reservoirs: current status and future directions.
Intercellular transmission of Seneca Valley virus mediated by exosomes.
Forcing Seasonality of Influenza-like Epidemics with Daily Solar Resonance.
Coxsackievirus B Tailors the Unfolded Protein Response to Favour Viral Amplification in Pancreatic $\beta$ Cells.
The proto-oncogene Bcl3, induced by Tax, represses Tax-mediated transcription via p300 displacement from the human T-cell leukemia virus type 1 promoter.
Frequent cross-species transmissions of foamy virus between domestic and wild felids.
Topical Application of Virus-Derived Immunomodulating Proteins and Peptides to Promote Wound Healing in Mouse Models.
Nucleocytoplasmic trafficking of Nipah virus W protein involves multiple discrete interactions with the nuclear import and export machinery.
Bo Zhong: Captive by the viral immune escape.
Tetherin antagonism by V proteins is a common trait among the genus Rubulavirus.
Antagonistic Coevolution of MER Tyrosine Kinase Expression and Function.
Merkel Cell Polyomavirus and Human Merkel Cell Carcinoma.
Coxsackievirus and Type 1 Diabetes: Diabetogenic Mechanisms and Implications for Prevention.

Targeting viperin to the mitochondrion inhibits the thiolase activity of the trifunctional enzyme complex.
Expression of all six human Torque teno virus (TTV) proteins in bacteria and in insect cells, and analysis of their IgG responses.
Betulonic Acid Derivatives Interfering with Human Coronavirus 229E Replication via the nsp15 Endoribonuclease.
Equine Arteritis Virus Elicits a Mucosal Antibody Response in the Reproductive Tract of Persistently Infected Stallions.
The role of M2-2 PDZ-binding motifs in pulmonary innate immune responses to human metapneumovirus.
The current status and future directions of myxoma virus, a master in immune evasion.
Inhibition of TNF receptor 1 internalization by adenovirus 14.7K as a novel immune escape mechanism.
L233P mutation in the bovine leukemia virus Tax protein has impact on annexin A3 and type I collagen secretion by host cells.
Advances in the Arms Race Between Silkworm and Baculovirus.
Enhanced alpha1 microglobulin secretion from Hepatitis E virus ORF3-expressing human hepatoma cells is mediated by the tumor susceptibility gene 101.
Endocytic activity of HIV-1 Vpu: Phosphoserine-dependent interactions with clathrin adaptors.
Equine infectious anemia.
A cornucopia of human polyomaviruses.
Rapid dynamics of polyomavirus type BK in renal transplant recipients.
Polyomavirus T Antigen Induces APOBEC3B Expression Using an LXCXE-Dependent and TP53-Independent Mechanism.
Diversity of ubiquitin and ISG15 specificity among nairoviruses' viral ovarian tumor domain proteases.
Structural basis for recruitment of host CypA and E3 ubiquitin ligase by maedi-visna virus Vif.
PHD domains and E3 ubiquitin ligases: viruses make the connection.
Potential chemotherapeutic targets for Japanese encephalitis: current status of antiviral drug development and future challenges.
Enteroviruses: A Gut-Wrenching Game of Entry, Detection, and Evasion.
Ancient and recent selective pressures shaped genetic diversity at AIM2-like nucleic acid sensors.
Immunobiology of Newcastle Disease Virus and Its Use for Prophylactic Vaccination in Poultry and as Adjuvant for Therapeutic Vaccination in Cancer Patients.
Enrichment of Zα domains at cytoplasmic stress granules is due to their innate ability to bind to nucleic acids.
Pseudorabies virus variant in Bartha-K61-vaccinated pigs, China, 2012.
Simian virus 40 inhibits differentiation and maturation of rhesus macaque DC-SIGN(+) dendritic cells.
Overexpression Bombyx mori HEXIM1 Facilitates Immune Escape of Bombyx mori Nucleopolyhedrovirus by Suppressing BmRelish-Driven Immune Responses.
Genome-wide analysis of codon usage bias in Ebolavirus.
Latency of viral expression in vivo is not related to CpG methylation in the U3 region and part of the R region of the long terminal repeat of bovine leukemia virus.
Detection of RNA modifications by HPLC analysis and competitive ELISA.
Transcriptional regulation of the bovine leukemia virus promoter by the cyclic AMP-response element modulator tau isoform.
Prediction of coronavirus 3C-like protease cleavage sites using machine-learning algorithms.

Differential lymphocyte and antibody responses in deer mice infected with Sin Nombre hantavirus or Andes hantavirus.
Zoonotic Potential of Emerging Paramyxoviruses: Knowns and Unknowns.
Structural mechanisms of DREAM complex assembly and regulation.
Genome-wide analysis of codon usage bias in Bovine Coronavirus.
An update on the clinical management of cutaneous molluscum contagiosum.
The dynamics of murid gammaherpesvirus 4 within wild, sympatric populations of bank voles and wood mice.
Bile Salts Alter the Mouse Norovirus Capsid Conformation: Possible Implications for Cell Attachment and Immune Evasion.
HTLV-I-infected breast milk macrophages inhibit monocyte differentiation to dendritic cells.
CD59 association with infectious bronchitis virus particles protects against antibody-dependent complement-mediated lysis.
In vitro antiviral activity of single domain antibody fragments against poliovirus.
Porcine deltacoronavirus accessory protein NS7a possesses the functional characteristics of a viroporin.
Binding to DCAF1 distinguishes TASOR and SAMHD1 degradation by HIV-2 Vpx.
Fugetaxis: active movement of leukocytes away from a chemokinetic agent.
Cytosolic Fc receptor TRIM21 inhibits seeded tau aggregation.
Torque teno virus microRNA detection in cerebrospinal fluids of patients with neurological pathologies.
SOCS3 Promotes ALV-J Virus Replication via Inhibiting JAK2/STAT3 Phosphorylation During Infection.
Generation of competent bone marrow-derived antigen presenting cells from the deer mouse ( <i>Peromyscus maniculatus</i> ).
Characterization of an RNA silencing suppressor encoded by maize yellow dwarf virus-RMV2.
Core binding factor beta plays a critical role by facilitating the assembly of the Vif-cullin 5 E3 ubiquitin ligase.
Study of the activation of the PI3K/Akt pathway by the motif of $\sigma A$ and $\sigma NS$ proteins of avian reovirus.
The Immune Response to Astrovirus Infection.
Molecular Mechanism of Adenovirus Late Protein L4-100K Chaperones the Trimerization of Hexon.
Ehrlichia SLiM ligand mimetic activates Hedgehog signaling to engage a BCL-2 anti-apoptotic cellular program.
Human endogenous retrovirus HERV-K(HML-2) encodes a stable signal peptide with biological properties distinct from Rec.
Molecular characterization of chronic-type adult T-cell leukemia/lymphoma.
Nonsense-containing mRNAs that accumulate in the absence of a functional nonsense-mediated mRNA decay pathway are destabilized rapidly upon its restitution.

Table A2-62, Cluster 61

Cluster 61 focuses on the tumor microenvironment, especially immune suppression in the tumor microenvironment, and its role in promoting tumor growth (1086)
The tumor microenvironment and its role in promoting tumor growth.
Pattern response of dendritic cells in the tumor microenvironment and breast cancer.
Tumor immune escape mechanisms that operate during metastasis.

Immune suppression in the tumor microenvironment: a role for dendritic cell-mediated tolerization of T cells.
Tumor-Infiltrating Dendritic Cells: Decisive Roles in Cancer Immunosurveillance, Immunoediting, and Tumor T Cell Tolerance.
Signaling defects in anti-tumor T cells. [Tumor-induced immunosuppression].
Acquired resistance to cancer immunotherapy: Role of tumor-mediated immunosuppression.
Immunoregulatory signal networks and tumor immune evasion mechanisms: insights into therapeutic targets and agents in clinical development.
Tumor-host immune interactions and dendritic cell dysfunction.
Tumor immunology--when a cancer cell meets the immune cells.
The role of immune cells in the tumor microenvironment.
Dendritic cells and tumor microenvironment: a dangerous liaison.
Camouflage and sabotage: tumor escape from the immune system.
Evaluation of Tumor-infiltrating Leukocyte Subsets in a Subcutaneous Tumor Model.
Role of dendritic cells in the regulation of antitumor immunity.
Immunotherapy: Reshape the Tumor Immune Microenvironment.
B cells promote tumor progression in a mouse model of HPV-mediated cervical cancer.
Regulation of tumor immunity by tumor/dendritic cell fusions.
Immunosuppressive cells in tumor immune escape and metastasis.
Tumor-altered dendritic cell function: implications for anti-tumor immunity.
Vaccines against human carcinomas: strategies to improve antitumor immune responses.
High immunosuppressive burden in cancer patients: a major hurdle for cancer immunotherapy.
Mechanisms of tumor-induced T cell immune suppression and therapeutics to counter those effects.
Tricks tumors use to escape from immune control.
The mechanisms tumor cells utilize to evade the host's immune system.
Paradoxical effects of cytokines in tumor immune surveillance and tumor immune escape.
Tumor Immune Microenvironment and Its Related miRNAs in Tumor Progression.
Mechanisms of tumor escape from the immune system: adenosine-producing Treg, exosomes and tumor-associated TLRs.
The tumor immune-microenvironment in gastric cancer.
Targeting Autophagy in the Tumor Microenvironment: New Challenges and Opportunities for Regulating Tumor Immunity.
Immunomodulatory Activity of VEGF in Cancer.
Dynamic cross-talk between tumor and immune cells in orchestrating the immunosuppressive network at the tumor microenvironment. [Immune system and tumors].
[Anti-tumor effect and impact on tumor immune microenvironment of tumor-targeted Salmonella VNP20009].
The Role of Autophagy in Tumor Immunology-Complex Mechanisms That May Be Explored Therapeutically.
Deciphering immune microenvironment and cell evasion mechanisms in human gliomas.
Using Flow Cytometry to Study Myc's Role in Shaping the Tumor Immune Microenvironment.
Predicting prognosis and therapeutic response from interactions between lymphocytes and tumor cells.

Tumor microenvironment-mediated immune tolerance in development and treatment of gastric cancer.
Induction of potent antitumor immunity by in situ targeting of intratumoral DCs.
Effector-phase tolerance: another mechanism of how cancer escapes antitumor immune response.
HPV16-associated tumors control myeloid cell homeostasis in lymphoid organs, generating a suppressor environment for T cells.
Tumor-induced modulation of dendritic cell function.
Unraveling How Tumor-Derived Galectins Contribute to Anti-Cancer Immunity Failure.
Cryo-ablation improves anti-tumor immunity through recovering tumor educated dendritic cells in tumor-draining lymph nodes.
The Role of the Tumor Vasculature in the Host Immune Response: Implications for Therapeutic Strategies Targeting the Tumor Microenvironment.
Oncolytic virus-mediated reversal of impaired tumor antigen presentation.
Crosstalk of Immune Cells and Platelets in an Ovarian Cancer Microenvironment and Their Prognostic Significance.
Dendritic cells in the cancer microenvironment.
T lymphocyte-derived TNF and IFN- $\gamma$ repress HFE expression in cancer cells.
Molecular Mechanism of Tumor Cell Immune Escape Mediated by CD24/Siglec-10.
Strategies of tumor immune evasion.
Glioblastoma progression is assisted by induction of immunosuppressive function of pericytes through interaction with tumor cells.
Tumors and the danger model.
Swords into plowshares: IL-23 repurposes tumor immune surveillance.
Mechanisms of tumor escape from immune system: role of mesenchymal stromal cells.
Role of p53 in the sensitization of tumor cells to apoptotic cell death.
The systemic-level repercussions of cancer-associated inflammation mediators produced in the tumor microenvironment.
T-cell mediated anti-tumor immunity after photodynamic therapy: why does it not always work and how can we improve it?
Dendritic cells in hematological malignancies.
Radiotherapy induced immunogenic cell death by remodeling tumor immune microenvironment.
Ovarian cancer biology and immunotherapy.
Tumor-driven evolution of immunosuppressive networks during malignant progression.
Antibodies to vascular endothelial growth factor enhance the efficacy of cancer immunotherapy by improving endogenous dendritic cell function.
Strategies to overcome DC dysregulation in the tumor microenvironment.
Tumor immune surveillance and ovarian cancer: lessons on immune mediated tumor rejection or tolerance.
STING, DCs and the link between innate and adaptive tumor immunity.
The immunomodulatory role of angiocidin, a novel angiogenesis inhibitor.
Lack of myeloid cell infiltration as an acquired resistance strategy to immunotherapy.
Adaptive Resistance to Immunotherapy Directed Against p53 Can be Overcome by Global Expression of Tumor-Antigens in Dendritic Cells.
Modulation of innate immunity in the tumor microenvironment.
Suppressive Myeloid Cells Shape the Tumor Immune Microenvironment.

Association between Inflammation and Function of Cell Adhesion Molecules Influence on Gastrointestinal Cancer Development.
Pas de Deux: Control of Anti-tumor Immunity by Cancer-Associated Inflammation.
Dendritic cells integrate signals from the tumor microenvironment to modulate immunity and tumor growth.
Enhancing the Efficacy of Tumor Vaccines Based on Immune Evasion Mechanisms.
Stat3 as a potential target for cancer immunotherapy.
[Vascular endothelial growth factor. From basic research to clinical application].
How to Hit Mesenchymal Stromal Cells and Make the Tumor Microenvironment Immunostimulant Rather Than Immunosuppressive.
Role of Oncogenic Pathways on the Cancer Immunosuppressive Microenvironment and Its Clinical Implications in Hepatocellular Carcinoma.
T-cell dysfunction in the glioblastoma microenvironment is mediated by myeloid cells releasing interleukin-10.
[The pattern recognition receptor-mediated immune tolerance is involved in tumor metastasis].
Bacterial immunotherapy of gastrointestinal tumors.
Tumor-derived factors impaired motility and immune functions of dendritic cells through derangement of biophysical characteristics and reorganization of cytoskeleton.
Transcriptomic signatures of tumors undergoing T cell attack.
The immune response to tumors.
[Cancer immunotherapy. Importance of overcoming immune suppression].
Adaptive Resistance to Cancer Immunotherapy.
Adaptive immune responses associated with breast cancer relapse.
Tumor-derived death receptor 6 modulates dendritic cell development.
Immunosuppressive Roles of Galectin-1 in the Tumor Microenvironment.
SIRT7 orchestrates melanoma progression by simultaneously promoting cell survival and immune evasion via UPR activation.
The Immunogenicity of Colorectal Cancer in Relation to Tumor Development and Treatment.
Dendritic Cells in Colorectal Cancer and a Potential for their Use in Therapeutic Approaches.
Epigenetic interplay between immune, stromal and cancer cells in the tumor microenvironment.
PLGA-polymer encapsulating tumor antigen and CpG DNA administered into the tumor microenvironment elicits a systemic antigen-specific IFN- $\gamma$ response and enhances survival.
Human embryo immune escape mechanisms rediscovered by the tumor.
The Role of miRNA in Tumor Immune Escape and miRNA-Based Therapeutic Strategies.
Interactions Between Anti-Angiogenic Therapy and Immunotherapy in Glioblastoma.
Restoring tumor immunogenicity with dendritic cell reprogramming.
Tumor-derived MUC1 mucins interact with differentiating monocytes and induce IL-10 <sup>high</sup> IL-12 <sup>low</sup> regulatory dendritic cell.
Immunogenicity of murine solid tumor models as a defining feature of in vivo behavior and response to immunotherapy.
Molecular profiling to identify relevant immune resistance mechanisms in the tumor microenvironment.
Current immunotherapeutic strategies in pancreatic cancer.
Interleukin-10 production by tumor infiltrating macrophages plays a role in Human Papillomavirus 16 tumor growth.
miRNAs as novel immunoregulators in cancer.

Melanoma-derived factors alter the maturation and activation of differentiated tissue-resident dendritic cells.
Local and systemic immunosuppression in pancreatic cancer: Targeting the stalwarts in tumor's arsenal.
4-1BBL as a Mediator of Cross-Talk between Innate, Adaptive, and Regulatory Immunity against Cancer.
A cancer treatment based on synergy between anti-angiogenic and immune cell therapies.
Anti-CD73 antibody therapy inhibits breast tumor growth and metastasis.
Roles of long noncoding RNAs on tumor immune escape by regulating immune cells differentiation and function.
How do tumors actively escape from host immunosurveillance?
The tumor microenvironment: key to early detection.
The Complexity of Targeting Chemokines to Promote a Tumor Immune Response.
Tumor resistance to specific lysis: a major hurdle for successful immunotherapy of cancer.
The immune response in cancer: from immunology to pathology to immunotherapy.
Immune manipulation of advanced breast cancer: an interpretative model of the relationship between immune system and tumor cell biology.
Calreticulin promotes tumor lymphocyte infiltration and enhances the antitumor effects of immunotherapy by up-regulating the endothelial expression of adhesion molecules.
CXCL8 Signaling in the Tumor Microenvironment.
Molecular and Cellular Mechanisms of Antitumor Immune Response Activation by Dendritic Cells.
The Janus face of dendritic cells in cancer.
Modulation of the tumor microenvironment by natural agents: implications for cancer prevention and therapy.
[Immune response and colorectal cancer].
Toll-like receptors on tumor cells facilitate evasion of immune surveillance.
Tumor associated regulatory dendritic cells.
GITR agonist enhances vaccination responses in lung cancer.
Extracellular Vesicles Secreted by Tumor Cells Promote the Generation of Suppressive Monocytes.
MAL2 drives immune evasion in breast cancer by suppressing tumor antigen presentation.
The administration of drugs inhibiting cholesterol/oxysterol synthesis is safe and increases the efficacy of immunotherapeutic regimens in tumor-bearing mice.
The role of p53 in anti-tumor immunity and response to immunotherapy.
Angiogenesis and the tumor vasculature as antitumor immune modulators: the role of vascular endothelial growth factor and endothelin.
Tumor immune escape mechanisms: impact of the neuroendocrine system.
The Role of Type 1 Conventional Dendritic Cells in Cancer Immunity.
Cancer immunotherapy based on killing of Salmonella-infected tumor cells.
How patients with an intact immune system develop head and neck cancer.
A new personalized vaccine strategy based on inducing the pyroptosis of tumor cells in vivo by transgenic expression of a truncated GSDMD N-terminus.
Immunosuppressive cells in cancer: mechanisms and potential therapeutic targets.
Study of galectins in tumor immunity: strategies and methods.
T-cell recognition of non-mutated tumor antigens in healthy individuals: connecting endogenous immunity and tumor dormancy.

Antibody-directed effector cell therapy of tumors: analysis and optimization using a physiologically based pharmacokinetic model.
Interleukin-17D promotes lung cancer progression by inducing tumor-associated macrophage infiltration via the p38 MAPK signaling pathway.
Cell Trafficking at the Intersection of the Tumor-Immune Compartments.
Tumor-host interactions: the role of inflammation.
Immune dysfunction in cancer patients.
Anti-tumor immunity induced by ectopic expression of viral antigens is transient and limited by immune escape.
Tumor-derived ILT4 induces T cell senescence and suppresses tumor immunity.
Autoantibodies against tumor-related antigens: incidence and biologic significance.
Small GTPase RBJ promotes cancer progression by mobilizing MDSCs via IL-6.
The therapeutic effect of an autologous and allogenic mixed glioma cell lysate vaccine in a rat model.
A team effort: natural killer cells on the first leg of the tumor immunity relay race.
Cancer acidity: An ultimate frontier of tumor immune escape and a novel target of immunomodulation.
Long Non-coding RNAs: Emerging Roles in the Immunosuppressive Tumor Microenvironment.
Modulation of the tumor vasculature and oxygenation to improve therapy.
Tumor's other immune targets: dendritic cells.
The Influence of Tumor Microenvironment on Immune Escape of Melanoma.
Role of Radiation Therapy in Modulation of the Tumor Stroma and Microenvironment.
Cancer/testis Antigen MAGEA3 Interacts with STAT1 and Remodels the Tumor Microenvironment.
Toll-like receptors expressed in tumor cells: targets for therapy.
Tumor-derived extracellular vesicles activate primary monocytes.
Interleukin-17, produced by lymphocytes, promotes tumor growth and angiogenesis in a mouse model of breast cancer.
Dendritic cell defects in the colorectal cancer.
Molecular Heterogeneity and Immunosuppressive Microenvironment in Glioblastoma.
Emergence of immunotherapy as a novel way to treat hepatocellular carcinoma.
The concept of immune surveillance against tumors. The first theories.
Local secretion/shedding of tumor-derived CD83 molecules as a novel tumor escape mechanism.
Yin and yang roles of B lymphocytes in solid tumors: Balance between antitumor immunity and immune tolerance/immunosuppression in tumor-draining lymph nodes.
Cytomegalovirus and tumors: two players for one goal-immune escape.
Downregulation of IFN-gammaR in association with loss of Fas function is linked to tumor progression.
Identification of Novel Tumor Microenvironment Regulating Factor That Facilitates Tumor Immune Infiltration in Cervical Cancer.
Minimal recruitment and activation of dendritic cells within renal cell carcinoma.
STAT3 Establishes an Immunosuppressive Microenvironment during the Early Stages of Breast Carcinogenesis to Promote Tumor Growth and Metastasis.
Crk adaptor protein promotes PD-L1 expression, EMT and immune evasion in a murine model of triple-negative breast cancer.
Immunosuppressive factors: role in cancer development and progression.
The two faces of IL-6 in the tumor microenvironment.
Canvassing Prospects of Glyco-Nanovaccines for Developing Cross-Presentation Mediated Anti-Tumor Immunotherapy.



Regulatory dendritic cells in the tumor immunoenvironment.
Insights Into Mechanisms of Tumor and Immune System Interaction: Association With Wound Healing.
Immune responses to malignancies.
LincRNA-immunity landscape analysis identifies EPIC1 as a regulator of tumor immune evasion and immunotherapy resistance.
Targeting Endocytosis and Cell Communications in the Tumor Immune Microenvironment.
Molecular mechanisms involved in dendritic cell dysfunction in cancer.
Generation of the tumor-suppressive secretome from tumor cells.
Mechanisms of local immunoresistance in glioma.
A Window of Opportunity: Targeting Cancer Endothelium to Enhance Immunotherapy.
Natural selection of tumor variants in the generation of "tumor escape" phenotypes.
Vascular endothelial growth factor (VEGF) impairs the motility and immune function of human mature dendritic cells through the VEGF receptor 2-RhoA-cofilin1 pathway.
An overview of current therapeutic strategies for glioblastoma and the role of CD73 as an alternative curative approach.
Mechanisms of immune evasion by tumors.
T cell-tumor interaction directs the development of immunotherapies in head and neck cancer.
Remodeling of tumor stroma and response to therapy.
The reciprocal function and regulation of tumor vessels and immune cells offers new therapeutic opportunities in cancer.
Altered expression of 15-hydroxyprostaglandin dehydrogenase in tumor-infiltrated CD11b myeloid cells: a mechanism for immune evasion in cancer.
Tumor-intrinsic signaling pathways: key roles in the regulation of the immunosuppressive tumor microenvironment.
Dynamic Patterns of Clonal Evolution in Tumor Vasculature Underlie Alterations in Lymphocyte-Endothelial Recognition to Foster Tumor Immune Escape.
Immunosuppressive strategies that are mediated by tumor cells.
N(6)-methyladenosine (m(6)A) RNA modification in tumor immunity.
Enhanced cancer immunotherapy using STAT3-depleted dendritic cells with high Th1-inducing ability and resistance to cancer cell-derived inhibitory factors.
The role of immune system exhaustion on cancer cell escape and anti-tumor immune induction after irradiation.
B7-H7: A potential target for cancer immunotherapy.
Potential targeting of B7-H4 for the treatment of cancer.
Immune Tumor Microenvironment in Ovarian Cancer Ascites.
[Molecular bases of cancer immunology].
Anti-Tumor Immunity in Head and Neck Cancer: Understanding the Evidence, How Tumors Escape and Immunotherapeutic Approaches.
The Impact of Chemotherapy, Radiation and Epigenetic Modifiers in Cancer Cell Expression of Immune Inhibitory and Stimulatory Molecules and Anti-Tumor Efficacy.
Prostaglandin E Receptor 4 Antagonist in Cancer Immunotherapy: Mechanisms of Action.
The lactate receptor GPR81 promotes breast cancer growth via a paracrine mechanism involving antigen-presenting cells in the tumor microenvironment.
Extrachromosomal DNA formation enables tumor immune escape potentially through regulating antigen presentation gene expression.

Role of the tumor microenvironment in PD-L1/PD-1-mediated tumor immune escape.
Escape from host-antitumor immunity.
Pro-tumoral immune cell alterations in wild type and Shb-deficient mice in response to 4T1 breast carcinomas.
Factors released by the tumor far microenvironment are decisive for pancreatic adenocarcinoma development and progression.
Multifaceted Role of the Placental Growth Factor (PlGF) in the Antitumor Immune Response and Cancer Progression.
Thrombin-PAR1 signaling in pancreatic cancer promotes an immunosuppressive microenvironment.
Tumor factors stimulate lysosomal degradation of tumor antigens and undermine their cross-presentation in lung cancer.
Immune evasion in cancer: Mechanistic basis and therapeutic strategies.
The Landscape and Clinical Application of the Tumor Microenvironment in Gastroenteropancreatic Neuroendocrine Neoplasms.
Hyperglycosylation of prosaposin in tumor DCs promotes immune escape in cancer.
Disruption of sialic acid metabolism drives tumor growth by augmenting CD8(+) T cell apoptosis.
MUC1-specific CTLs are non-functional within a pancreatic tumor microenvironment.
The flip side of immune surveillance: immune dependency.
cC1qR/CR and gC1qR/p33: observations in cancer.
Targeting Toll-Like Receptors for Cancer Therapy.
The tyrp1-Tag/tyrp1-FGFR1-DN bigenic mouse: a model for selective inhibition of tumor development, angiogenesis, and invasion into the neural tissue by blockade of fibroblast growth factor receptor activity.
CD73-generated adenosine: orchestrating the tumor-stroma interplay to promote cancer growth.
Strategies for tumor immune escape.
HIF-1 $\alpha$ Inhibition Improves Anti-Tumor Immunity and Promotes the Efficacy of Stereotactic Ablative Radiotherapy (SABR).
CD40-induced countercurrent conduits for tumor escape or elimination?
Tumor counterattack: fact or fiction?
Cutaneous tumors cease CXCL9/Mig production as a result of IFN- $\gamma$ -mediated immunoediting.
Immune-mediated eradication of tumors through the blockade of transforming growth factor-beta signaling in T cells.
Cleaver-1/stabilin-1 controls cancer growth and metastasis.
MicroRNAs and tumor vasculature normalization: impact on anti-tumor immune response.
Delivery of foreign cytotoxic T lymphocyte epitopes to tumor tissues for effective antitumor immunotherapy against pre-established solid tumors in mice.
Monocytes in the Tumor Microenvironment.
Coordination of intratumoral immune reaction and human colorectal cancer recurrence.
Tracking the common ancestry of antigenically distinct cancer variants.
Phases of apoptosis of melanoma cells, but not of normal melanocytes, differently affect maturation of myeloid dendritic cells.
p53 mutation and deletion contribute to tumor immune evasion.
Molecular genetics of brain tumors.
Type III TGF- $\beta$ receptor downregulation generates an immunotolerant tumor microenvironment.
In vivo Effects in Melanoma of ROCK Inhibition-Induced FasL Overexpression.
Anti-cancer therapies targeting the tumor stroma.

ER Stress Sensor XBP1 Controls Anti-tumor Immunity by Disrupting Dendritic Cell Homeostasis.
Recombinant tumor-associated MUC1 glycoprotein impairs the differentiation and function of dendritic cells.
5-Azacytidine Potentiates Anti-tumor Immunity in a Model of Pancreatic Ductal Adenocarcinoma.
Intelligent photothermal dendritic cells restart the cancer immunity cycle through enhanced immunogenic cell death.
Signal transducer and activator of transcription 3 signaling in tumor immune evasion.
Reovirus virotherapy overrides tumor antigen presentation evasion and promotes protective antitumor immunity.
The role of CD200-CD200R in tumor immune evasion.
Opportunities and challenges to engineer 3D models of tumor-adaptive immune interactions.
Radio-Immune Response Modelling for Spatially Fractionated Radiotherapy.
The interplay between innate and adaptive immunity in cancer shapes the productivity of cancer immunosurveillance.
Cyclooxygenase-2-Prostaglandin E2 pathway: A key player in tumor-associated immune cells.
Role of VEGFs/VEGFR-1 Signaling and its Inhibition in Modulating Tumor Invasion: Experimental Evidence in Different Metastatic Cancer Models.
Cancer Acidity and Hypertonicity Contribute to Dysfunction of Tumor-Associated Dendritic Cells: Potential Impact on Antigen Cross-Presentation Machinery.
Toll-like Receptor-4 Activation Boosts the Immunosuppressive Properties of Tumor Cells-derived Exosomes.
Radio-immune response modelling for spatially fractionated radiotherapy.
HMGB1 mediates endogenous TLR2 activation and brain tumor regression.
Pannexin1 channel-dependent secretome from apoptotic tumor cells shapes immune-escape microenvironment.
Chemokines in tumor-associated angiogenesis.
Dual character of Toll-like receptor signaling: pro-tumorigenic effects and anti-tumor functions.
Natural immunity to cancer in humans.
Induction of lymphoidlike stroma and immune escape by tumors that express the chemokine CCL21.
Commentary: Immune escape versus tumor tolerance: how do tumors evade immune surveillance?
Cancer immunoediting: integrating immunity's roles in cancer suppression and promotion.
Defective generation and maturation of dendritic cells from monocytes in colorectal cancer patients during the course of disease.
B cells and tumor immune escape.
New dimensions in tumor immunology: what does 3D culture reveal?
The Dynamics of Interactions Among Immune and Glioblastoma Cells.
On-chip modeling of tumor evolution: Advances, challenges and opportunities.
Epigenetic Changes of the Immune System with Role in Tumor Development.
Oncogenic growth factor signaling mediating tumor escape from cellular immunity.
Tumor-intrinsic LKB1-LIF signaling axis establishes a myeloid niche to promote immune evasion and tumor growth.
The Regulatory Effects of MicroRNAs on Tumor Immunity.
The importance of N6-methyladenosine modification in tumor immunity and immunotherapy.
Genetic and modifying factors that determine the risk of brain tumors.
Tumor-educated CD11b <sup>high</sup> IL10 <sup>high</sup> regulatory dendritic cells suppress T cell response through arginase I.

Altered Lipid Tumor Environment and Its Potential Effects on NKT Cell Function in Tumor Immunity.
Role of B7-H1 and B7-H4 molecules in down-regulating effector phase of T-cell immunity: novel cancer escaping mechanisms.
Tumor stroma-associated antigens for anti-cancer immunotherapy.
Anticancer potential of naturally occurring immunoepigenetic modulators: A promising avenue?
An epigenetically altered tumor cell vaccine.
BiTE secretion from in situ-programmed myeloid cells results in tumor-retained pharmacology.
The immunomodulating roles of glycoproteins in epithelial ovarian cancer.
Identification of tumorigenic cells and therapeutic targets in pancreatic neuroendocrine tumors.
A stressful microenvironment: opposing effects of the endoplasmic reticulum stress response in the suppression and enhancement of adaptive tumor immunity.
Immune modulatory microRNAs as a novel mechanism to revert immune escape of tumors.
Immune Landscape of Thyroid Cancers: New Insights.
Hijacker of the Antitumor Immune Response: Autophagy Is Showing Its Worst Facet.
Tumor blood vessels, a difficult hurdle for infiltrating leukocytes.
Fine-tuning antitumor responses through the control of galectin-glycan interactions: an overview. [Escape Strategies of Tumors from Immune Surveillance].
Human cancer evolution in the context of a human immune system in mice.
Better understanding tumor-host interaction in head and neck cancer to improve the design and development of immunotherapeutic strategies.
Tumor-induced endothelial cell surface heterogeneity directly affects endothelial cell escape from a cell-mediated immune response in vitro.
Immune System in Action.
Transient downregulation of monocyte-derived dendritic-cell differentiation, function, and survival during tumoral progression and regression in an in vivo canine model of transmissible venereal tumor.
Tumor immunoediting and immunosculpting pathways to cancer progression.
Autologous tumor rejection in humans: trimming the myths.
CD73 and adenosine generation in the creation of regulatory microenvironments.
Dual tumor targeting with pH-sensitive and bioreducible polymer-complexed oncolytic adenovirus.
Gaucher disease: chemotactic factors and immunological cell invasion in a mouse model.
RNA expression of complement regulatory proteins in human brain tumors.
The role of lysosomal peptidases in glioma immune escape: underlying mechanisms and therapeutic strategies.
CD155, an onco-immunologic molecule in human tumors.
IL-33 and ST2 mediate FAK-dependent antitumor immune evasion through transcriptional networks.
A murine model for the development of melanocytic nevi and their progression to melanoma.
Immune escape mechanism: defective resting and stimulated leukocyte-endothelium interaction in hepatocellular carcinoma of the rat.
Gimme shelter: the immune system during pregnancy.
Chemokine modulation of the tumor microenvironment.
Failure of blood-thymus barrier as a mechanism of tumor and trophoblast escape.
Salivary duct carcinoma with striking neutrophil-tumor cell cannibalism.
Distinctive Subpopulations of Stromal Cells Are Present in Human Lymph Nodes Infiltrated with Melanoma.

Treatment and prevention of rat glioblastoma by immunogenic C6 cells expressing antisense insulin-like growth factor I RNA.
API5 confers tumoral immune escape through FGF2-dependent cell survival pathway.
A Role for MAIT Cells in Colorectal Cancer.
Epigenetic inactivation of follistatin-like 1 mediates tumor immune evasion in nasopharyngeal carcinoma.
Onco-immunomodulatory properties of pharmacological interference with RAS-RAF-MEK-ERK pathway hyperactivation.
Glioma cells display complex cell surface topographies that resist the actions of cytolytic effector lymphocytes.
Proteolytic single hinge cleavage of pertuzumab impairs its Fc effector function and antitumor activity in vitro and in vivo.
Cancer Immune Equilibrium and Schizophrenia Have Similar Interferon- $\gamma$ , Tumor Necrosis Factor- $\alpha$ , and Interleukin Expression: A Tumor Model of Schizophrenia.
High-throughput molecular and histopathologic profiling of tumor tissue in a novel transplantable model of murine neuroblastoma: new tools for pediatric drug discovery.
Progress in research into the role of abnormal glycosylation modification in tumor immunity.
Evasion of immunosurveillance by genomic alterations of PPAR $\gamma$ /RXR $\alpha$ in bladder cancer.
The RB tumor suppressor at the intersection of proliferation and immunity: relevance to disease immune evasion and immunotherapy.
The role of Fusobacteria in oral cancer and immune evasion.
Neopterin, a prognostic marker in human malignancies.
Use of surrogate antigens as vaccines against cancer.
Proteomic profiling reveals antitumor effects of RT2 peptide on a human colon carcinoma xenograft mouse model.
ADAR and Immune Silencing in Cancer.
Lineage tracing of mutant granulosa cells reveals in vivo protective mechanisms that prevent granulosa cell tumorigenesis.
Immunocytochemical detection of members of the caspase cascade of apoptosis in high-grade astrocytomas.
Increased blood spermine levels decrease the cytotoxic activity of lymphokine-activated killer cells: a novel mechanism of cancer evasion.
Tim-3 promotes tube formation and decreases tight junction formation in vascular endothelial cells.
Aberrant miRNAs Regulate the Biological Hallmarks of Glioblastoma.
The Good, The Bad, and The Unexpected: Roles of DUX4 in Health and Disease.
Excretory/secretory products from plerocercoids of <i>Spirometra erinaceieuropaei</i> suppress gene expressions and production of tumor necrosis factor-alpha in murine macrophages stimulated with lipopolysaccharide or lipoteichoic acid.
[Immunological analogies between ovarian cancer and pregnancy].
Serum Tumor Necrosis Factor-Alpha Levels in Acute Leukemia and Its Prognostic Significance.
The sialoglycan-Siglec-E checkpoint axis in dexamethasone-induced immune subversion in glioma-microglia transwell co-culture system.
Heme oxygenase and carbon monoxide as an immunotherapeutic approach in transplantation and cancer.
Cancer-initiating cells from colorectal cancer patients escape from T cell-mediated immunosurveillance in vitro through membrane-bound IL-4.
Patterns of immune equilibrium and escape in indolent and progressing tumors.

Targeting activation-induced cytidine deaminase overcomes tumor evasion of immunotherapy by CTLs.
The Role of Pericytes in Regulation of Innate and Adaptive Immunity.
APOBEC Enzymes: Mutagenic Fuel for Cancer Evolution and Heterogeneity.
Controlling the Immune Suppressor: Transcription Factors and MicroRNAs Regulating CD73/NT5E.
Wnt1 silences chemokine genes in dendritic cells and induces adaptive immune resistance in lung adenocarcinoma.
PARP mediated DNA damage response, genomic stability and immune responses.
Extensive conservation of genomic imbalances in canine transmissible venereal tumors (CTVT) detected by microarray-based CGH analysis.
Gene therapy with GM-CSF, interleukin-4 and herpes simplex virus thymidine kinase shows strong antitumor effect on lung cancer.
PPAR $\gamma$ agonist inhibits c-Myc-mediated colorectal cancer tumor immune escape.
GARP: A Key Target to Evaluate Tumor Immunosuppressive Microenvironment.
Targeting PIM Kinases to Improve the Efficacy of Immunotherapy.
A study of the possible role of Fab-glycosylated IgG in tumor immunity.
Intricate confrontation: Research progress and application potential of TRIM family proteins in tumor immune escape.
Immune evasion of tumor cells using membrane-bound complement regulatory proteins.
Pericyte Secretome.
Alternative versus classical activation of macrophages.
Morphological evidence of neutrophil-tumor cell phagocytosis (cannibalism) in human gastric adenocarcinomas.
In situ tuning proangiogenic factor-mediated immunotolerance synergizes the tumoricidal immunity via a hypoxia-triggerable liposomal bio-nanoreactor.
Secreted protein acidic and rich in cysteine produced by human melanoma cells modulates polymorphonuclear leukocyte recruitment and antitumor cytotoxic capacity.
Oncogene MYCN regulates localization of NKT cells to the site of disease in neuroblastoma.
Culture expanded primary chondrocytes have potent immunomodulatory properties and do not induce an allogeneic immune response.
Human polyomaviruses and cancer: an overview.
Pyridoxal kinase and poly(ADP-ribose) affect the immune microenvironment of locally advanced cancers.
Targeted genomic sequencing of follicular dendritic cell sarcoma reveals recurrent alterations in NF- $\kappa$ B regulatory genes.
Specific ADAM10 inhibitors localize in exosome-like vesicles released by Hodgkin lymphoma and stromal cells and prevent sheddase activity carried to bystander cells.
Immune evasion of the CD1d/NKT cell axis.
Antitumor and immune-modulatory efficacy of dual-treatment based on levamisole and/or taurine in Ehrlich ascites carcinoma-bearing mice.
Tn glycosylation of the MUC6 protein modulates its immunogenicity and promotes the induction of Th17-biased T cell responses.
Tumor-derived endothelial cells evade apoptotic activity of the interferon-inducible IFI16 gene.
L-arginine supplementation does not affect chemically induced carcinogenesis and tumor growth in BALB-c mice.
Spontaneous generation of germline characteristics in mouse fibrosarcoma cells.
ATRX Alteration Contributes to Tumor Growth and Immune Escape in Pleomorphic Sarcomas.

Tyk2 is a tumor suppressor in colorectal cancer.
miR-155 expression in antitumor immunity: The higher the better?
Immune activation is essential for the antitumor activity of EZH2 inhibition in urothelial carcinoma.
Ferumoxyl- $\beta$ -glucan Inhibits Melanoma Growth via Interacting with Dectin-1 to Polarize Macrophages into M1 Phenotype.
Emerging Role of the Macrophage Migration Inhibitory Factor Family of Cytokines in Neuroblastoma. Pathogenic Effectors and Novel Therapeutic Targets?
Endogenous retroviruses and cancer.
Isolation and culture of decidual natural killer cells from term placenta and complete hydatidiform mole.
Epigenetic control of the immune escape mechanisms in malignant carcinomas.
Immunomodulating effect of ethyl pyruvate on nonsyngenic transplanted tumor in mice.
IL-10 Controls Early Microglial Phenotypes and Disease Onset in ALS Caused by Misfolded Superoxide Dismutase 1.
Gangliosides as Siglec ligands.
Tumor Targeting by Fusobacterium nucleatum: A Pilot Study and Future Perspectives.
Predator-Prey in Tumor-Immune Interactions: A Wrong Model or Just an Incomplete One?
Cheater-altruist synergy in public goods games.
Bounded-noise-induced transitions in a tumor-immune system interplay.
[The role of IL411 in immunoregulation: An update].
Identification and characterization of blocking nanobodies against human CD70.
Efficacy of anti-CD47 antibody-mediated phagocytosis with macrophages against primary effusion lymphoma.
Subversion of calreticulin exposure as a strategy of immune escape.
Association of Canine Osteosarcoma and Monocyte Phenotype and Chemotactic Function.
Complement controls the immune synapse and tumors control complement.
Primary peripheral nerve tumors associated with nerve-territory herpes zoster.
Case Report: Microglia Composition and Immune Response in an Immunocompetent Patient With an Intracranial Syphilitic Gumma.

Table A2-63, Cluster 62

Cluster 62 focuses on cancer, especially the mechanisms of cancer progression and metastasis and their relation to epithelial-mesenchymal plasticity and immune evasion (869)
Immunological Consequences of Epithelial-Mesenchymal Transition in Tumor Progression.
Epithelial-Mesenchymal Transition: Role in Cancer Progression and the Perspectives of Antitumor Treatment.
The role of inflammation in head and neck cancer.
Roles of toll-like receptors: From inflammation to lung cancer progression.
Revisiting epithelial-mesenchymal transition in cancer metastasis: the connection between epithelial plasticity and stemness.
TLR/WNT: A Novel Relationship in Immunomodulation of Lung Cancer.
Prostaglandin E2 and Cancer: Insight into Tumor Progression and Immunity.
Inflammation and lung carcinogenesis: applying findings in prevention and treatment.
Platelets in cancer metastasis: To help the "villain" to do evil.

Central Roles of STAT3-Mediated Signals in Onset and Development of Cancers: Tumorigenesis and Immunosurveillance.
Mechanistic insights into the interplays between neutrophils and other immune cells in cancer development and progression.
Sestrin2 in cancer: a foe or a friend?
Microenvironmental Influences on Metastasis Suppressor Expression and Function during a Metastatic Cell's Journey.
New insights into the role of EMT in tumor immune escape.
Role of the tumor microenvironment in regulating apoptosis and cancer progression.
The role of tumor microenvironment in therapeutic resistance.
Targeting Oncoimmune Drivers of Cancer Metastasis.
The mechanisms of cancer immunoescape and development of overcoming strategies.
On the origin of cancer metastasis.
Steps in metastasis: an updated review.
Role of STAT3 in lung cancer.
Tumor Immune Microenvironment during Epithelial-Mesenchymal Transition.
[A Review of the Roles of Endoplasmic Reticulum Stress in Cancer Cell Metastasis].
Therapeutic implications of cancer epithelial-mesenchymal transition (EMT).
The Landscape of Circulating Tumor Cell Research in the Context of Epithelial-Mesenchymal Transition.
Classical epithelial-mesenchymal transition (EMT) and alternative cell death process-driven blebbistatin metastatic-witch (BMW) pathways to cancer metastasis.
The metastatic niche formation: focus on extracellular vesicle-mediated dialogue between lung cancer cells and the microenvironment.
Influence of Aberrant Epigenetic Changes and the Tumor Microenvironment in Ovarian Cancer Metastasis.
Chemokines driven ovarian cancer progression, metastasis and chemoresistance: Potential pharmacological targets for cancer therapy.
Regulation of antitumor immunity by inflammation-induced epigenetic alterations.
Autophagy in cancer metastasis.
SNAIL1: Linking Tumor Metastasis to Immune Evasion.
STAT3 as a central regulator of tumor metastases.
Assessing the epithelial-to-mesenchymal plasticity in a small cell lung carcinoma (SCLC) and lung fibroblasts co-culture model.
Involvement of platelets in tumor cell metastasis.
Axis of evil: molecular mechanisms of cancer metastasis.
A mechanism for epithelial-mesenchymal heterogeneity in a population of cancer cells.
The Role of the Innate Immune System in Cancer Dormancy and Relapse.
Interplay Between Inflammation and Epigenetic Changes in Cancer.
Regulation of Cell Death Mechanisms by Melatonin: Implications in Cancer Therapy.
TLR signaling by tumor and immune cells: a double-edged sword.
Epithelial-Mesenchymal Plasticity in Cancer Progression and Metastasis.
Toll-like receptors: A pathway alluding to cancer control.
The role of toll-like receptors in colorectal cancer progression: evidence for epithelial to leucocytic transition.
The resolution of inflammation and cancer.



[Epithelial-mesenchymal transitions in cancer onset and progression].
Platinum sensitivity of ovarian cancer cells does not influence their ability to induce M2-type macrophage polarization.
Progranulin and its biological effects in cancer.
Epithelial-to-Mesenchymal Transition in Metastasis: Focus on Laryngeal Carcinoma.
Regulation of metastases by signal transducer and activator of transcription 3 signaling pathway: clinical implications.
Cell death under epithelial-mesenchymal transition control in prostate cancer therapeutic response.
Integrins and Epithelial-Mesenchymal Cooperation in the Tumor Microenvironment of Muscle-Invasive Lethal Cancers.
Could senescence phenotypes strike the balance to promote tumor dormancy?
Pro-Resolving Factor Administration Limits Cancer Progression by Enhancing Immune Response Against Cancer Cells.
Microparticle Phosphatidylserine Mediates Coagulation: Involvement in Tumor Progression and Metastasis.
Crucial roles of circulating tumor cells in the metastatic cascade and tumor immune escape: biology and clinical translation.
Epithelial-mesenchymal transition as strategic microenvironment mimicry for cancer cell survival and immune escape?
Targeting the tumor microenvironment as a potential therapeutic approach in colorectal cancer: Rational and progress.
Epithelial-mesenchymal transition: Initiation by cues from chronic inflammatory tumor microenvironment and termination by anti-inflammatory compounds and specialized pro-resolving lipids.
Epithelial-Mesenchymal Transition and Immune Evasion during Lung Cancer Progression: The Chicken or the Egg?
Cytokine-chemokine network driven metastasis in esophageal cancer; promising avenue for targeted therapy.
Role of the calcium-sensing receptor (CaSR) in cancer metastasis to bone: Identifying a potential therapeutic target.
Platelet adherence to cancer cells promotes escape from innate immune surveillance in cancer metastasis.
The role of platelet activation in tumor metastasis.
Exosome-encapsulated ncRNAs: Emerging yin and yang of tumor hallmarks.
Delineating the twin role of autophagy in lung cancer.
The Role of the Microbiome in Cancer Initiation and Progression: How Microbes and Cancer Cells Utilize Excess Energy and Promote One Another's Growth.
Targeting STAT3 Signaling Pathway in Colorectal Cancer.
WNT Signaling in Tumors: The Way to Evade Drugs and Immunity.
The Lymph Node Microenvironment May Invigorate Cancer Cells With Enhanced Metastatic Capacities.
Targeting STAT3 affects melanoma on multiple fronts.
Colon cancer and apoptosis.
Role of apoptosis resistance in immune evasion and metastasis of colorectal cancer.
Sweetening the hallmarks of cancer: Galectins as multifunctional mediators of tumor progression.
Tumor-Associated Inflammation: The Tumor-Promoting Immunity in the Early Stages of Tumorigenesis.

Inhibition of STAT3 Signaling Pathway by Terphenyllin Suppresses Growth and Metastasis of Gastric Cancer.
Unpacking the Complexity of Epithelial Plasticity: From Master Regulator Transcription Factors to Non-Coding RNAs.
Cell stemness, epithelial-to-mesenchymal transition, and immunoevasion: Intertwined aspects in cancer metastasis.
Cancer invasion and resistance: interconnected processes of disease progression and therapy failure.
Future of circulating tumor cells in the melanoma clinical and research laboratory settings.
[Contribution to tumor escape and chemotherapy response: A choice between senescence and apoptosis in heterogeneous tumors].
Communication between EMT and PD-L1 signaling: New insights into tumor immune evasion.
Mechanosensitive Ion Channel PIEZO1 Signaling in the Hall-Marks of Cancer: Structure and Functions.
Innate and acquired immune surveillance in the postdissemination phase of metastasis.
Osteopontin is involved in TLR4 pathway contributing to ovarian cancer cell proliferation and metastasis.
PAUF as a Target for Treatment of High PAUF-Expressing Ovarian Cancer.
The Immune Microenvironment and Cancer Metastasis.
[Molecular genetic principles of progression of malignant diseases].
The Role of Circulating Tumor Cells in the Metastatic Cascade: Biology, Technical Challenges, and Clinical Relevance.
Metastasis-Initiating Cells and Ecosystems.
AXL Receptor Tyrosine Kinase as a Promising Therapeutic Target Directing Multiple Aspects of Cancer Progression and Metastasis.
Cancer immunologists and cancer biologists: why we didn't talk then but need to now.
Epithelial-to-mesenchymal transition and autophagy induction in breast carcinoma promote escape from T-cell-mediated lysis.
Immune escape mechanisms in colorectal cancer pathogenesis and liver metastasis.
Overview of Evidence-Based Chemotherapy for Oral Cancer: Focus on Drug Resistance Related to the Epithelial-Mesenchymal Transition.
Dormancy, stemness, and therapy resistance: interconnected players in cancer evolution.
Molecular Characterization of Circulating Tumor Cells to Study Cancer Immunoevasion.
The role of complement in the tumor microenvironment.
Translational Regulation of Cancer Metastasis.
Tumor microenvironment: a main actor in the metastasis process.
Role of chemokines in tumor growth.
Neuropilins Controlling Cancer Therapy Responsiveness.
Prostate cancer cell-intrinsic interferon signaling regulates dormancy and metastatic outgrowth in bone.
Tumor-derived exosomes: Nanovesicles made by cancer cells to promote cancer metastasis.
The Role of Tumor Inflammatory Microenvironment in Lung Cancer.
Three-dimensional in vitro tumor models for cancer research and drug evaluation.
Shaping the Immune Landscape in Cancer by Galectin-Driven Regulatory Pathways.
Targeting STAT3 signaling pathway in cancer by agents derived from Mother Nature.
Platelet Microparticles and miRNA Transfer in Cancer Progression: Many Targets, Modes of Action, and Effects Across Cancer Stages.

The Hedgehog-Gli pathway in embryonic development and cancer: implications for pulmonary oncology therapy.
Ubiquitination and deubiquitination in the regulation of epithelial-mesenchymal transition in cancer: Shifting gears at the molecular level.
Targeting Mononuclear Phagocyte Receptors in Cancer Immunotherapy: New Perspectives of the Triggering Receptor Expressed on Myeloid Cells (TREM-1).
Chemokines in progression, chemoresistance, diagnosis, and prognosis of colorectal cancer.
Cooperation between Prostaglandin E2 and Epidermal Growth Factor Receptor in Cancer Progression: A Dual Target for Cancer Therapy.
Nuclear-localized costimulatory molecule 4-1BBL promotes colon cancer cell proliferation and migration by regulating nuclear Gsk3 $\beta$ , and is linked to the poor outcomes associated with colon cancer.
Role of Type I and II Interferons in Colorectal Cancer and Melanoma.
IL6/STAT3 Signaling Orchestrates Premetastatic Niche Formation and Immunosuppressive Traits in Lung.
Role of Calcium Homeostasis in Modulating EMT in Cancer.
Death receptors in chemotherapy and cancer.
Intra- and Extra-Cellular Events Related to Altered Glycosylation of MUC1 Promote Chronic Inflammation, Tumor Progression, Invasion, and Metastasis.
Cancer cell states recur across tumor types and form specific interactions with the tumor microenvironment.
The role of Toll-like receptor 4 signaling pathway in ovarian, cervical, and endometrial cancers.
Cooperative and Escaping Mechanisms between Circulating Tumor Cells and Blood Constituents.
Interaction of autophagy and Toll-like receptors: a regulatory cross-talk--even in cancer cells?
A mathematical model for pancreatic cancer growth and treatments.
Parasite and Cancer Relationship.
Galectins dysregulation: A way for cancer cells to invade and pervade.
A small molecule compound targeting STAT3 DNA-binding domain inhibits cancer cell proliferation, migration, and invasion.
Turning-Off Signaling by Siglecs, Selectins, and Galectins: Chemical Inhibition of Glycan-Dependent Interactions in Cancer.
The role of inflammation in kidney cancer.
The molecular biology of pulmonary metastasis.
Targeting cancer's weaknesses (not its strengths): Therapeutic strategies suggested by the atavistic model.
Systemic metastasis-targeted nanotherapeutic reinforces tumor surgical resection and chemotherapy.
Non-genetic adaptive resistance to KRAS(G12C) inhibition: EMT is not the only culprit.
Dialog beyond the Grave: Necrosis in the Tumor Microenvironment and Its Contribution to Tumor Growth.
Involvement of platelet-tumor cell interaction in immune evasion. Potential role of podocalyxin-like protein 1.
A membrane penetrating peptide aptamer inhibits STAT3 function and suppresses the growth of STAT3 addicted tumor cells.
IL-23R in laryngeal cancer: a cancer immunoediting process that facilitates tumor cell proliferation and results in cisplatin resistance.
Platelets in cancer development and diagnosis.
Crossing paths in Human Renal Cell Carcinoma (hRCC).

Galectins in the Tumor Microenvironment: Focus on Galectin-1.
ANGPTL3 affects the metastatic potential and the susceptibility of ovarian cancer cells to natural killer cell-mediated cytotoxicity.
CytomégaloVirus humain et cancers.
Antithrombotic Agents and Cancer.
Anti-cancer therapeutic strategies based on HGF/MET, EpCAM, and tumor-stromal cross talk.
Contribution of TLR signaling to the pathogenesis of colitis-associated cancer in inflammatory bowel disease.
The PKC universe keeps expanding: From cancer initiation to metastasis.
Immune Evasion Mechanism and AXL.
Intracellular galectins in cancer cells: potential new targets for therapy (Review).
Stat3 is activated in skin lesions by the local application of imiquimod, a ligand of TLR7, and inhibited by the recombinant peptide aptamer rS3-PA.
Osteopontin at the Crossroads of Inflammation and Tumor Progression.
Cytomegalovirus is a tumor-associated virus: armed and dangerous.
Complex cross-talk between EZH2 and miRNAs confers hallmark characteristics and shapes the tumor microenvironment.
Metastatic Colonization: Escaping Immune Surveillance.
A temporal examination of calcium signaling in cancer- from tumorigenesis, to immune evasion, and metastasis.
Capitalizing on the immunogenicity of dying tumor cells.
Galectins: Multitask signaling molecules linking fibroblast, endothelial and immune cell programs in the tumor microenvironment.
The role of cancer-derived microRNAs in cancer immune escape.
TLR4 signaling promotes immune escape of human lung cancer cells by inducing immunosuppressive cytokines and apoptosis resistance.
The role of toll-like receptor 4 (TLR4) in cancer progression: A possible therapeutic target?
Cancer-associated mesothelial cells are regulated by the anti-Müllerian hormone axis.
Complement as a Biological Tool to Control Tumor Growth.
EMT impairs breast carcinoma cell susceptibility to CTL-mediated lysis through autophagy induction.
Platelets: the point of interconnection among cancer, inflammation and cardiovascular diseases.
Stepping out of the flow: capillary extravasation in cancer metastasis.
Molecular mechanisms of tumor metastasis and angiogenesis.
The multifaceted role of autophagy in tumor evasion from immune surveillance.
Extrinsic cell death pathway plasticity: a driver of clonal evolution in cancer?
Interleukin-30/IL27p28 Shapes Prostate Cancer Stem-like Cell Behavior and Is Critical for Tumor Onset and Metastasis.
Effects on tumor development and metastatic dissemination by the NKG2D lymphocyte receptor expressed on cancer cells.
Small Extracellular Vesicles and Metastasis-Blame the Messenger.
CCRK is a novel signalling hub exploitable in cancer immunotherapy.
Wnt/ $\beta$ -catenin signaling in melanoma: Preclinical rationale and novel therapeutic insights.
The multifaceted role of reactive oxygen species in tumorigenesis.
Targeted Single-cell Isolation of Spontaneously Escaping Live Melanoma Cells for Comparative Transcriptomics.

Systemic Regulation of Cancer Development by Neuro-Endocrine-Immune Signaling Network at Multiple Levels.
IGF2: A Role in Metastasis and Tumor Evasion from Immune Surveillance?
Acquisition of tumor cell phenotypic diversity along the EMT spectrum under hypoxic pressure: Consequences on susceptibility to cell-mediated cytotoxicity.
circ-Keratin 6c Promotes Malignant Progression and Immune Evasion of Colorectal Cancer through microRNA-485-3p/Programmed Cell Death Receptor Ligand 1 Axis.
Hedgehog signaling: modulation of cancer properties and tumor microenvironment.
Blocking Muscarinic Receptor 3 Attenuates Tumor Growth and Decreases Immunosuppressive and Cholinergic Markers in an Orthotopic Mouse Model of Colorectal Cancer.
Putting the brakes on tumorigenesis with snake venom toxins: New molecular insights for cancer drug discovery.
Autophagy facilitates TLR4- and TLR3-triggered migration and invasion of lung cancer cells through the promotion of TRAF6 ubiquitination.
An updated review on molecular mechanisms underlying the anticancer effects of capsaicin.
B7 homologue 3 as a prognostic biomarker and potential therapeutic target in gastrointestinal tumors.
Chemokine receptor CXCR4: role in gastrointestinal cancer.
Tumor Plasticity and Resistance to Immunotherapy.
MicroRNAs play a central role in molecular dysfunctions linking inflammation with cancer.
Cell dormancy and tumor refractory.
The Emerging Role of Platelets in the Formation of the Micrometastatic Niche: Current Evidence and Future Perspectives.
Galectins in Glioma: Current Roles in Cancer Progression and Future Directions for Improving Treatment.
Stat3 Signaling Pathway: A Future Therapeutic Target for Bone-Related Diseases.
Targeting glycoprotein VI to disrupt platelet-mediated tumor cell extravasation.
Is complement good or bad for cancer patients? A new perspective on an old dilemma.
Platelets contribute to the initiation of colitis-associated cancer by promoting immunosuppression.
Old and New Players of Inflammation and Their Relationship With Cancer Development.
3D Tumor Models for Breast Cancer: Whither We Are and What We Need.
Roles of toll-like receptors in cancer: a double-edged sword for defense and offense.
Insights into the Role of Sialylation in Cancer Metastasis, Immunity, and Therapeutic Opportunity.
Failsafe program escape and EMT: a deleterious partnership.
Wnts and the hallmarks of cancer.
[Advances in the Study of Chemokine-like Factor Superfamily Members in Tumors].
Immunologic consequences of signal transducers and activators of transcription 3 activation in human squamous cell carcinoma.
Role of Stat3 in suppressing anti-tumor immunity.
MYC as a Multifaceted Regulator of Tumor Microenvironment Leading to Metastasis.
ANO1: More Than Just Calcium-Activated Chloride Channel in Cancer.
Decoding Strategies to Evade Immunoregulators Galectin-1, -3, and -9 and Their Ligands as Novel Therapeutics in Cancer Immunotherapy.
MiR-221-induced PUMA silencing mediates immune evasion of bladder cancer cells.
The role of STAT3 in glioblastoma progression through dual influences on tumor cells and the immune microenvironment.

Promotion of cell proliferation and inhibition of ADCC by cancerous immunoglobulin expressed in cancer cell lines.
The Vicious Cross-Talk between Tumor Cells with an EMT Phenotype and Cells of the Immune System.
Expression of Tumor-mediated CD137 ligand in human colon cancer indicates dual signaling effects.
Impact of Galectins in Resistance to Anticancer Therapies.
The countercurrent principle in invasion and metastasis of cancer cells. Recent insights on the roles of chemokines.
Wnt Signaling and Drug Resistance in Cancer.
Changing mutational and adaptive landscapes and the genesis of cancer.
A review of mathematical models of cancer-immune interactions in the context of tumor dormancy.
Tumor progression locus 2 (TPL2) in tumor-promoting Inflammation, Tumorigenesis and Tumor Immunity.
Aspirin, platelets, and cancer: The point of view of the internist.
The NRSF/REST transcription factor in hallmarks of cancer: From molecular mechanisms to clinical relevance.
To kill a cancer: Targeting the immune inhibitory checkpoint molecule, B7-H3.
Targeting Cancer Cell Dormancy.
Rapamycin suppresses TLR4-triggered IL-6 and PGE(2) production of colon cancer cells by inhibiting TLR4 expression and NF-kappaB activation.
Platelets: linking hemostasis and cancer.
Interleukin-5 facilitates lung metastasis by modulating the immune microenvironment.
CD11b(+)CTLA4(+) myeloid cells are a key driver of tumor evasion in colorectal cancer.
The Sweet Side of Immune Evasion: Role of Glycans in the Mechanisms of Cancer Progression.
Surgery-mediated tumor-promoting effects on the immune microenvironment.
Manipulating the chemokine-chemokine receptor network to treat cancer.
On the role of galectin-3 in cancer apoptosis.
Targeting transforming growth factor- $\beta$ signaling for enhanced cancer chemotherapy.
$\beta$ -Catenin Drives Butyrophilin-like Molecule Loss and $\gamma\delta$ T-cell Exclusion in Colon Cancer.
WEE1 inhibition alleviates resistance to immune attack of tumor cells undergoing epithelial-mesenchymal transition.
A specific STAT3-binding peptide exerts antiproliferative effects and antitumor activity by inhibiting STAT3 phosphorylation and signaling.
A FBXO7/EYA2-SCF(FBXW7) axis promotes AXL-mediated maintenance of mesenchymal and immune evasion phenotypes of cancer cells.
Oxidative stress, inflammation, and cancer: how are they linked?
Altered Cell Adhesion and Glycosylation Promote Cancer Immune Suppression and Metastasis.
The Role of Galectins in Cervical Cancer Biology and Progression.
Epigenetic drugs as pleiotropic agents in cancer treatment: biomolecular aspects and clinical applications.
Extracellular matrix remodeling in tumor progression and immune escape: from mechanisms to treatments.
Perivascular niches: critical hubs in cancer evolution.
The roles of CD73 in cancer.
AXL Targeting Overcomes Human Lung Cancer Cell Resistance to NK- and CTL-Mediated Cytotoxicity.
Insights into the Relationship between Pentraxin-3 and Cancer.
Complement inhibition in cancer therapy.

Functional genomic landscape of cancer-intrinsic evasion of killing by T cells.
SIRPy-expressing cancer stem-like cells promote immune escape of lung cancer via Hippo signaling.
Enzymatic Inactivation of Oxysterols in Breast Tumor Cells Constrains Metastasis Formation by Reprogramming the Metastatic Lung Microenvironment.
Recent advances in the development of sialyltransferase inhibitors to control cancer metastasis: A comprehensive review.
Apoptosis: Activation and Inhibition in Health and Disease.
Clusterin: the missing link in the calcium-dependent resistance of cancer cells to apoptogenic stimuli.
Tumor accomplice: T cell exhaustion induced by chronic inflammation.
Small molecule inhibitors of STAT3 for cancer therapy.
Galectins in the tumor endothelium: opportunities for combined cancer therapy.
Potential Co-Relation Between Chronic Periodontitis And Cancer - An Emerging Concept.
TLR4 signaling in cancer cells promotes chemoattraction of immature dendritic cells via autocrine CCL20.
Cell surface glycan-lectin interactions in tumor metastasis.
Role of Focal Adhesion Kinase in Small-Cell Lung Cancer and Its Potential as a Therapeutic Target.
Cytokine-Mediated Dysregulation of Signaling Pathways in the Pathogenesis of Multiple Myeloma.
Introduction to a review series on platelets and cancer.
Aberrant Sialylation in Cancer: Biomarker and Potential Target for Therapeutic Intervention?
The Role of HDACs in the Response of Cancer Cells to Cellular Stress and the Potential for Therapeutic Intervention.
Osteopontin: regulation in tumor metastasis.
Epithelial immune cell-like transition (EIT): a proposed transdifferentiation process underlying immune-suppressive activity of epithelial cancers.
Cancer Immunology and Immunotherapy.
Cancer therapy targeting the fibrinolytic system.
[Galectines as prognostic factors for the malignant lymphoproliferative diseases].
Mechanisms of disease: Insights into the emerging role of signal transducers and activators of transcription in cancer.
miRNAs in melanoma: a defined role in tumor progression and metastasis.
Relation of immune semaphorin/plexin signaling to carcinogenesis.
Roles of RNA-binding proteins in immune diseases and cancer.
Tissue factor: a neglected role in cancer biology.
Modulation of Immune Responses by Platelet-Derived ADAM10.
Fusion between Intestinal epithelial cells and macrophages in a cancer context results in nuclear reprogramming.
Emerging roles of platelets in cancer biology and their potential as therapeutic targets.
Inflammation: what role in pediatric cancer?
The regulatory effect of hyaluronan on human mesenchymal stem cells' fate modulates their interaction with cancer cells in vitro.
Small-molecule compounds targeting the STAT3 DNA-binding domain suppress survival of cisplatin-resistant human ovarian cancer cells by inducing apoptosis.
CXCL13/CXCR5 signaling axis in cancer.
Clinical and experimental approaches to the pathophysiology of interleukin-18 in cancer progression.
Hijacking the Hexosamine Biosynthetic Pathway to Promote EMT-Mediated Neoplastic Phenotypes.

Tumor microenvironment and epithelial-mesenchymal transition in bladder cancer: Cytokines in the game?
Newly identified form of phenotypic plasticity of cancer: immunogenic mimicry.
Metastatic cells can escape the proapoptotic effects of TNF- $\alpha$ through increased autocrine IL-6/STAT3 signaling.
Short-term EGFR blockade enhances immune-mediated cytotoxicity of EGFR mutant lung cancer cells: rationale for combination therapies.
Extracellular Vesicles and Tumor-Immune Escape: Biological Functions and Clinical Perspectives.
Cryptotanshinone inhibits constitutive signal transducer and activator of transcription 3 function through blocking the dimerization in DU145 prostate cancer cells.
Nuclear morphometry, nucleomics and prostate cancer progression.
Role of circular RNAs in disease progression and diagnosis of cancers: An overview of recent advanced insights.
Immunosuppressive Signaling Pathways as Targeted Cancer Therapies.
Beyond the Oncogene Revolution: Four New Ways to Combat Cancer.
Extracellular vesicle-mediated EBAG9 transfer from cancer cells to tumor microenvironment promotes immune escape and tumor progression.
Roles of matrix metalloproteinases in tumor metastasis and angiogenesis.
Resistance of cancers to immunologic cytotoxicity and adoptive immunotherapy via X-linked inhibitor of apoptosis protein expression and coexisting defects in mitochondrial death signaling.
PI3K functions in cancer progression, anticancer immunity and immune evasion by tumors.
Host manipulation by cancer cells: Expectations, facts, and therapeutic implications.
Systems-Level Proteomics Evaluation of Microglia Response to Tumor-Supportive Anti-Inflammatory Cytokines.
AXL Receptor in Cancer Metastasis and Drug Resistance: When Normal Functions Go Askew.
Epithelial/mesenchymal plasticity: how have quantitative mathematical models helped improve our understanding?
Understanding and Targeting Apoptotic Pathways in Ovarian Cancer.
Tumorigenic bacteria in colorectal cancer: mechanisms and treatments.
Modulation of apoptotic pathways by human papillomaviruses (HPV): mechanisms and implications for therapy.
Tracking the fates of iron-labeled tumor cells in vivo using magnetic particle imaging.
Position of STAT-1 alpha in cycloheximide-dependent apoptosis triggered by TNF-alpha in human colorectal COLO 205 cancer cell line; role of polyphenolic compounds.
Cholangiocarcinoma: recent progress. Part 2: molecular pathology and treatment.
Modulation of the tumor microenvironment by Epstein-Barr virus latent membrane protein 1 in nasopharyngeal carcinoma.
Orthogonal targeting of osteoclasts and myeloma cells for radionuclide stimulated dynamic therapy induces multidimensional cell death pathways.
The regulatory and modulatory roles of TRP family channels in malignant tumors and relevant therapeutic strategies.
IL-4 Counteracts the Cytotoxic Effects of Peripheral Blood Mononuclear Cells on Hormone-sensitive Prostate Cancer Cells.
Single-Cell Technologies to Decipher the Immune Microenvironment in Myeloid Neoplasms: Perspectives and Opportunities.
Can killers be saviors?



DNA damage repair kinase DNA-PK and cGAS synergize to induce cancer-related inflammation in glioblastoma.
Mer Tyrosine Kinase Regulates Disseminated Prostate Cancer Cellular Dormancy.
Effect of crosstalk among conspirators in tumor microenvironment on niche metastasis of gastric cancer.
1,3,4-Oxadiazole: An Emerging Scaffold to Inhibit the Thymidine Phosphorylase as an Anticancer Agent.
Sialyltransferase inhibition leads to inhibition of tumor cell interactions with E-selectin, VCAM1, and MADCAM1, and improves survival in a human multiple myeloma mouse model.
Role of the inflammatory protein serine protease inhibitor Kazal in preventing cytolytic granule granzyme A-mediated apoptosis.
Abrogation of TLR4 and CD14 expression and signaling in human adrenocortical tumors.
Regulation of cellular immunity by activating transcription factor 4.
[VEGF-C and lymphatic vessels: a double-edged sword in tumor development and metastasis].
Cyclophilin A Inhibitor Debio-025 Targets Crk, Reduces Metastasis, and Induces Tumor Immunogenicity in Breast Cancer.
Enzymatic synthesis of non-natural trisaccharides and galactosides; Insights of their interaction with galectins as a function of their structure.
Dynamic switch of immunity and antitumor effects of metformin in rat spontaneous esophageal carcinogenesis.
Does the Use of the "Proseek(®) Multiplex Inflammation I Panel" Demonstrate a Difference in Local and Systemic Immune Responses in Endometriosis Patients with or without Deep-Infiltrating Lesions?
Complex roles of the old drug aspirin in cancer chemoprevention and therapy.
NOS1 inhibits the interferon response of cancer cells by S-nitrosylation of HDAC2.
Gain-of-'endocytic' function in mutant p53 cancer cells.
miR-34a induces immunosuppression in colorectal carcinoma through modulating a SIRT1/NF-κB/B7-H3/TNF-α axis.
Anti-inflammatory cytokines in endometriosis.
Sodium butyrate sensitizes human colon adenocarcinoma COLO 205 cells to both intrinsic and TNF-alpha-dependent extrinsic apoptosis.
The Polycomb Repressor Complex 1 Drives Double-Negative Prostate Cancer Metastasis by Coordinating Stemness and Immune Suppression.
Three-dimensional cell culture: a breakthrough in vivo.
Kaposi's sarcoma: a model of both malignancy and chronic inflammation.
Activation of pro-uPA is critical for initial escape from the primary tumor and hematogenous dissemination of human carcinoma cells.
Dissecting the signaling pathways that mediate cancer in PTEN and LKB1 double-knockout mice.
Unphosphorylated STAT1 promotes sarcoma development through repressing expression of Fas and bad and conferring apoptotic resistance.
The expanding role of innate lymphoid cells and their T-cell counterparts in gastrointestinal cancers.
Induced phase separation of mutant NF2 imprisons the cGAS-STING machinery to abrogate antitumor immunity.
The regulatory roles of calcium channels in tumors.
Pentoxifylline and the proteasome inhibitor MG132 induce apoptosis in human leukemia U937 cells through a decrease in the expression of Bcl-2 and Bcl-XL and phosphorylation of p65.
Malignant pirates of the immune system.

Kynurenic acid is a potent endogenous aryl hydrocarbon receptor ligand that synergistically induces interleukin-6 in the presence of inflammatory signaling.
Effect of celecoxib on capecitabine-induced hand-foot syndrome and antitumor activity.
Micro-masters of glioblastoma biology and therapy: increasingly recognized roles for microRNAs.
Cytokine-regulated Th17 plasticity in human health and diseases.
Heparan Sulfate and Heparan Sulfate Proteoglycans in Cancer Initiation and Progression.
Sustained compensatory p38 MAPK signaling following treatment with MAPK inhibitors induces the immunosuppressive protein CD73 in cancer: combined targeting could improve outcomes.
Merkel Cell Carcinoma and Immune Evasion: Merkel Cell Polyomavirus Small T-Antigen-Induced Surface Changes Can Be Reverted by Therapeutic Intervention.
Targeting bromodomain and extra-terminal proteins to inhibit neuroblastoma tumorigenesis through regulating MYCN.
Spatiotemporal view of malignant histogenesis and macroevolution via formation of polyploid giant cancer cells.
High-density neutrophils in MGUS and multiple myeloma are dysfunctional and immune-suppressive due to increased STAT3 downstream signaling.
Role of sialyl Lewis X in liver metastasis in view of liver-associated immunity.
Centrosome dysfunction contributes to chromosome instability, chromoanagenesis, and genome reprogramming in cancer.
Testing the theory of immune selection in cancers that break the rules of transplantation.
Pathological Role of HDAC8: Cancer and Beyond.
Heat Shock Proteins and PD-1/PD-L1 as Potential Therapeutic Targets in Myeloproliferative Neoplasms.
Targeting Opposing Immunological Roles of the Junctional Adhesion Molecule-A in Autoimmunity and Cancer.
VEGFR2 Promotes Metastasis and PD-L2 Expression of Human Osteosarcoma Cells by Activating the STAT3 and RhoA-ROCK-LIMK2 Pathways.
Beyond DNA Repair: DNA-PKcs in Tumor Metastasis, Metabolism and Immunity.
Cancer and meiotic gene expression: Two sides of the same coin?
The Taiman Transcriptional Coactivator Engages Toll Signals to Promote Apoptosis and Intertissue Invasion in Drosophila.
Multifunctional antitumor molecule 5'-triphosphate siRNA combining glutaminase silencing and RIG-I activation.
APE/Ref-1 makes fine-tuning of CD40-induced B cell proliferation.
O-GlcNAcylation in Chronic Lymphocytic Leukemia and Other Blood Cancers.
RAD21 amplification epigenetically suppresses interferon signaling to promote immune evasion in ovarian cancer.
Cancer cell transmission via the placenta.
Globular C1q Receptor (gC1qR/p32/HABP1) Suppresses the Tumor-Inhibiting Role of C1q and Promotes Tumor Proliferation in 1q21-Amplified Multiple Myeloma.
The Deadly Bite of STAT3.
Bisindolylmaleimide IX facilitates extrinsic and initiates intrinsic apoptosis in TNF-alpha-resistant human colon adenocarcinoma COLO 205 cells.
Epilipidomics of Senescent Dermal Fibroblasts Identify Lysophosphatidylcholines as Pleiotropic Senescence-Associated Secretory Phenotype (SASP) Factors.
Clinically relevant radioresistant rhabdomyosarcoma cell lines: functional, molecular and immune-related characterization.

Colorectal cancer cells induce lymphocyte apoptosis by an endothelial monocyte-activating polypeptide-II-dependent mechanism.
Poly Combs the Immune System: PRC2 Loss in Malignant Peripheral Nerve Sheath Tumors Can Dampen Immune Responses.
Immunology: glyco-engineering 'super-self'.
Carbohydrate Sulfation As a Mechanism for Fine-Tuning Siglec Ligands.
Glycosylation of Siglec15 promotes immunoescape and tumor growth.
Phosphorothioate-modified TLR9 ligands protect cancer cells against TRAIL-induced apoptosis.
Interleukin (IL)-18, a biomarker of human ovarian carcinoma, is predominantly released as biologically inactive precursor.
Epigenomic Reordering Induced by Polycomb Loss Drives Oncogenesis but Leads to Therapeutic Vulnerabilities in Malignant Peripheral Nerve Sheath Tumors.
Adhesion-mediated intracellular redistribution of c-Fas-associated death domain-like IL-1-converting enzyme-like inhibitory protein-long confers resistance to CD95-induced apoptosis in hematopoietic cancer cell lines.
Reversibility and recurrence of IGF-IR-induced mammary tumors.
Tetraspanins as therapeutic targets in hematological malignancy: a concise review.
Astrocytic Chitinase-3-like protein 1 in neurological diseases: Potential roles and future perspectives.
IFN unresponsiveness in LNCaP cells due to the lack of JAK1 gene expression.
Bortezomib restores stroma-mediated APO2L/TRAIL apoptosis resistance in multiple myeloma.
Role of CD38 Expression in Diagnosis and Pathogenesis of Chronic Lymphocytic Leukemia and Its Potential as Therapeutic Target.
Deubiquitinating enzyme OTUB1 in immunity and cancer: Good player or bad actor?
MSP-RON Pathway: Potential Regulator of Inflammation and Innate Immunity.
Decellularization of the mouse ovary: comparison of different scaffold generation protocols for future ovarian bioengineering.
Selenium, Selenoproteins, and Immunity.
STAG2 regulates interferon signaling in melanoma via enhancer loop reprogramming.
Why does the gut synthesize glucocorticoids?
Echinococcus multilocularis: inflammatory and regulatory chemokine responses in patients with progressive, stable and cured alveolar echinococcosis.
IL-18 enhances the migration ability of murine melanoma cells through the generation of ROI and the MAPK pathway.
Serum levels of the progesterone induced blocking factor do not precipitously rise in women with gynecologic cancer in contrast to women exposed to progesterone.
The Role of Peritoneal Immunity in Peritoneal Endometriosis and Related Infertility.
Extra-Adrenal Glucocorticoid Synthesis in the Intestinal Mucosa: Between Immune Homeostasis and Immune Escape.
The smoking gun: many conditions associated with tobacco exposure may be attributable to paradoxical compensatory autonomic responses to nicotine.
Publisher Correction: Landscape of B cell immunity and related immune evasion in human cancers.
Immunological dysfunction in chronic arsenic exposure: From subclinical condition to skin cancer.

Table A2-64, Cluster 63

Cluster 63 focuses on mechanisms associated with evasion of the host immune system by bacterial and viral pathogens (1301)
Molecular basis of host specificity in human pathogenic bacteria.
Anti-immunology: evasion of the host immune system by bacterial and viral pathogens.
Intracellular Pathogens: Host Immunity and Microbial Persistence Strategies.
Strategies Used by Bacteria to Grow in Macrophages.
Subversion of Macrophage Functions by Bacterial Protein Toxins and Effectors.
Intracellular innate resistance to bacterial pathogens.
Autophagy and burkholderia.
Remodeling of host glycoproteins during bacterial infection.
The Immunological Synapse: An Emerging Target for Immune Evasion by Bacterial Pathogens.
Out-Smarting the Host: Bacteria Maneuvering the Immune Response to Favor Their Survival.
Contrasting Lifestyles Within the Host Cell.
Neutrophils and Bacterial Immune Evasion.
Formation and Maturation of the Phagosome: A Key Mechanism in Innate Immunity against Intracellular Bacterial Infection.
Within-Host Envelope Remodelling and its Impact in Bacterial Pathogen Recognition.
Innate recognition of bacteria by a macrophage cytosolic surveillance pathway.
The Recruitment and Activation of Plasminogen by Bacteria-The Involvement in Chronic Infection Development.
Interactions of <i>Listeria monocytogenes</i> with the autophagy system of host cells.
Bacteria fighting back: how pathogens target and subvert the host innate immune system.
T Cell Immunity to Bacterial Pathogens: Mechanisms of Immune Control and Bacterial Evasion.
Interaction of <i>Chlamydiae</i> with human macrophages.
Modulation of the host innate immune and inflammatory response by translocated bacterial proteins.
<i>Shigella</i> are versatile mucosal pathogens that circumvent the host innate immune system.
Outsmarting the host: bacteria modulating the immune response.
Host-microbe communication within the GI tract.
A cross-disciplinary perspective on the innate immune responses to bacterial lipopolysaccharide.
From Gene to Protein-How Bacterial Virulence Factors Manipulate Host Gene Expression During Infection.
Manipulation of Host Cell Organelles by Intracellular Pathogens.
Subversion of inflammasome activation and pyroptosis by pathogenic bacteria.
Silent control: microbial plant pathogens evade host immunity without coding sequence changes.
Type VI Secretion System in Pathogenic <i>Escherichia coli</i> : Structure, Role in Virulence, and Acquisition.
The Mechanisms of Disease Caused by <i>Acinetobacter baumannii</i> .
Hijacking Host Cell Highways: Manipulation of the Host Actin Cytoskeleton by Obligate Intracellular Bacterial Pathogens.
Recognition of bacterial pathogens and mucosal immunity.
Urocanate as a potential signaling molecule for bacterial recognition of eukaryotic hosts.
From the Cradle to the Grave of an Infection: Host-Pathogen Interaction Visualized by Intravital Microscopy.
Within-host evolution decreases virulence in an opportunistic bacterial pathogen.
The Exploration of Complement-Resistance Mechanisms of Pathogenic Gram-Negative Bacteria to Support the Development of Novel Therapeutics.

To catch a thief: regulated RIPK1 post-translational modifications as a fail-safe system to detect and overcome pathogen subversion of immune signaling.
Beyond host-pathogen interactions: microbial defense strategy in the host environment.
Bacterial TIR-containing proteins and host innate immune system evasion.
Bacterial strategies for immune systems - Role of the type VI secretion system.
Subverting Toll-Like Receptor Signaling by Bacterial Pathogens.
Molecular battle between host and bacterium: recognition in innate immunity.
Bacterial glycans and their interactions with lectins in the innate immune system.
Shigella manipulates host immune responses by delivering effector proteins with specific roles.
Virulence factors of gram-negative bacteria in sepsis with a focus on Neisseria meningitidis.
Manipulation of the host cell death pathway by Shigella.
Glycointeractions in bacterial pathogenesis.
Masquerading microbial pathogens: capsular polysaccharides mimic host-tissue molecules.
Acinetobacter baumannii: An Ancient Commensal with Weapons of a Pathogen.
Modulation of innate lymphoid cells by enteric bacterial pathogens.
The induction of apoptosis by bacterial pathogens.
Origins of symbiosis: shared mechanisms underlying microbial pathogenesis, commensalism and mutualism of plants and animals.
Host inflammasome defense mechanisms and bacterial pathogen evasion strategies.
Immune Evasion Strategies of Pathogens in Macrophages: the Potential for Limiting Pathogen Transmission.
Innate immune encounters of the (Type) 4th kind: Brucella.
Pattern Recognition Receptors in Innate Immunity to Obligate Intracellular Bacteria.
Comparative phylogenomics and evolution of the Brucellae reveal a path to virulence.
Modulation of host immune defenses by Aeromonas and Yersinia species: convergence on toxins secreted by various secretion systems.
A Stenotrophomonas maltophilia Strain Evades a Major Caenorhabditis elegans Defense Pathway.
Apoptosis in infectious diseases as a mechanism of immune evasion and survival.
Avoidance and Subversion of Eukaryotic Homeostatic Autophagy Mechanisms by Bacterial Pathogens.
A review on host-pathogen interactions: classification and prediction.
Manipulation of rab GTPase function by intracellular bacterial pathogens.
Glycan-mediated molecular interactions in bacterial pathogenesis.
Pyroptosis modulation by bacterial effector proteins.
Extracellular Matrix Interactions with Gram-Positive Pathogens.
A New perspective: How Pathogens Manipulate Phagocytosis?
Chronic Bacterial Pathogens: Mechanisms of Persistence.
Role of innate immunity in respiratory mycoplasma infection.
Microbiota and Its Role on Viral Evasion: Is It With Us or Against Us?
Host-microbe interactions in the small bowel.
Extracellular Vesicles: Recent Insights Into the Interaction Between Host and Pathogenic Bacteria.
An Emerging Approach for Parallel Quantification of Intracellular Protozoan Parasites and Host Cell Characterization Using TissueFAXS Cytometry.
Structure and function: Lipid A modifications in commensals and pathogens.
Effector-triggered immunity and pathogen sensing in metazoans.
Antigenic Variation in Bacterial Pathogens.
The lipopolysaccharide core of Brucella abortus acts as a shield against innate immunity recognition.

Bacterial Evasion of Host Antimicrobial Peptide Defenses.
Role of Cyclic di-GMP in the Bacterial Virulence and Evasion of the Plant Immunity.
The ubiquitin system: a critical regulator of innate immunity and pathogen-host interactions.
The roles of microRNAs in regulation of autophagy during bacterial infection.
Innate Lymphoid Cells and Natural Killer Cells in Bacterial Infections: Function, Dysregulation, and Therapeutic Targets.
Mechanisms of Epithelial Immunity Evasion by Respiratory Bacterial Pathogens.
TH1/TH2 paradigm extended: macrophage polarization as an unappreciated pathogen-driven escape mechanism?
Persistent bacterial infections, antibiotic tolerance, and the oxidative stress response.
Host response to leptospira infection.
Serving the new masters - dendritic cells as hosts for stealth intracellular bacteria.
Acinetobacter baumannii phenylacetic acid metabolism influences infection outcome through a direct effect on neutrophil chemotaxis.
Male genital tract immune response against Chlamydia trachomatis infection.
Subterfuge and sabotage: evasion of host innate defenses by invasive gram-positive bacterial pathogens.
Fatal attraction: how bacterial adhesins affect host signaling and what we can learn from them.
Membrane rafts: a potential gateway for bacterial entry into host cells.
Bacterial effectors mimicking ubiquitin-proteasome pathway tweak plant immunity.
Identification of a peptide-pheromone that enhances Listeria monocytogenes escape from host cell vacuoles.
[Host Defense against Bacterial Infection and Bacterial Toxin-induced Impairment of Innate Immunity].
Host defense mechanisms against pathogens.
Emerging Evasion Mechanisms of Macrophage Defenses by Pathogenic Bacteria.
Host-Pathogen Interactions of Marine Gram-Positive Bacteria.
Identification of Brucella abortus virulence proteins that modulate the host immune response.
Neutrophils and aquatic pathogens.
Phytopathogen effectors subverting host immunity: different foes, similar battleground.
Macrophages in Microbial Pathogenesis: Commonalities of Defense Evasion Mechanisms.
Uncovering complex molecular networks in host-pathogen interactions using systems biology.
Vaccine development: From concept to early clinical testing.
Bacterial Metabolism Shapes the Host-Pathogen Interface.
Common Differences: The Ability of Inflammasomes to Distinguish Between Self and Pathogen Nucleic Acids During Infection.
Genetic Screen in Chlamydia muridarum Reveals Role for an Interferon-Induced Host Cell Death Program in Antimicrobial Inclusion Rupture.
Plant-Pathogen Effectors: Cellular Probes Interfering with Plant Defenses in Spatial and Temporal Manners.
Avoiding death by autophagy: interactions of Listeria monocytogenes with the macrophage autophagy system.
PANoptosis: A New Insight Into Oral Infectious Diseases.
The role of autophagy in intracellular pathogen nutrient acquisition.
The Bacterial and Viral Agents of BRDC: Immune Evasion and Vaccine Developments.
The well-evolved pathogen.

Manipulation of autophagy by bacteria for their own benefit.
Bacterial differentiation, development, and disease: mechanisms for survival.
The impact of Toll-like receptors on bacterial virulence strategies.
Systems Biology Analysis of Temporal In vivo Brucella melitensis and Bovine Transcriptomes Predicts host:Pathogen Protein-Protein Interactions.
Brucella spp. Virulence Factors and Immunity.
Challenges in Drug Discovery for Intracellular Bacteria.
Chlamydia psittaci: update on an underestimated zoonotic agent.
Community behavior and spatial regulation within a bacterial microcolony in deep tissue sites serves to protect against host attack.
Immune-evasion Strategies of Mycobacteria and Their Implications for the Protective Immune Response.
Subversion of innate immune responses by bacterial hindrance of NF- $\kappa$ B pathway.
The arsenal of virulence factors deployed by Listeria monocytogenes to promote its cell infection cycle.
Mechanisms of Bacterial Colonization of the Respiratory Tract.
Thermal control of virulence factors in bacteria: a hot topic.
The Listeria monocytogenes ChiA chitinase enhances virulence through suppression of host innate immunity.
Targeting of plant pattern recognition receptor-triggered immunity by bacterial type-III secretion system effectors.
Evolving Bacterial Envelopes and Plasticity of TLR2-Dependent Responses: Basic Research and Translational Opportunities.
Galectins in Host-Pathogen Interactions: Structural, Functional and Evolutionary Aspects.
Chlamydia Infection Across Host Species Boundaries Promotes Distinct Sets of Transcribed Anti-Apoptotic Factors.
Establishment of Chronic Infection: Brucella's Stealth Strategy.
The LOV protein of Xanthomonas citri subsp. citri plays a significant role in the counteraction of plant immune responses during citrus canker.
To con protection: TIR-domain containing proteins (Tcp) and innate immune evasion.
An evolutionary strategy for a stealthy intracellular Brucella pathogen.
The PathoChip, a functional gene array for assessing pathogenic properties of diverse microbial communities.
Role of mitochondria in regulating immune response during bacterial infection.
Immunology of the porcine respiratory disease complex.
Subversion of mouse dendritic cell subset function by bacterial pathogens.
Targeting Autophagy in Innate Immune Cells: Angel or Demon During Infection and Vaccination?
The Phagosome-Lysosome Fusion Is the Target of a Purified Quillaja saponin Extract (PQSE) in Reducing Infection of Fish Macrophages by the Bacterial Pathogen Piscirickettsia salmonis.
The sst1 resistance locus regulates evasion of type I interferon signaling by Chlamydia pneumoniae as a disease tolerance mechanism.
Group B streptococcal infections in the postgenomic era.
Bacterial inhibition of inflammatory responses via TLR-independent mechanisms.
Fibrinolysis and host response in bacterial infections.
TLR4-mediated podosome loss discriminates gram-negative from gram-positive bacteria in their capacity to induce dendritic cell migration and maturation.

Zooming in on common immune evasion mechanisms of pathogens in phagolysosomes: potential broad-spectrum therapeutic targets against infectious diseases.
A critical role for peptidoglycan N-deacetylation in <i>Listeria</i> evasion from the host innate immune system.
BtpB, a novel <i>Brucella</i> TIR-containing effector protein with immune modulatory functions.
MicroRNAs and bacterial infection.
Network-based analysis of virulence factors for uncovering <i>Aeromonas veronii</i> pathogenesis.
Speaking the language of lipids: the cross-talk between plants and pathogens in defence and disease.
Infection strategies of mycoplasmas: Unraveling the panoply of virulence factors.
CTLs: Killers of intracellular bacteria.
Bacterial pathogens activate plasminogen to breach tissue barriers and escape from innate immunity.
Sensing Gram-negative bacteria: a phylogenetic perspective.
Genomic analysis of <i>Acinetobacter baumannii</i> prophages reveals remarkable diversity and suggests profound impact on bacterial virulence and fitness.
Modified horseshoe crab peptides target and kill bacteria inside host cells.
Lipid Droplets: A Significant but Understudied Contributor of Host~Bacterial Interactions.
Microbial recognition and evasion of host immunity.
Screening of Virulence-Related Transcriptional Regulators in <i>Streptococcus suis</i> .
Intracellular lifestyles and immune evasion strategies of uropathogenic <i>Escherichia coli</i> .
Persistent bacterial infections and persister cells.
Incoming pathogens team up with harmless 'resident' bacteria.
The Chlamydia effector CpoS modulates the inclusion microenvironment and restricts the interferon response by acting on Rab35.
Immune interactions between <i>Drosophila</i> and the pathogen <i>Xenorhabdus</i> .
The role of NLRP3 and AIM2 in inflammasome activation during <i>Brucella abortus</i> infection.
When the Going Gets Rough: The Significance of <i>Brucella</i> Lipopolysaccharide Phenotype in Host-Pathogen Interactions.
RAB11-mediated trafficking in host-pathogen interactions.
Bacteriophage-encoded bacterial virulence factors and phage-pathogenicity island interactions.
Surface polysaccharides enable bacteria to evade plant immunity.
Neutrophil apoptosis: relevance to the innate immune response and inflammatory disease.
Progress and obstacles in vaccine development for the ehrlichioses.
Proteomic Approaches to Unravel Mechanisms of Antibiotic Resistance and Immune Evasion of Bacterial Pathogens.
Macrophages Infected by a Pathogen and a Non-pathogen Spotted Fever Group Rickettsia Reveal Differential Reprogramming Signatures Early in Infection.
Comparative genomics analysis and virulence-related factors in novel <i>Aliarcobacter faecis</i> and <i>Aliarcobacter lanthieri</i> species identified as potential opportunistic pathogens.
Intrinsic disorder in pathogen effectors: protein flexibility as an evolutionary hallmark in a molecular arms race.
Review of CRISPR-Cas Systems in <i>Listeria</i> Species: Current Knowledge and Perspectives.
Cell death programs in <i>Yersinia</i> immunity and pathogenesis.
Epigenetic regulation in bacterial infections: targeting histone deacetylases.
Molecular mechanisms of bacterial pathogenicity.
Nanocarriers for the delivery of antibiotics into cells against intracellular bacterial infection.
Interactions of the human pathogenic <i>Brucella</i> species with their hosts.



The bacterial pathogen <i>Listeria monocytogenes</i> and the interferon family: type I, type II and type III interferons.
Chlamydia and Its Many Ways of Escaping the Host Immune System.
Immunopathogenesis of streptococcal deep tissue infections.
<i>Edwardsiella piscicida</i> YccA: A novel virulence factor essential to membrane integrity, mobility, host infection, and host immune response.
LAPped in Proof: LC3-Associated Phagocytosis and the Arms Race Against Bacterial Pathogens.
Epigenetics and miRNA during bacteria-induced host immune responses.
LC3-Associated Phagocytosis in Bacterial Infection.
Using Epidemiology, Immunology, and Genomics to Study the Biology of <i>Chlamydia trachomatis</i> .
The inflammasome: Learning from bacterial evasion strategies.
Electroporation of functional bacterial effectors into mammalian cells.
Gut commensal <i>Enterocloster</i> species host inoviruses that are secreted in vitro and in vivo.
The <i>Legionella pneumophila</i> EnhC protein interferes with immunostimulatory muramyl peptide production to evade innate immunity.
Reorganization of the host cytoskeleton by the intracellular pathogen <i>Chlamydia trachomatis</i> .
Type VI secretion and anti-host effectors.
Proteinaceous effector discovery and characterization in filamentous plant pathogens.
An overview on phase variation, mechanisms and roles in bacterial adaptation.
GAPDH, as a virulence factor.
Cas9-dependent endogenous gene regulation is required for bacterial virulence.
The highest priority: what microbial genomes are telling us about immunity.
Pathogens MentORing Macrophages and Dendritic Cells: Manipulation of mTOR and Cellular Metabolism to Promote Immune Escape.
Immune subversion and quorum-sensing shape the variation in infectious dose among bacterial pathogens.
Who's really in control: microbial regulation of protein trafficking in the epithelium.
Recognition of Intracellular Bacteria by Inflammasomes.
Subversion of the Immune Response by Human Pathogenic Mycoplasmas.
Inflammasome/IL-1 $\beta$ Responses to Streptococcal Pathogens.
Beyond the antibody: B cells as a target for bacterial infection.
Dendritic cell development in infection.
Advances in the study of myeloid-derived suppressor cells in infectious lung diseases.
Antigenic variation and transmission fitness as drivers of bacterial strain structure.
The surface lipoproteins of gram-negative bacteria: Protectors and foragers in harsh environments.
Bacterial Pathogens Hijack the Innate Immune Response by Activation of the Reverse Transsulfuration Pathway.
Melioidosis and glanders modulation of the innate immune system: barriers to current and future vaccine approaches.
Epstein Barr virus-induced 3 (EBI3) together with IL-12 negatively regulates T helper 17-mediated immunity to <i>Listeria monocytogenes</i> infection.
Microbial manipulation of receptor crosstalk in innate immunity.
Bacterial secreted effectors and caspase-3 interactions.
<i>Brucella</i> evasion of adaptive immunity.
The bacterial metabolite 2,3-butanediol ameliorates endotoxin-induced acute lung injury in rats.
Do symbiotic bacteria subvert host immunity?

Activation and pathogenic manipulation of the sensors of the innate immune system.
Identification and Preliminary Characterization of Novel Type III Secreted Effector Proteins in <i>Chlamydia trachomatis</i> .
Purinergic modulation of the immune response to infections.
Protein glycosylation: Sweet or bitter for bacterial pathogens?
Neutrophils are essential for containment of <i>Vibrio cholerae</i> to the intestine during the proinflammatory phase of infection.
<i>Chlamydia</i> Lipooligosaccharide Has Varied Direct and Indirect Roles in Evading both Innate and Adaptive Host Immune Responses.
A new avenue to investigate: the autophagic process. From Crohn's disease to <i>Chlamydia</i> .
Reverse protein arrays applied to host-pathogen interaction studies.
The role of the <i>vacB</i> gene in the pathogenesis of <i>Brucella abortus</i> .
Evasion and interference: intracellular pathogens modulate caspase-dependent inflammatory responses.
Neutrophil serine proteases in antibacterial defense.
Host immunity and cellular responses to bacterial outer membrane vesicles.
Analysis of host-pathogen gene association networks reveals patient-specific response to streptococcal and polymicrobial necrotising soft tissue infections.
The Role of Glycosylation in Infectious Diseases.
An Update on Host-Pathogen Interplay and Modulation of Immune Responses during <i>Orientia tsutsugamushi</i> Infection.
A Bittersweet Kiss of Gram-Negative Bacteria: The Role of ADP-Heptose in the Pathogenesis of Infection.
Mechanisms of pathogen entry through the endosomal compartments.
The function of plant PR1 and other members of the CAP protein superfamily in plant-pathogen interactions.
A CRISPR/Cas system mediates bacterial innate immune evasion and virulence.
Modeling of variables in cellular infection reveals CXCL10 levels are regulated by human genetic variation and the <i>Chlamydia</i> -encoded CPAF protease.
Recent Advances on the Innate Immune Response to <i>Coxiella burnetii</i> .
Fibrin and fibrinolysis in infection and host defense.
Commensal and Pathogenic Biofilms Alter Toll-Like Receptor Signaling in Reconstructed Human Gingiva.
Avoiding the trap: Mechanisms developed by pathogens to escape neutrophil extracellular traps.
Genome sequencing and comparative analysis of three <i>Chlamydia pecorum</i> strains associated with different pathogenic outcomes.
A CRISPR-Cas system enhances envelope integrity mediating antibiotic resistance and inflammasome evasion.
Infection, Transmission, Pathogenesis and Vaccine Development against <i>Mycoplasma gallisepticum</i> .
Non-traditional Antibacterial Therapeutic Options and Challenges.
The role of lipids in host microbe interactions.
Extracellular vesicles and infectious diseases: new complexity to an old story.
Transcriptional portrait of <i>Actinobacillus pleuropneumoniae</i> during acute disease--potential strategies for survival and persistence in the host.
Herpes Simplex Virus Type 1 and Other Pathogens are Key Causative Factors in Sporadic Alzheimer's Disease.

Breathing life into pathogens: the influence of oxygen on bacterial virulence and host responses in the gastrointestinal tract.
Pathogen manipulation of cIL-10 signaling pathways: opportunities for vaccine development?
Cells within cells: Rickettsiales and the obligate intracellular bacterial lifestyle.
A systems biological view of intracellular pathogens.
Views of immunology: effector T cells.
Francisella novicida pathogenicity island encoded proteins were secreted during infection of macrophage-like cells.
Understanding GroEL and DnaK Stress Response Proteins as Antigens for Bacterial Diseases.
The coordination of anti-phage immunity mechanisms in bacterial cells.
CD4+ T cell-derived IL-10 promotes Brucella abortus persistence via modulation of macrophage function.
Evasion of autophagy mediated by Rickettsia surface protein OmpB is critical for virulence.
Extracellular vesicles: An emerging platform in gram-positive bacteria.
Comparative analysis of the growth and biological activity of a respiratory and atheroma isolate of Chlamydia pneumoniae reveals strain-dependent differences in inflammatory activity and innate immune evasion.
Type 1 pilated uropathogenic Escherichia coli hijack the host immune response by binding to CD14.
Infection in a dish: high-throughput analyses of bacterial pathogenesis.
Metabolic inhibitors of bacterial glycan biosynthesis.
A Fluorescence-based Method to Study Bacterial Gene Regulation in Infected Tissues.
A 'pathogenic needle' in a 'commensal haystack': Genetic virulence signatures of Corynebacterium glucuronolyticum that may drive its infectious propensity for the male urogenital system.
Immune evasion strategies of major tick-transmitted bacterial pathogens.
T cell responses to Chlamydia.
The Role of the Host in Driving Phenotypic Heterogeneity in Salmonella.
Evolution of bopA Gene in Burkholderia: A Case of Convergent Evolution as a Mechanism for Bacterial Autophagy Evasion.
Virulence-targeted Antibacterials: Concept, Promise, and Susceptibility to Resistance Mechanisms.
Interaction between Intracellular Bacterial Pathogens and Host Cell Mitochondria.
Peptidoglycan Deacetylases in Bacterial Cell Wall Remodeling and Pathogenesis.
Processivity in Bacterial Glycosyltransferases.
Toward the use of genomics to study microevolutionary change in bacteria.
Mini-review: Strategies for Variation and Evolution of Bacterial Antigens.
CRISPR-Cas systems: new players in gene regulation and bacterial physiology.
Molecular strategies used by fish pathogens to interfere with host-programmed cell death.
Evasion of inflammasome activation by microbial pathogens.
Listeria motility increases the efficiency of epithelial invasion during intestinal infection.
Dissecting the immune response to the entomopathogen Photobacterium.
Antigenic escape selects for the evolution of higher pathogen transmission and virulence.
Leptospira and inflammation.
Principles of Vaccination.
Persistence Alters the Interaction between Chlamydia trachomatis and Its Host Cell.
Chlamydia trachomatis Lipopolysaccharide Evades the Canonical and Noncanonical Inflammatory Pathways To Subvert Innate Immunity.

microRNAs in human brucellosis: A promising therapeutic approach and biomarker for diagnosis and treatment.
Immune evasion through Toll-like receptor 4: The role of the core oligosaccharides from $\alpha$ 2-Proteobacteria atypical lipopolysaccharides.
Phase and antigenic variation in bacteria.
Host response to Brucella infection: review and future perspective.
Interplay of host-pathogen microvesicles and their role in infectious disease.
Bordetella evades the host immune system by inducing IL-10 through a type III effector, BopN.
Within-host evolution of a gut pathobiont facilitates liver translocation.
Immune evasion by pathogens of bovine respiratory disease complex.
Repression of bacterial lipoprotein production by Francisella novicida facilitates evasion of innate immune recognition.
ALV-J strain SCAU-HN06 induces innate immune responses in chicken primary monocyte-derived macrophages.
Three small molecule pan activator families of Ras-related GTPases.
Trichomoniasis: What's New?
Nucleic Acid Nanostructure Assisted Immune Modulation.
The immune response to Naegleria fowleri amebae and pathogenesis of infection.
The adenovirus E3-6.7K protein adopts diverse membrane topologies following posttranslational translocation.
Bartonella infection in immunocompromised hosts: immunology of vascular infection and vasoproliferation.
Proinflammatory cytokine expression by Theileria annulata infected cell lines correlates with the pathology they cause in vivo.
Zwitterionic 3D-Printed Non-Immunogenic Stealth Microrobots.
Hide and Seek: Nanomaterial Interactions With the Immune System.
Non-pharmaceutical interventions and the emergence of pathogen variants.
The effects of environmental factors on the virulence of Trichomonas vaginalis.
Exploring the Immune-Boosting Functions of Vitamins and Minerals as Nutritional Food Bioactive Compounds: A Comprehensive Review.
B-1 cell-mediated modulation of M1 macrophage profile ameliorates microbicidal functions and disrupt the evasion mechanisms of Encephalitozoon cuniculi.
Light-triggered switching of liposome surface charge directs delivery of membrane impermeable payloads in vivo.
Stabilized albumin coatings on engineered xenografts for attenuation of acute immune and inflammatory responses.
Inhibitory ITAMs: a matter of life and death.
Phenotypic alterations and survival of monocytes following infection by human herpesvirus-6.
Differential response of porcine immature monocyte-derived dendritic cells to virulent and inactivated transmissible gastroenteritis virus.
Weil's Disease-Immunopathogenesis, Multiple Organ Failure, and Potential Role of Gut Microbiota.
Siglec receptors and hiding plaques in Alzheimer's disease.
Hybrid Nanogels: Stealth and Biocompatible Structures for Drug Delivery Applications.
A Comparative Analysis of Edwardsiella tarda-Induced Transcriptome Profiles in RAW264.7 Cells Reveals New Insights into the Strategy of Bacterial Immune Evasion.
Toward a comparison of microelectrodes for acute and chronic recordings.

The <i>Haemophilus ducreyi</i> cytolethal distending toxin activates sensors of DNA damage and repair complexes in proliferating and non-proliferating cells.
Comparative Pathogenesis, Genomics and Phylogeography of Mousepox.
Cell targeting by a generic receptor-targeted polymer nanocontainer platform.
Isolation and characterization of a neprilysin-like protein from <i>Venturia canescens</i> virus-like particles.
<i>Trichinella pseudospiralis</i> -secreted 53 kDa protein ameliorates imiquimod-induced psoriasis by inhibiting the IL-23/IL-17 axis in mice.
Microbial evasion of the immune system: structural modifications of enterobactin impair siderocalin recognition.
The clinical impact of glycobiology: targeting selectins, Siglecs and mammalian glycans.
Exploring the functions of polymers in adenovirus-mediated gene delivery: Evading immune response and redirecting tropism.
Correction: Bacterial immune evasion through manipulation of host inhibitory immune signaling.
Mechanisms by which mastitis affects reproduction in dairy cow: A review.
Vaccination with <i>Strongyloides ratti</i> heat shock protein 60 increases susceptibility to challenge infection by induction of Th1 response.
Soluble Immune Response Suppressor (SIRS): Reassessing the immunosuppressant potential of an elusive peptide.
The Gut-Immune-Brain Axis: An Important Route for Neuropsychiatric Morbidity in Inflammatory Bowel Disease.
Deficiency of Selected Cathepsins Does Not Affect the Inhibitory Action of ECTV on Immune Properties of Dendritic Cells.
Phylogenomics of nontuberculous mycobacteria respiratory infections in people with cystic fibrosis.
Invasion and destruction of mucosal plasma cells by <i>Tropheryma whippelii</i> .
Iron metabolism disorders of patients with chronic paracoccidioidomycosis.
Distinct Transcriptional Profile of PDZ Genes after Activation of Human Macrophages and Dendritic Cells.
Impact of growth pH and glucose concentrations on the CodY regulatory network in <i>Streptococcus salivarius</i> .
<i>Treponema pallidum</i> promoted microglia apoptosis and prevented itself from clearing by human microglia via blocking autophagic flux.
Duchenne Muscular Dystrophy Gene therapy.
Cloning, expression, purification, crystallization and preliminary crystallographic studies of BceC, a UDP-glucose dehydrogenase from <i>Burkholderia cepacia</i> IST408.
Immune evasion by designer microrobots.
The sound of silence.
Low molecular-weight chitosan as a pH-sensitive stealth coating for tumor-specific drug delivery.
This Is the End: Regulation of Rab7 Nucleotide Binding in Endolysosomal Trafficking and Autophagy.
'There and back again': revisiting the pathophysiological roles of human endogenous retroviruses in the post-genomic era.
New frontiers in gene targeting and cloning: success, application and challenges in domestic animals and human embryonic stem cells.
Post-traumatic anxiety associates with failure of the innate immune receptor TLR9 to evade the pro-inflammatory NFkB pathway.
Mycotoxins: Emerging toxic mechanisms, and unanswered research questions.
Corrigendum: Advancements in understanding the molecular and immune mechanisms of <i>Bartonella</i> pathogenicity.

[Current research on picornavirus 3C protease].
Stealth Iron Oxide Nanoparticles for Organotropic Drug Targeting.
Differences in Immune Defense Evasion of Selected Inbred Lines of <i>Heterorhabditis</i> Bacteriophora in Two White Grub Species.
A Higher Abundance of O-Linked Glycans Confers a Selective Advantage to High Fertile Buffalo Spermatozoa for Immune-Evasion From Neutrophils.
Biobehavioral Mechanisms Associated With Nonhealing Wounds and Psychoneurologic Symptoms (Pain, Cognitive Dysfunction, Fatigue, Depression, and Anxiety) in Older Individuals With Chronic Venous Leg Ulcers.
Sensing their plasma membrane curvature allows migrating cells to circumvent obstacles.
Effect of partial PEGylation on particle uptake by macrophages.
Questioning the Use of PEGylation for Drug Delivery.
Tuft cells are key mediators of interkingdom interactions at mucosal barrier surfaces.
Inactivated tetanus as an immunological smokescreen: A major step towards harnessing tetanus-based therapeutics.
Intravenous hemostats: challenges in translation to patients.
Effects of mRNA Modifications on Translation: An Overview.
Hypotheses on the evolution of hyaluronan: a highly ionic acid.
An update on T-2 toxin and its modified forms: metabolism, immunotoxicity mechanism, and human exposure assessment.
Is <i>Necator americanus</i> approaching a mutualistic symbiotic relationship with humans?
Therapeutic efficacy of a systemically delivered oncolytic adenovirus - biodegradable polymer complex.
Design of Engineered Cyclodextrin Derivatives for Spontaneous Coating of Highly Porous Metal-Organic Framework Nanoparticles in Aqueous Media.
So many ways of getting in the way: diversity in the molecular architecture of superantigen-dependent T-cell signaling complexes.
A Bioartificial Pancreas with "Immune Stealth" And Continuous Oxygen Supply for Islet Transplantation.
Pentapeptide sharing between <i>Corynebacterium diphtheria</i> toxin and the human neural protein network.
ADP-ribosylation in evasion, promotion and exacerbation of immune responses.
How Corona Formation Impacts Nanomaterials as Drug Carriers.
[Septic arthritis: what is the role for the rheumatologist?].
Pollen-Inspired Shell-Core Aerosol Particles Capable of Brownian Motion for Pulmonary Vascularization.
Medical and Veterinary Importance of the Moonlighting Functions of Triosephosphate Isomerase.
Primary Amoebic Meningoencephalitis: Neurochemotaxis and Neurotropic Preferences of <i>Naegleria fowleri</i> .
<i>Acanthamoeba castellanii</i> Genotype T4: Inhibition of Proteases Activity and Cytopathic Effect by Bovine Apo-Lactoferrin.
Immunohistochemical study of Langerhans cells in cutaneous lesions of the Jorge Lobo's disease.
Resistance to granzyme B-mediated cytochrome c release in Bak-deficient cells.
The Entomopathogenic Nematodes <i>H. bacteriophora</i> and <i>S. carpocapsae</i> Inhibit the Activation of proPO System of the Nipa Palm Hispid <i>Octodonta nipae</i> (Coleoptera: Chrysomelidae).
Application of polyglycerol coating to plasmid DNA lipoplex for the evasion of the accelerated blood clearance phenomenon in nucleic acid delivery.

An oral commensal associates with disease: chicken, egg, or red herring?
Cholesterol sulfate is a DOCK2 inhibitor that mediates tissue-specific immune evasion in the eye.
The link between immunity, autoimmunity and endometriosis: a literature update.
[Research progress in immunopathogenesis of cystic echinococcosis].
Evaluation of the innate immune response in pups during canine parvovirus type 1 infection.
Zwitterionic PEG-PC Hydrogels Modulate the Foreign Body Response in a Modulus-Dependent Manner.
Immunophysiology and pathology of inflammation in the testis and epididymis.
The presence of teleost-type angiotensin components in lamprey buccal gland suggests a role in endocrine mimicry.
Chemokinesis-driven accumulation of active colloids in low-mobility regions of fuel gradients.
Study on the allosteric activation mechanism of SHP2 via elastic network models and neural relational inference molecular dynamics simulation.
Gelatin carriers for drug and cell delivery in tissue engineering.
Immunomodulatory function of seminal catecholamines may be an adaptation for reproduction.
Sialylation: fate decision of mammalian sperm development, fertilization, and male fertility†.
Immunotherapy of Equine Sarcoids-From Early Approaches to Innovative Vaccines.

### Appendix 3 – Contributing Factors to Immune System Dysfunction

This appendix contains the methodology for retrieving dysfunctional immune system CFs from our 2020 monograph (<https://repository.gatech.edu/entities/publication/a5c9c8e9-8cc9-4588-a065-2be90b43b8dc>), as well as a listing of these CFs.

#### METHODOLOGY

The methodology, reproduced from Chapter 2 of the monograph, is as follows:

“The main analytical objective of this study is to identify factors that contribute to weakening of the immune system. Toward this end, a query was generated that would retrieve records with high probability of containing such factors. The query retrieved 10,733 records from Medline. Some additional records (identified from those records retrieved with the query) were also used for the analysis. The retrieved records were inspected visually, and contributing factors were extracted manually.....The query used to retrieve these records from SCI-Medline is shown boxed below.”

#### FINAL QUERY USED FOR STUDY SCI-MEDLINE

MESH--- (/adverse effects OR /toxicity OR /poisoning) AND TITLE--- (immun\* AND (expos\* OR induc\* OR exacerbat\* OR aggravat\* OR trigger\* OR disrupt\* OR impair\* OR dysfunction\* OR suppress\*))

OR

MESH--- (/adverse effects OR /toxicity OR /poisoning)

AND

TITLE--- immun\*

AND

TOPIC--- IMMUNOTOXIC\*

OR

TITLE--- IMMUNOTOXIC\* OR IMMUN\* NEAR/3 IMPAIR\*



“Initially, a much broader query was developed, and it retrieved tens of thousands of records. However, given that the projected analysis would be labor intensive, with visual inspection of the retrieved records and manual extraction of contributing factors, it was decided to generate a more focused query that would retrieve fewer records, but with higher probability of relevance (precision). The boxed query achieved that goal.”

## SUMMARY RESULTS

The following is a reproduction of Chapter 3 of the 2020 monograph, which contains the summary results. The numbering of tables is that used in the monograph.

“Approximately 65% of the retrieved records were evaluated. The extracted contributing factors were assigned to one of five categories (Lifestyle, Iatrogenic, Biotoxin/Biomaterial, Occupational/Environmental, Psychosocial/Socioeconomic), even though many of the contributing factors could have been assigned to multiple categories. Table 3-1 contains contributing factors that have been shown repeatedly to weaken the immune system. They include:

TABLE 3-1

### HIGH-FREQUENCY CONTRIBUTING FACTORS TO WEAKENED IMMUNE SYSTEM

CATEGORY	CONTRIBUTING FACTOR
LIFESTYLE	smoking, excess alcohol, substance abuse, high-fat diet, protein-deficient diet, high-cholesterol diet, Western-style diets, chronic sleep restriction, etc.
IATROGENIC	immunosuppressive drugs, gamma radiation treatments, nanomedicinal products, adjuvanted vaccines, acetaminophen, non-steroidal antiinflammatory drugs (NSAIDs), surgical stress, serotonin reuptake inhibitors, selected anesthetics, selected antibiotics, highly active antiretroviral therapy drugs, etc.
BIOTOXIN/ BIOMATERIAL	aflatoxin, ochratoxin, T-2 toxin, anatoxin-A, mycotoxins, microcystin-LR, dietary toxic cyanobacteria, yessotoxin, scorpion venom; Streptomyces californicus; Pseudomonas aeruginosa; Rhinovirus, respiratory syncytial virus, etc.
OCCUPATIONAL/ ENVIRONMENTAL	microplastics, endocrine-disrupting chemicals, heavy metals, pesticides/insecticides/herbicides, nanoparticles, perfluorooctanoic acid (PFOA), polychlorinated biphenyls (PCBs), polyaromatic hydrocarbons (PAHs), perfluorooctanesulfonate (PFOS), fine particulate matter, air pollution, acrylamide, aromatic halogenated disinfection byproducts, benzene, benzo(a)pyrene, crude oil, corexit, ultraviolet (UV) radiation, wireless radiation-cell phones/cell towers/WiFi, sodium fluoride, etc.
PSYCHOSOCIAL/ SOCIOECONOMIC	depression, chronic stress, restraint stress, social isolation, stressful life events, childhood adversity, etc.

Eliminating/ameliorating these toxic exposures/behaviors will require a combination of individual motivations/efforts and government efforts, especially at the regulatory level. The factors in the Lifestyle category mainly require motivation and willpower to eliminate, although government regulation would be beneficial for controlling food additives and labelling contents of processed foods. For the Iatrogenic category, government regulation is necessary for ensuring treatment safety. There is room for individual motivation in eliminating excessive or unnecessary use of painkillers, such as NSAIDs or opioids, and unnecessary/elective surgeries.

Members of the Biotoxin/Biomaterial category (especially the Biotoxin component) are more difficult for individuals to eliminate. As we are seeing with COVID-19, virus exposure is difficult to control (as is bacterial exposure). There are many mycotoxins listed in Table 3-1. Those found in food may result from improper storage and insufficient processing to eliminate mycotoxins. Those in indoor environments may result from insufficient moisture/humidity control. Some of these problems can be addressed by stricter government regulations.

The Occupational/Environmental category could benefit substantially from more rigorous government regulation. Most of the exposures are beyond the control of the individual; in fact, the individual most probably doesn't know he/she is being exposed to these substances.

For example, the Occupational Safety and Health Administration (OSHA) has responsibility for regulating most workplace toxic exposures. Out of the more than 85,000 chemicals registered with the EPA, OSHA issues Federally-enforceable Permissible Exposure Limits (PELs) for about 500 of these chemicals. In 2018, the first author published a study of the adequacy of OSHA's PELs [20], using a sampling technique. Of those substances that were sampled, their PELs were one-four orders of magnitude higher than exposures shown in the biomedical literature to cause damage.

As another example, the radiation exposure limits for wireless radiation (cell phone/cell towers/WiFi, etc.) approved by the FCC are from three-six orders of magnitude higher than exposures shown in the biomedical literature to cause damage, the discrepancy varying with the level of damage [21].

But, even in this category, individual choice and motivation play a role. People who want to strengthen their immune systems can choose (especially in the home environment, and partly in the work environment) to reduce exposure to wireless radiation, water with sodium fluoride, strong pesticides, strong disinfectants, etc.

The PsychoSocial/SocioEconomic category could benefit from some government interventions that reduce stressful situations for the individual (e.g., providing economic/health/occupational security, providing more protections for the most vulnerable (very young, elderly, disabled), etc.). Some of the types of adverse events and stresses are beyond the control of government or the individual, but here again, the individual can take steps to improve their responses to many of these types of stress.

Most of the studies, especially the laboratory studies, that gathered the data reported in Table 3-1 and the more extensive [Table A4-1](#) (shown in [Appendix 4](#)) focused on the effects of one stressor/toxic stimulus in isolation. In real-life, people are exposed to myriad toxic stimuli, both in parallel and over different time periods. Typically, when toxic stimuli are combined, less of each constituent of the combination is required to cause damage compared to the amount determined from single stressor

experiments [22]. Thus, real-life weakening of the immune system from myriad exposures to the toxic stimuli in Table 3-1 (and Table A4-1) is far greater than that reported in the single toxic stimulus studies.

To summarize, repurposed (mainly) antiviral treatments can only be expected to have very limited results in controlling SARS-CoV-2 viral load of the most severely impacted, based on trials conducted so far. Further, based on the findings of [Appendix 3](#), it is difficult to see how safe COVID-19 vaccines can be developed and fully tested on time scales of one or two years, as proposed presently.

Finally, the only real protection against a future COVID-19 pandemic or any other viral pandemic is the one that was demonstrated to work in the SARS, MERS, COVID-19 and annual influenza pandemics: a healthy immune system capable of neutralizing incoming viruses as nature intended. We need an Operation Warp Speed (the Federal Administration Program currently targeted to produce a vaccine in a record short time period in the USA) to identify and eliminate those factors that weaken the immune system as thoroughly, comprehensively, and rapidly as possible!"

## DETAILED RESULTS

Appendix 4 of the 2020 monograph is reproduced in the following section. The numbering is that used in the 2020 monograph. A4-A lists the CFs to a dysfunctional immune system, and A4-B lists the titles of the records from which the CFs were derived.

### "A4-A. Contributing Factors to Weakened Immune System

This section presents the complete list of factors that contribute to weakening of the immune system based on our analysis of the retrieved data base. About 10,800 records were retrieved using the query shown in Chapter 2 (and other records identified in the course of the analysis), about 7,000 of these records were analyzed, and about 4,700 (of the 7,000) records identified factors that contributed to weakening of the immune system. The contributing factors from these ~4,700 records were aggregated, and slightly over 2,200 contributing factors were entered into the table below. Not all redundancies were eliminated, but if even on the order of 30% redundancy exists, this means that about 1,500 contributing factors to weakening the immune system have been identified.

This is a very conservative estimate. The query in [Chapter 2](#) was written to retrieve those articles most relevant to immune-weakening factors, given the labor-intensive approach taken to identify these factors from the retrieved articles. A much more expansive query, albeit with less precision, would have retrieved tens of thousands of articles, and produced many more contributing factors.

But, the 1,500 constitute a large number, are far more than we have seen in any one article, and probably cover all the major contributing factors. Any measures that an individual or government regulator can take for protecting the health of the immune system will only be able to deal with a small fraction of the 1,500 already identified, at least initially.

**Commented [MA1]:** Make a reference to the Table in this section, and also qualify that though comprehensive it is unlikely to include all all factors

TABLE A4-1

## CONTRIBUTING FACTORS TO WEAKENED IMMUNE SYSTEM

LIFESTYLE	
advanced glycation end products	high-cholesterol diet
age	high-fat-diet
alcohol abuse	hygiene hypothesis
aniline-adulterated rapeseed oil	ice cream
Argemone oil	isobutyl nitrite
Biomass smoke exposure	isoflavones
Caesarian delivery	linoleic acid
cannabinoid	linolenic acid
cheese	Low dietary n – 3 fatty acid
chronic Sleep Restriction	Low prenatal-neonatal animal bacterial exposures
cigarette smoking	Maize prolamins
cinnamaldehyde	Malnutrition
cocaine	Maternal separation in early life
cow's milk insulin	Maternal smoking
crack	Maternal undernutrition
dietary biotin deficiency	medication use
dietary excess lipids	methamphetamine
dietary fatty acid intake	methylenedioxymethamphetamine
dietary Iron Deficiency	Microbial depletion in early environment
dietary iron excess	minced pork
dietary low lipids	minced turkey
dietary niacin deficiency	mobile phone
dietary oxidized lipids	morphine dependence
dietary phosphorus deficiency	n-propyl gallate
dietary protein restriction	Obese pregnancy
dietary pyridoxine deficiency	opioid
dietary thiamin deficiency	paracetamol
dietary valine deficiency	peanuts
Dietary zinc deficiency	peppermint oil
diet-induced obesity	phencyclidine
drug abuse	phloretin-contained fruits
Early antibiotic use	Prenatal nutrient energy overload
Early life diet	Prenatal nutrient-insufficiency
Early weaning stress	Protein malnutrition
E-cigarette	protein-deficient diet
Ecstasy	recreational drugs
ethanol consumption	rectal insemination
ethanol intoxication	Red meat consumption
excess dietary cholesterol	Reduced neonatal microbial environment
excessive Se	sedentary

extended contact lens wearing	Selenium deficiency
fast-food burger	snuff
fatty acid anilides	Sodium nitrite
fetal alcohol exposure	synthetic food additive antioxidants
fish extracts	Tetrahydrocannabinol
fish oil supplementation	tobacco smoking
Flavored e-cigarette liquids	traffic pollution
folic acid deficiency	vitamin A deficiency
formula feeding	vitamin B-12 deficiency
genistein	vitamin B-6 deficiency
Glue solvent	vitamin C deficiency
gluten	vitamin D deficiency
heavy competition	Western-style diets
heavy training	Wheat Gluten
high fat diet	yoghurt
high glucose exposure	zinc deficiency
high n – 6 fatty acid-trans fat	
<b>IATROGENIC</b>	
[(2-fluoro-2',5'-dimethyl-4'-[6-(3-methyl-2-butenyloxy) pyridin-3-yl] biphenyl-4-yl)-(3-methyl-2-butenyl) amine)	indomethacin
(S)-1-(3-hydroxy-2-phosphonmethoxy propyl) cytosine	Infliximab
145-2C11	influenza vaccination
17alpha-ethinyl estradiol	influenza vaccine
17beta-estradiol	Inteferon alpha/beta
18beta-Glycyrrhetic acid	Interferon alfacon-1
1-chloro-2,4-dinitrobenzene	Interferon alfa-n3
2,2'-dichlorodiethyl sulfide	interferon alpha
2,4-Dichlorophenoxybutyric acid	interferon alpha therapy
2-[bis(2-chloroethyl)amino]tetrahydro-2H-1,3,2-oxazophosphorine 2-oxide	Interferon beta-1a
2'3'-Dideoxyinosine	interferon type 1
2-amino-2-(2-[4-octylphenyl]ethyl)-1,3-propanediol hydrochloride	Interferon-alpha
2-aryl carbapenem	interleukin 2 receptor antibody
2'-deoxycoformycin	intravenous gammaglobulin
3,3'-Diindolylmethane	Intravenous immunoglobulin
3'-Azido-2',3'-dideoxythymidine	Intravenous immunoglobulin G
3-methyl-2-quinoxalin benzenevinylketo-1, 4-dioxide	ipilimumab
4AZA1378	iproniazid
5 alpha-dihydrotestosterone	iproniazide
5-azacytidine	irinotecan
5'-deoxy-5-fluorouridine	isoflurane
5'-DFUR	Isoniazid

5'-dFUrd	isoproterenol
5-fluorouracil	ivermectin
6-fluoro-8-nitro-4-oxo-1,4-dihydroquinoline-3-carboxylate	Ixabepilone
9-(N-methyl-L-isoleucine)-cyclosporin A	ixazomib
abacavir	JAK-inhibitors
abatacept	ketorolac
abciximab	Keyhole limpet haemocyanin
Abeta immunization	L. acidophilus
Abetimus	lamivudine
acetaminophen	Lamotrigine
Acetylsalicylate	L-arginine
acetylsalicylic-acid	L-canaline
Acteoside	L-canavanine
Actinomycin D	Leflunomide
acyclic retinoid	left ventricular assist device
acyclovir	Lenalidomide
Adalimumab	leuprolide acetate
adenosine 5'-monophosphate (AMP) Quantum dots (QDs) (AMP-QD)	levamisole
Adjuvanted vaccines	Linezolid
adjuvants	lipidated TN-ApoA-I
Adriamycin	liposomal drugs
ADTP vaccine	liquamycin 200
Advagraf	Lomustine
Afelimomab	lopinavir
Aldesleukin	Lorabid
Aldosterone	loracarbef
Alefacept	lorazepam
alemtuzumab	L-tryptophan
alendronate	lymphocyte immune globulin
alfathesin	M344
alkylating chemotherapy	mAbs
allogeneic bone marrow transplantation	major surgery
allogeneic hematopoietic stem cell transplant	mammary silicone implants
allopurinol	m-carboxy-cinnamicacidbis-hydroxamide
Alloxan	measles, mumps, and rubella vaccine
Alovudine	Mechlorethamine
alpha-ethinylestradiol	medical devices in contact with the blood stream
alpha-methyldopa	medroxyprogesterone
alpha-momorcharin	medroxyprogesterone acetate
Altretamine	mefenamic acid
Ambisome	meglumine antimoniate
amiodarone	Melphalan
Amlodipine	Mepolizumab
Amoxicillin	Mercaptopurine

amphenicols	Meropenem
Amphotericin	methadone
Amprenavir	methimazole
Amsacrine	Methotrexate
anabolic androgenic steroid	Methylprednisolone
Anakinra	Midazolam
anesthesia	minocycline
antibiotics	mirtazapine
anticalcineurin	Misoprostol
Anti-CD20	mitoxantron
anti-CD3 monoclonal antibody	Mitoxantrone
anticonvulsants	mizoribine
anti-CTLA-4 antibody	Mometasone
Anti-D immunoglobulin	Mometasone furoate
anti-d(beta)h-saporin	monoclonal anti-16/6 Id antibody
antilymphocyte globulin	monoclonal antibodies
anti-programmed death 1	Monoclonal antilymphocyte antibodies
antiretroviral agents	Monomethyl fumarate
antithymocyte globulin	Morphine
anti-thymocyte globulin treatment	morphine sulfate
Antituberculosis drug	moxifloxacin
antiviral treatment	murine-derived monoclonal CD3 antibody
antiviral vaccines	muromonab
Apicidin	Mycophenolate Mofetil
Apremilast	mycophenolate sodium
Arctigenin	Mycophenolic acid
Aripiprazole	Mycophenolic acid
artemisinin	N4-Trimethoxybenzoyl-5'-deoxy-5-fluorocytidine
aspirin	N-acylhydrazone
Atgam	nanomedicinal products
ATG-Fresenius S	nanoparticle-formulated drugs
atropine	naproxen
AZA	Natalizumab
Azacitidine	N-desmethyloclozapine
Azathioprine	Nelarabine
azidothymidine	neomycin
azithromycin	nevirapine
bacille Calmette-Guerin vaccine	N-hydroxy-7-(-4-dimethylaminobenzol)aminoheptanamide
Bariatric Surgery	nifedipine
basiliximab	Nilotinib
Begelomab	nitrofurantoin
Belatacept	nitrogen mustard
Belimumab	Nivolumab
Belinostat	nomifensine
Bendamustine	non-steroidal anti-inflammatory drugs

benzocaine	norfloxacin
Berberine	Norfloxacin nicotinate
Beta-diketone antibiotics	novobiocin
beta-lactam antibiotics	NSAIDs
betalactams	nucleic acid-based therapeutics (NATs) by a nanocarrier
Betamethasone	Obinutuzumab
Bevacizumab	Ocrelizumab
Bexarotene	ofatumumab
bezafibrate	Ofloxacin eyedrops
Biopin	Olaparib
Bisphosphonates	omega interferon
Bleomycin	open-heart surgery
Blinatumomab	operation
Blood transfusion	oral antibiotic treatment
bone marrow transplantation	Oryza sativa
Bortezomib	oseltamivir
Bosutinib	Oxaliplatin
BR96 sFv-PE40	Oxcarbazepine
breast implant	oxytetracycline
Brentuximab vedotin	Ozanimod
Briakinumab	Paclitaxel
Brodalumab	Palbociclib
Bucillamine	Panitumumab
Budesonide	Panobinostat
buprenorphine	Paramethasone
Busulfan	Paroxetine
butalbital	Pazopanib
Cabazitaxel	Peficitinib
calcineurin inhibitors	Pefloxacin
Calcium folinate	Pegaspargase
calcium hydroxylapatite	Peginterferon alfa-2a
Canakinumab	Peginterferon alfa-2b
Capecitabine	Peginterferon beta-1a
Carbamazepine	pembrolizumab
Carbendazim	Pemetrexed
Carboplatin	Penicillamine
carboxyphosphamide	penicillin
cardiac surgery	penicillin G
cardiopulmonary bypass	pentamidine
carfilzomib	Pentamidine isethionate
Carmofur	Pentosan
Carmustine	Pentostatin
Castanospermine	pentylene tetrazole
catecholamine	peplomycin
cecal ligation and puncture	peritoneal dialysis



cefotetan	permethrin
Ceftazidime	Phenprocoumon
ceftriaxone	Phenylalanine
Cefuroxime	phenytoin
Celecoxib	Phospholipid-based phase separation gel
cephalosporin	photodynamic therapy
Certolizumab pegol	Photoimmunotherapy
Cervus korean TEMMINCK var. mantchuricus Swinhoe	pimecrolimus
chemoradiation therapy	Piperacillin
chemotherapeutics	piperine
chemotherapy	Piracetam
Chinese medicine injection	Pirarubicin
chlorambucil	Pirfenidone
chloramphenicol	Plasma-derived Human serum albumin p
chlordecone	pneumococcal 13-valent/23-valent polysaccharide vaccine
Chlorhexidine	Pneumococcal vaccine
chloroquine	poliomyelitis vaccine
chlormpromazine	polivalent immunoglobulins
cholinotropic preparations	polyclonal antilymphocyte antibodies
cidofovir	polyclonal antithymocyte antibody
Cimetidine	Poly(lactic-co-glycolic acid) nanoparticle
ciprofloxacin	polymerized porcine hemoglobin
cisplatin	pomalidomide
Cis-platinum	Ponatinib
citalopram	Poppers
Cladribine	post-bone marrow transplant
clavulanic acid	Pralatrexate
Clemastine	Prednicarbate
Clofarabine	prednisolone
clometacin	prednisone
Clonazepam	Prednylidene
clozapine	Preoperative Chemoradiotherapy
colchicine	previcur
combination antiretroviral therapy	procainamide
Complete Freund's adjuvant	Procarbazine
congenital heart disease surgery	progesterone
cord blood transplantation	Prograf
Coriandrum sativum	propranolol
corticoid	Propylthiouracil
corticosteroids	propyphenazone
corticosterone	Protamine
Corticotropin	psoralen
cortisol implants	PUVA treatment
Cortisone acetate	PYCNOGENOL

cotrimoxazole	Pyrazinamide
cranial radiotherapy	pyrimethamine
Cremophor-EL	Qingkailing
CsA	quinine
CT20126	rAAV gene medicine
cupravit	rabbit antithymocyte globulin
cuprizone	radio-immunotherapy
Cyclophosphamide	radioiodine therapy
Cyclophosphamide	radiotherapy
cyclosporin	Raltitrexed
cyclosporine	rapamycin
cyclosporine G	Ravulizumab
Cytarabine	recombinant 19-kDa antigen
cytokine gene therapy	Recombinant human granulocyte-macrophage colony-stimulating factor
cytostatic drugs	renal allografts
cytotoxic drug therapy	rhGM-CSF
cytoxan	Rhododenol
Dacarbazine	rHuIFN-alpha
dacizumab	ribavirin
Dactinomycin	Rifabutin
daptomycin	Rilonacept
Dasatinib	rimantadine
Daunorubicin	Rimexolone
DaunoXome	Risankizumab
death-ligand 1 antibodies	Risperidone
Decitabine	ritonavir
Deflazacort	Rituximab
Delta(9)-tetrahydrocannabinol	RN6G
Deoxyspergualin	Ro 09-1390
derivative of gamma-aminobutyric acid	romidepsin
dexamethasone	Rosuvastatin
Dexrazoxane	Rozanolixizumab
DHPG	rubella vaccine
diazepam	Ruxolitinib
diclofenac	S-2678
didanosine	Sanguinarine
diethylstilbestrol	Sarilumab
Digoxin	Secukinumab
dihydralazine	Seocalcitol
dilantin	serotonin reuptake inhibitors
Dimethyl fumarate	sevoflurane
Dinutuximab	silicon breast implant
diphenylhydantoin	Siltuximab
diphtheria-tetanus-poliomyelitis vaccine	Siponimod
Docetaxel	Sirolimus

dolutegravir	Sirukumab
Dow2	SK&F 95654
Doxifluridine	Sorafenib
Doxil	splenectomy
Doxorubicin	statins
d-penicillamine	stavudine
drosiprenone	stem cell transplantation
drug G1	Stepronin
Eculizumab	steroids
Efalizumab	Streptozocin
efavirenz	SU5416
enemas	suberoylanilide hydroxamic acid
enfuvirtide	sulfamethazole
Epirubicin	sulfamethoxazole
equine antithymocyte globulin	sulfapyridine
Eribulin	Sulfasalazine
erythromycin	sulfisoxazole
Estradiol	sulfur mustard
Estramustine	sulindac
etanercept	sulphadiazine
ETC-216	sulphamethoxazole
ethinylestradiol	Sulphonylurea
Ethyl 4-isothiocyanatobutanoate	Sunitinib
etoposide	Supplemental oxygen
Everolimus	Surgery
fenofibrate	Surgical damage
fenofibric acid	surgical stress
Feraheme	surgical trauma
fibrates	Synthetic human chorionic gonadotropin-related oligopeptides
Fingolimod	tabalumab
FK 506	Tacrolimus
FK-50	Tamoxifen
flavivirus vaccination	tazobactam
Florfenicol	Tedizolid phosphate
Floxuridine	tegafur
Fluazuron	teicoplanin
flucloxacillin	Temozolomide
fluconazole	Temsirolimus
Flucytosine	Teniposide
fludarabine	tenofovir
Fludrocortisone	Tepoxalin
fluocinolone acetonide	teriflunomide
Fluoroquinolones	Tetracyclines
Fluorouracil	Tetrandrine
Fluoxetine	TGN1412

foscarnet	thalidomide
FTY720	thiamphenicol
Fusidic acid	thiazolidinedione
Gadobutrol	thiopental
Gallium nitrate	Thiopronine
ganciclovir	Thiopurine agents
gastrectomy	Thiotepa
gemcitabine	thoracic transplantation
gemfibrozil	thymectomy
Gemtuzumab ozogamicin	thymoglobulin
gentamycin	ticlopidine
Glatiramer	tienilic acid
Glimepiride	Tioguanine
Glivec	tirofiban
glucocorticoids	Tixocortol
GM-CSF vaccine	tobramycin
Golimumab	Tocilizumab
gosereline acetate	Tofacitinib
Granulocyte colony stimulating factor	Topotecan
Gusperimus	Tositumomab
HAART	total-body gamma radiation
haloperidol	total-body irradiation
halothane	Trabectedin
hCG-related oligopeptides	Trastuzumab emtansine
HDAC inhibitors	tremelimumab
hematopoietic stem cell transplantation	trenbolone acetate
hemodialysis	Treponema spp.
Heparin	Tretinoin
Hepatectomy	Triamcinolone
Hepatitis B immunoglobulin	triamterene
hepatitis B vaccine	triazene compounds
hgd40	trichosanthin
highly active antiretroviral therapy	Trichostatin A
Histone deacetylase inhibitors	Trifluridine
Human immunodeficiency virus protease inhibitors	Trilostane
human papilloma virus vaccine	trimethoprim
Hydantoin	Triptolide
hydralazine	Trofosfamide
hydrochlorothiazide	troglitazone
hydrocortisone	Trovafloxacin
Hydroxychloroquine	tumor necrosis factor alpha antagonists
hydroxyethyl starch	Tyrosine kinase inhibitors
hydroxylamine	Ulobetasol
hydroxyurea	Umbilical cord blood transplantation
Hypericin	Umirolimus

Ibritumomab tiuxetan	ustekinumab
Ibrutinib	vaccinations
ibuprofen	vaccines
icotinib	valproic acid
ID35	vancomycin
Idarubicin	Vedolizumab
Idelalisib	Venlafaxine
IFN-alpha	Venocuran
IFN-beta	verapamil
Ifosfamide	Verapamil hydrochloride
IgG immune complex	vesnarinone
IK blockers	vigabatrin
IL-15 superagonist	Vilanterol
Imatinib	Vinblastine
Imiquimod	Vincristine
immune checkpoint inhibitors	Vindesine
immunoglobulin G immune-complex	Vinorelbine
immunoglobulins	Voclosporin
immunomodulatory monoclonal antibodies	Vorinostat
immunostimulatory oligodeoxynucleotide	Warfarin
immunosuppressants	Wortmannin
Immunosuppressive acid protein	YF vaccine
immunosuppressive agents	Zalcitabine
immunosuppressive drugs	zanamivir
immunosuppressive induction therapy	zidovudine
Indinavir	
<b>BIOTOXINS/BIOMATERIALS</b>	
09 LPS	Human immunodeficiency virus
192 IgG-saporin	Human leucocyte antigen G
ATCC 25285	Human Pappilomavirus
A.actinomyetemcomitans	Human respiratory syncytial virus
Aahl	Human Rhinovirus
AahlI	Indoxacarb
Abscisic acid	Influenza A virus protein PB1-F2
aerosolized porcine brain	influenza virus
aflatoxin	Interleukin-10
Aflatoxin B	kainic acid
Aflatoxin B1	kaliotoxin
aflatoxin M1	Kaposi's sarcoma-associated herpesvirus
aflatoxins	ketones
Airborne fungus	lacto-N-fucopentaose3
alcohols	Lichen planus
Amylosin	Lipopolysaccharides
Anatoxin-a	lymphocytic choriomeningitis virus
Androctonus australis hector scorpion venom	M. tuberculosis
Anthrax lethal toxin	malondialdehyde

Anti-dopamine beta-hydroxylase immunotoxin	Marek's disease virus
antigenic peptide P815E	measles virus infection
Antx-a	microcystin-LR
Asobara japonica venom	microcystins
Aspergillus fumigatus spores	milky sap of F. carica
Aspergillus versicolor	Mitomycin
Ataxin-1 oligomers	mitomycin C
atranone A	Mixed mold mycotoxicosis
atranone C	moisture-damaged buildings
B. forsythus	moniliformin
Bacillus Calmette-Guerin	Monocrotaline
Bacillus thuringiensis israelensis	Moraxella catarrhalis
bacterial	mould
bacterial infections during pregnancy	Murine gammaherpes-virus-68
Bacteroides fragilis	Mus m 1
beauvericin	Mycobacteria tuberculosis adjuvant
Berthellina citrina	Mycobacterium marinum
beta-(1,3)-glucan	Mycobacterium tuberculosis
bilirubin	Mycobacterium xenopi
Bordetella pertussis	Mycolactone
Borrelia burgdorferi	Mycotoxicoeses
Bovine viral diarrhea virus	mycotoxins
brevetoxin	Neonatal infections
brown recluse spider venom	nivalenol
Brucella melitensis	Nodularin
bursal disease virus	Nomuraea rileyi
butylated hydroxytoluene	NTHI
C. albicans	ochratoxin A
C. perfringens	ochratoxin B
C. rectus	okadaic acid
Candida albicans water soluble fraction	oleic acid
Capnocytophaga	opportunistic infections
cardiac glycoside	organic dusts
carrageenan	ovalbumin
catalase	P. gingivalis
chitosan	P. intermedia
Chlamydia pneumoniae	Pam3Cys
cholera toxin	Pasteurella haemolytica A1 lipopolysaccharide
Chronic Viral Infection	pathogen-associated molecular patterns
Chrysaora quinquecirrha venom	Patulin
Cis-urocanic acid	peptidoglycan
citrinin	peptidoglycan-polysaccharide
clade B human immunodeficiency virus	pertussis toxin
Clostridium perfringens	phorbol ester
CMV	phytoestrogen
cobra venom factor	Plasmodium

Coinfection with an intestinal helminth	pollen
collagen	Polymethoxylated flavone extract
concanavalin A	Polyomavirus BK
Contact allergens	porcine serum
Cotesia chilonis venom	Porphyromonas gingivalis
coxsackievirus B3	Premolis semirufa
cryopreserved mesenchymal stromal cells	Prevotella intermedia
cyanobacteria	Prolactin-induced protein
cyanobacterial metabolites	Proteoglycan
cyanotoxin	protozoan parasites
Cyclopiazonic acid	Pseudomonas aeruginosa
Cylindrospermopsin	Pseudomonas exotoxin A
Cytomegalovirus	Pteridium aquilinum
daphnane diterpene esters	Pteromalus puparum venom
Demodex mites	ragweed
Dengue Infection	Recombinant interleukin-2
deoxynivalenol	respiratory pathogens
Dermatophagoides pteronyssinus group 5	Rhein
D-galactosamine	Rhinovirus
D-galactose	ribosome inhibiting protein
dinoflagellate genus Alexandrium	ricin toxin A
domoic acid	Rotavirus
dust	Royal jelly
dust mite	rubella
E. coli endotoxin	rubratoxin B
E. nodatum	S. intermedius
EBV	S. mutans
Echinacea	S.aureus
echinococcal	sporin
endotoxin	Schistosoma mansoni
enniatin B	Scleroderma guani venom
Epstein-Barr virus	sepsis
Escherichia coli O:14 lipopolysaccharide	settled dust
Escherichia coli lipopolysaccharide	Shiga toxin
estrogen	Simian immunodeficiency virus
estrogenic xenobiotics	SIVcpz
excess Growth hormone	soybean dust
extractable organic matter	spores of basidiomycete fungi
Fasciola hepatica	squalene
flour dust	Stachybotrys chartarum
Francisella tularensis	staphylococcal enterotoxin B
	Staphylococcus aureus
Friend leukemia virus	Staphylococcus epidermidis
FtoxG50	Streptococcus pneumoniae
fumonisin B1	Streptococcus pyogenes
fungi bioaerosols	Streptomyces californicus

fusarenon-X	sulfur-fumigated polysaccharides from Smilax glabra Roxb
Fusaric Acid	Superantigens
fusariotoxins	Swainsonine
fusarium T-2 toxin	T-2 toxin
Fusobacterium nucleatum	Taenia solium metacestode
galactosamine	Tat 1-72
Gibberellic acid	tetanus toxin
Ginkgo leaf extracts	Thalassophryne nattereri fish venom
gliadin	thermophilic actinomycete
gliotoxin	Tityustoxin
glucose-6-phosphate isomerase	TLR ligands lipopolysaccharide
Glutamate	toxic cyanobacteria
grain dust	Toxoplasma gondii
gram-negative bacteria	tree pollen
grass pollen	Trichothecenes
Group-A beta-Haemolytic Streptococcus	Tripterygium wilfordii Hook f
H1N1	Tulipalin A
HBV	type II collagen
HCV	Umbelliferae
heat shock protein 65	Ureaplasma parvum
Hepatitis B virus	vaccinia virus
Hepatitis C Virus	viral infections during pregnancy
Histamine	virus-associated hematologic malignancies
HIV	vomitoxin
HIV-1	wheat alpha-amylase inhibitor
house dust mite	xanthine oxidase
HPV-16 E7	yessotoxins
HT-2	zanthoxylamide protoalkaloids
human cytomegalovirus	zearalenone
Human herpesvirus 8	zymosan A
<b>OCCUPATIONAL/ENVIRONMENTAL</b>	
(90)Sr	hypoxanthine
(Z)-cis-bifenthrin	hypoxia
1,1,1-trichloro-2,2-bis(4-methoxyphenyl)ethane	Imidacloprid
1,1,1-trichloro-2,2-bis(p-chlorophenyl)ethane	imiprothrin
1,1,1-trichloroethane	indeno[1,2,3,c,d]pyrene
1,1-dichloro-2,2-bis(p-chlorophenyl)ethane	indium
1,1-dichloro-2,2-bis(p-chlorophenyl)ethene	Indole-3-butyric acid
1,1-dichloroethylene	Indoxyl sulfate
1,2,3,4,6,7,8-heptachlorodibenzo-p-dioxin	industrial aerosols
1,2,3,4,7-pentachlorodibenzofuran	industrial noise
1,2,3,7,8-pentabromodibenzo-p-dioxin	Industrial organic solvents
1,2,3,7,8-pentachlorodibenzo-p-dioxin	Inhaled particles
1,2,3,7,9-PeCDF	intense sound
1,2,4-benzenetriol	iodine excess



1,2,5,6-dibenzanthracene	iodine gadital
1,2-dibromopropane	ionizing radiation
1,2-dichloroethane	iron
1,2-dimethylhydrazine	iron nanoparticles
1,3,6,8-TCDF	iron oxide nanoparticles
1,3-butadiene	iron-containing compounds
1,3-dibromopropane	irradiation
1,4-bis [( 2-aminoethyl)amino]-5,8-dihydroxy-9,10-anthracenedione dihydrochloride	Isocyanates
1.8 GHz electromagnetic fields	Isophorone diisocyanate
12-methylbenz[a]anthracene	jet fuel
17alpha-ethynylestradiol	Jet-A
17alpha-Methyltestosterone	JP-8
1-Bromobutane	lactic acid
1-bromopropane	lambda-cyhalothrin
1-Chloro-4-(Trifluoromethyl)Benzene	lanthanoids
1-methyl-3-octylimidazolium bromide	lead
1-methyl-4-phenyl pyridinium	lead acetate
1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine	lead nitrate
2,2',4,4'-tetra-bromodiphenyl ether	leptophos
2,2',4,6,6'-Pentachlorobiphenyl	lewisite
2,3',4,4',5-pentachlorobiphenyl	lindane
2,3,4,7,8-PeCDF	Lipid-based nanoparticles
2,3,4,7,8-pentachlorodibenzofuran	lithium
2,3,5,6-tetrachloro-4-(methylsulphonyl)pyridine	Lithium carbonate
2,3,7,8-tetrabromodibenzo-p-dioxin	lithium chloride
2,3,7,8-tetrachlorinated dibenzo-p-dioxin	long-haul flight
2,3,7,8-tetrachlorodibenzodioxin	low-frequency high-intensity magnetic field
2,3,7,8-tetrachlorodibenzofuran	low-frequency noise
2,3,7,8-tetrachlorodibenzo-p-dioxin	Magnesium sulfate
2,3-dichloro-1-propanol	Malathion
2',3'-dideoxyinosine	mancozeb
2,3-dihydro-2,2-dimethyl-7-benzo-furanol N-methylcarbamate	manganese
2,4,6-tribromophenol	matacil
2,4,6-trichlorophenol	Mechanical ventilation
2,4,6-triiodophenol	Mecoprop
2,4,6-trinitrobenzene sulfonic acid	Melamine
2,4,6-trinitrophenyl-N-methyl-nitramine	Mercuric chloride
2,4,6-trinitrotoluene	mercury
2,4,6-triphenyl-4H-selenopyrane	Meso-2,3-dimercaptosuccinic acid
2,4-Diaminotoluene	Mesoporous silica nanoparticles
2,4-dichlorophenol	metal nanoparticles
2,4-dichlorophenoxyacetic acid	metallic mercury vapor
2,4-dinitrofluorobenzene	metal-oxide nanoparticles
2,5-hexanedione	metals

2,6,10,14-tetramethylpentadecane	metalworking fluid
2,7-dichlorodibenzo-p-dioxin	meta-nitrotoluene
2100-MHz radiofrequency radiation	methacrylate
2450-MHz microwave	methamidophos
2-acetyl-4(5)-(1,2,3,4-tetrahydroxybutyl)-imidazole	Methanol
2-Acetylaminofluorene	methoxychlor
2-amino-1-methyl-6-phenylimidazo[4,5-b]pyridine	methyl bromide
2-Aminoanthracene	methyl chloride
2-bromopropane	methyl ethyl ketone
2-Mercaptobenzothiazole	Methyl eugenol
2-methoxyacetic acid	methyl isocyanate
2-methoxyethanol	methyl methanesulfonate
2-methoxyethyl acetate	methyl tertiary-butyl ether
2-octynoic acid	methylmercury
3-(2-chloroethyl)-2(2-mesyloxyethylamino)-tetrahydro-2H-1,3,2-oxazaphosphorine-2-oxide	methylmercury chloride
3-(3,4-dichlorophenyl)-1,1-dimethylurea	methylmercury sulfide
3,3',4,4',5,5'-hexabromobiphenyl	methylnitrosourea
3,3',4,4',5,5'-hexaCB	methyl-N-nitrosourea
3,3',4,4',5-pentaCB	methylparathion
3,3',4,4',5-pentachlorobiphenyl	methyltetrahydrophthalic anhydride
3,3',4,4'-tetraCB	metribuzin
3,3',4,4'-tetrachloroazobenzene	Microcrystalline cellulose
3,3',4,4'-tetrachloroazoxybenzene	microgravity
3,4,3',4'-tetrachlorobiphenyl	Microparticles
3,4,5,3',4',5'-hexabromobiphenyl	microplastics
3,4,5,3',4',5'-hexachlorobiphenyl	microwaves
3,4-dichloroaniline	millimeter-wave
3,4-dichloropropionanilide	mineral dust
3,4-dichloropropionaniline	mineral oil distillate D-11
3-methyl-4-nitrophenol	mineral oils
3-methylcholanthrene	mining
3-Monochloro-1,2-propanediol	mirex
3-phenoxybenzoic acid	m-nitrotoluene
4-(methylnitrosamino)-1-(3-pyridyl)-1-butanone	moisture-damage
4,4'-thiobis-(6-t-butyl-m-cresol)	mono-2-ethylhexyl phthalate
460-MHz microwaves	mono-iso-decyl phthalate
4-dimethylamino-3-methyl-N-carbamate	mono-iso-nonyl phthalate
4-hydroxy-2-nonenal	Monomethoxy(polyethyleneglycol)-poly (D,L-lactic-co-glycolic acid)-monomethoxy (PELGE) nanoparticles
4-hydroxydiclofenac	monomethylarsonous acid
4-Methylanisole	mono-n-octyl phthalate
4-nitroquinoline-1-oxide	monophthalates

4-nonylphenol	monosodium urate monohydrate crystal
4-pentylphenol	multiwalled carbon nanotubes
5-(3-3-dimethyl-1-triazeno)imidazole-4-carboxamide	Municipal wastewaters
50 Hz amplitude-modulated 2.45 GHz microwaves	mustard oil
6:2 fluorotelomer sulfonamide alkylbetaine	N,N,-diethyl-m-toluamide
6:2 FTAB	N,N1-bis(dichloroacetyl)-1,8-octamethylene diamine
60Cogamma radiation	N,N-diethylaniline
6-hydroxydopamine	N,N-diethyl-m-toluamide
6-mercaptopurine	N-acetylneuraminic acid
6-propyl-2-thiouracil	Nanoceria nanoparticles
7,12-dimethylbenz(a)anthracene	nanomaterials
8:2 Fluorotelomer alcohol	Nanoparticle protein corona (NP-PC)
835 MHz radiofrequency	nanoparticles
9-(2-hydroxyethoxymethyl)guanine	nanoparticulate TiO2
9,10-dimethylantracene	nanoplastic particles
900 MHz radiofrequency radiation	Nanoshells
915-MHz microwave radiation	nanosilicas
abamectin	nanosilver
abandoned oil refinery	naphthalenes
abestos	naphthenic acids
acephate	N-bis (2-hydroxypropyl) nitrosamine
Acer3 deficiency	N-Butylbenzene sulfonamide
Acetaldehyde	N-butyl-N-(4-hydroxybutyl) nitrosamine
acetamiprid	NdCl3
Acetochlor	Neb-colloostatin
acetofenate	negatively charged-particle dominant indoor air-conditions
acetone	neonicotinoids
acetoxydimethylnitrosamine	N-ethyl, N-nitrosourea
acetyldinaline	neutron irradiation
acid water	NH4VO3
acrolein	n-hexane
acrylamides	nickel
acrylonitrile	nickel aerosols
activated carbon	Nickel nanoparticles
activated-sludge	nickel oxide
Adamsite	nickel subsulfide
Agent Orange	nickel sulfate hexahydrate
agrochemicals	NiCl2
air pollutants	Nicotinamide
air pollution	nicotine
airborne particles	niridazole
airborne particulate matter	nitenpyram

airborne suspended matter	nitrite
Al citrate	Nitrobenzene
alachlor	Nitrofurazone
aldicarb	nitrogen dioxide
alkylphenol polyethoxylates	nitrogen oxides
alkylphenols	nitrosamines
alpha3(IV) collagen	nitrosomethylurea
alpha-cypermethrin	nitrous oxide
alpha-naphthyl-AS-acetate esterase	N-methyl-D-aspartate
alpha-naphthyl-butyrate esterase	N-methyl-N'-nitro-N-nitrosoguanidine
altered oxygen supply	N-methyl-N-nitrosourea
Alumina nanoparticles	N-nitrosodibutylamine
aluminium	N-nitrosodiethylamine
Aluminium chloride	N-nitrosodimethylamine
Aluminum	N-nitrosodipropylamine
aluminum chloride	N-nitrosomethylbutylamine
aluminum hydroxide	N-nitrosomethylethylamine
aluminum sulfate	N-nitrosomethylpropylamine
aluminum trichloride	N-nitroso-N-methylurea
aluminum-based nanoparticles	noise
ambient particulate matter	non-halogenated p-benzo-1,4-quinone
Aminocarb	nonoxynol-9
amitraz	nonylphenol
ammonia	O,O,S-trimethyl phosphorothioate
Ammonium hexachloroplatinate	O,O-diethyl-O-4-nitro-phenylthiophosphate
ammonium metavanadate	O,O-dimethyl, S-ethyl phosphorothioate
amyl nitrite	o,p-dichlorodiphenyldichloroethylene
anhydride	O,S-dimethyl-acetylphosphoramidothioate
anthracene	O3-degraded AA
antigravitary posture	ocean acidification
antimony	oil exposure
aqueous fraction of PW	oil fly ash
Arabian crude oil	Oil mist
arc welding fume	oil sands process water
arecoline	Oil Sands tailings water
Aroclor 1254	oil sands-derived naphthenic acids
Aroclor 1260	oil spills
Aroclor-1242	O-isopropyl methylphosphonofluoridate
Aroclor-1248	Oleic acid anilide
Aroclors	omethoate
aromatic amino compounds	o-Nitroanisole
aromatic halogenated disinfection byproducts	Orange II
aromatic hydrocarbon receptor agonists	ordnance
aromatic hydrocarbons	Organic acid anhydrides
aromatic nitro compounds	organic hydrocarbons
arsenate	organic pollutants

arsenic	organic solvents
arsenic trioxide	Organochlorine compounds
Arsenite	organochlorine contaminants
arsenobetaine	organochlorine insecticide
aryl-triazene derivatives	Organochlorine pesticides
asbestos	organohalogen flame retardant
atmospheric pollution	organophosphate flame retardants
atrazine	organophosphorous pesticides
aurothiopropanol-sulfonate	Organophosphorus Compounds
aurothiopropanol-sulfonate sodium salt	organosilicon compounds
Avermectin	organotin compounds
BDE-99	Ortho-phenylphenol salt
Be aluminum	ortho-phthalaldehyde
Be oxide	OS leachate
bendiocarbamate	organic perfluorochemicals
Bentazon TP	ouabain
bentazone	Oxazolone
benz(a)anthracene	oxidant gases
benz(a)pyrene	ozone
benzalkonium chloride	p,p'-dichlorodiphenyldichloroethylene
Benzanthrone	paints
benzene	palladium
benzenes	palmitoleic acid
Benzethonium chloride	palmyrah
benzo [a]pyrene	PAMAM dendrimers
benzo[a]pyrene-7,8-dihydrodiol	pancreatic enzymes
benzo[b]fluoranthene	Paper and pulp mill effluent
benzo[k]fluoranthene	parabens
benzo-1,4-quinones	para-Nitrotoluene
benzodiazepines	paraquat
Benzothiazole	parathion
benzyl 3-quinuclidylate	particulate matter
benzyl alcohol	passive smoking
beryllium	Pb-acetate
Beta-chlorolactic acid	PBDE-209 exposure during pregnancy
beta-cypermethrin	p-bromophenylacetylurea
beta-N-Methylamino-L-alanine	PCB118
Bi2Se3 nanoparticles	PCB126
Bifenthrin	PCB153
biological	PCB169
biomass smoke	PCBs
BioProtein nucleic acid reduced variant	p-chloroaniline
biphenyls	p-chloronitrobenzene
bis(2-chloroethyl) sulfide (HD)	P-dichlorobenzene
bis(tri-n-butyltin)oxide	pendimethalin
bismuth selenide	pentabromodiphenyl ether

bisphenol A	pentaCB
Bisphenol F	pentachlorophenol
bisphenol S	perchloroethylene
bivalent ionic mercury	Perfluorinated alkylated substances
bleached kraft pulp mill effluent	Perfluorinated compounds
Boric acid	per-fluoroalkyl
boron nitride nanoflakes	perfluoroalkyl acids
branched polyethylenimine nanoparticles	perfluoroalkyl carboxylates
Brominated diphenyl ethers	perfluoroalkyl substances
Brominated Tetra-BDE	perfluoro-n-decanoic acid
bromodichloromethane	perfluorononanoic acid
burn	perfluorooctane sulfate
butachlor	perfluorooctane sulfonate
butyltins	Perfluorooctane sulfonic acid
C60 nanoparticles	perfluorooctanoate
cadmium	Perfluorooctanoic acid
cadmium acetate	perfluorooctanoic sulfonic acid
cadmium chloride	perinatal chemical exposures
Cadmium nanoparticles	Permanone
cadmium oxide	permethrin exposure
cadmium telluride quantum dots	persistent organic pollutants
Caramel Colour III	persistent organochlorine compounds
carbamate pesticides	pesticides
carbaryl	petrochemical waste sites
Carbendazim	petrochemical wastes
carbofuran	petrochemicals
carbon black nanoparticles	petroleum
carbon disulfide	PFAS
Carbon fullerenes	PFOA
carbon ion radiation	PFOS
carbon monoxide	PHAHs
Carbon nanoparticles	pharmaceuticals
carbon nanotubes	Phenanthrene
carbon tetrachloride	phenol
carbosulfan	phenols
carboxylic acid-coated QDs	phenoxypyrenols
carpet shampoo	phenyl arsenic compounds
cathelicidin	phosgene
caustic soda	phosphamidon
CdS/CdTe quantum dots	phthalates
CdSe/ZnS quantum dots	physical fatigue
CdTe quantum dots	physiological factors
CeCl3	physiological stress
Cellphone electromagnetic radiation	picoxystrobin
Cement kiln dusts	piperonyl butoxide
chemical pollutants	pirimicarb

chemical stress	Pirimiphos-methyl (O-2-diethylamino-6-methylpyrimidin-4-yl O,O-dimethyl phosphorothioate)
chemical warfare agents	Platinum nanoparticles
chemical weapon	Plutonium
chemically dispersed oil	PM2.5
Chiral organophosphorus pesticides	p-Nitroaniline
Chit	p-nitrotoluene
chlordanes	pollutants
chlorfenvinphos	poly (lactic-co-glycolic acid) nanoparticles
chloridazone	poly(carboxybetaine) nanoparticles
Chlorinated aromatic hydrocarbons	Poly(D,L-Lactic Acid) Nanoparticle
chlorinated carbohydrates	poly(ethylene glycol) nanoparticles
chlorinated dibenzodioxins	polyamidoamine dendrimers
chlorinated dibenzofurans	polyaromatic hydrocarbons
chlorinated dibenzo-p-dioxins	polybrominated biphenyls
Chlorinated Flame Retardant	polybrominated diphenyl ethers
chlorinated hydrocarbons	Polybrominated flame retardants
Chlorinated pesticides	Polycarbonate
chlorine	polychlorinated biphenyl 153
chloroacetophenone	polychlorinated biphenyl congeners
chlorobenzenes	polychlorinated biphenyl mixtures
chloroform	Polychlorinated biphenyls
Chloronitrobenzenes	polychlorinated dibenzofurans
Chlorophenols	polychlorinated dibenzo-p-dioxins
Chloroplatinic acid	polychlorinated dioxins
chlorothalonil	polychlorinated diphenyl ethers
chlorpyrifos	polychlorinated naphthalenes
chromate (VI)	Polycyclic aromatic hydrocarbons
Chromium	polycyclic aromatics
Chrysene	polycyclic hydrocarbons
CI-994	polyethylene microplastics
cigarette smoke	polyfluorinated compounds
Cis-bifenthrin	polyfluoroalkyl
cis-nonachlor	polyhalogenated aromatic hydrocarbons
Citral	polyhalogenated biphenyls
citral dimethyl acetal	polyhalogenated hydrocarbons
Clark I	polyinosinic:polycytidylic acid
Clethodim	Polymeric nanoparticles
clothianidin	polynuclear aromatic hydrocarbon
CO2	Polyphosphoester-based nanoparticles
coal fly ash	polyriboinosinic-polyribocytidilic acid
coal tar	polystyrene
Cobalt	polyurethane
Cobalt chloride	polyvinyl chloride
cocaethylene	pond sediments

Cold	potassium bromate
cold stress	potassium salt
cold water stress	Potassium sorbate
Constant illumination	potassium tetrachloroplatinate
Constant light	p-phenylenediamine
contact lenses	prallethrin 1.6%
coplanar polychlorinated biphenyls	prenatal arsenic exposure
Copolymer-I	prenatal trichloroethylene
copper	Pretilachlor
copper alginate fibers	produced water
copper carbonate nanoparticles	propanil
copper chloride	Propoxur
Copper nanoparticles	propoxur
Copper oxide nanoparticles	propyl gallate
copper sulfate	Prowl 400
Corexit 9500	pulp mill effluent
Corexit/oil mixture	pulse modulated Radiofrequency
cosmic radiation overexposure	punicalagin
cotton dust	Pyraclofos
crassin acetate	pyraclostrobin
crotonaldehyde	pyrene
crude oil	pyrethroid insecticides
crystalline silica	pyrethroid metabolites
cumene hydroperoxide	pyrethroids
Cyanuric acid	pyridostigmine bromide
cyclosporin A	pyrocatechol
cypermethrin	pyrrolizidine alkaloids
Dapsone	quartz
d-cyphenothrin	Quinocetone
DDT	Quinolinic Acid
DE-71	Quinones
decabrominated diphenyl ether	radio frequencies
Dechlorane 602	radioactive isotopes
decimeter waves	radioactivity
decimeter-band electromagnetic field	radiocontrast media
dehydroabietic acid	radiofrequency radiation
Deltamethrin	radioiodine-131
dental amalgam	radiotelevision broadcasting stations
DEP benzene extracts	radiotherapy
dermal bioimplants	residual oil fly ash
dextran sodium sulfate	Resin glass ionomers
dextran stabilized iron oxide nanoparticles	Respiratory Syncytial Virus
dextran sulfate	Retinal laser burn
di-(2-ethylhexyl) phthalate	Roundup
diazinon	Salinity change
dibenz(a,c)anthracene	salt dust



dibenz(a,h)anthracene	sand dust
Dibenzo[def,p]chrysene	Sarin
dibenzofuranes	scum of waste crankcase oil
dibenzofurans	Secondhand Smoke
dibenzo-p-dioxins	selenium
Dibenzopyrene	Sephadex
dibromoacetic acid	sertraline
Dibutyl phthalate	silesian dust
dibutyltin	silica
dichlofluanid	silica nanoparticles
dichloroacetic acid	silicates
dichlorodiphenyldichloroethylene	Silicon dioxide nanoparticles
dichlorodiphenyltrichloroethane	Silicone
dichloroethane	silver
dichloromethane	silver acetate
dichlorophene	silver nanoparticles
dichloropropanol	simazine
diclofop-methyl	single-walled carbon nanotubes
Dicyclohexylphthalate	SiO2 nanocomposites
dieldrin	sodium acetate
diesel exhaust	Sodium arsenite
Dietary Nickle Chloride	Sodium bromate
Diethanolamine	sodium carbonate
Diethyldithiophosphate	sodium chromate(VI)
dihydroxy-di-n-propylnitrosamine	Sodium dodecyl sulfate
diisononylphthalate	Sodium fluoride
diisopropyl ether	sodium methyldithiocarbamate
dim light at night	sodium Ortho-phenylphenol salt
dimethoate	sodium tungstate
Dimethyl carbonate	solar-simulated radiation
dimethyl dichlorovinyl phosphate	solvents
dimethyl nitrosamine	soman
Dimethyl sulfoxide	space radiation
dimethylbenz(a)anthracene	spinal cord injury
dimethylformamide	spinetoram
dimethylnitrosamine	stack gas condensates from waste incineration plants
dimethylsulphoxide	Stannous chloride
di-n-butyltin dichloride	Stomp
dinocap	strobilurins
di-n-octyltin dichloride	styrene
dinotefuran	styrene-butadiene rubber production
Dioxin-like polychlorinated biphenyls	sulfluramid
dioxins	Sulfoxaflor
diphenols	sulfur dioxide
diphenyl ethers	sulfur yperite

diphenylmethane diisocyanate	sulfuric acid
disinfection byproducts	Sunset yellow FCF
Disperse Red 1	supercypermethrin forte
dithiocarbamates	suspended particulates
Diuron	sweet crude oil
Domestic wastewaters	tannery effluent
d-phenothrin	tannins
electromagnetic field	tartrazine
electromagnetic radiation	TCMS pyridine
electromagnetic waves	tebuconazole
electronic waste	Terahertz radiation
Endocrine disrupting compounds	tert-butylhydroquinone
endosulfan	Tertiary-butyl acetate
endosulfane	tetrabromobisphenol A
endrin	Tetrabromoethylcyclohexane
engine exhausts	tetrachlorodiarylmethanes
engineered nanomaterials	tetrachloroethylene
environmental chemicals	tetrachloromethane
environmental estrogens	tetraethyl lead
environmental pollutants	tetramethoxypropane
epichlorohydrin	tetraphenyl arsonium chloride
Erionite	tetryl
esfenvalerate	thermal injury
Ethanol	thiabendazole
ethephon	thiamethoxam
ethyl benzene	thimerosal
ethyl carbamate	thioacetamide
ethyl t-butyl ether	thymidine dinucleotides
ethylene dibromide	thymol
ethylene glycol	Ti ascorbate
ethylene oxide	Ti oxalate
e-waste disposing areas	tin
exhaust gases	TiO2 composite
extremely high frequency	Titanium
famoxadone-cymoxanil	titanium dioxide
Fenitrothion	titanium dioxide nanoparticles
fenthion	Titanium implants
Fenvalerate	titanocene
ferrocenes	tobacco smoke
fine particulate matter	toluene
fipronil	toluene 2,4-diisocyanate
firefighting overhaul environment	toluene diisocyanate
fluoranthene	total particulate matter
Fluorides	toxic waste sites
fluoro-edenite fibers	traffic-related particles
fluorotelomer	trans-nonachlor

fluvalinate	trauma
fly ash	treated urban wastewaters
footshock	trehalose dimycolate
Forafac1157	triallate
Formaldehyde	triazine Irgarol 1051
FR167653	Tributyltin
fuel oil	Tributyltin chloride
Fullerene nanoparticles	tributyltin oxide
furans	Tributyltin-sulfhydryl
furfurol	trichloroacetic acid
furfuryl alcohol	trichloroethane
furniture dust	Trichloroethylene
furosemide	Triclosan
gallium	Triethanolamine
gallium arsenide	trifloxystrobin
gamma radiation	trimellitic anhydride
gamma-chlordane	Trimethyltin
gamma-hexachlorocyclohexane	tri-n-butyltin chloride
gasoline	tri-n-butyltin oxide
Gasoline exhaust particles	trinitrobenzene sulphonic acid
Geogenic dust	trinitrotoluene
geogenic particulate matter	tri-n-propyltin chloride
global warming	tri-ortho-cresyl phosphate
Glucan Nanoparticles	tri-ortho-tolyl phosphate
glufosinate-ammonium	Triphenylphosphate
Glutaraldehyde	triphenyltin
glycidamide	triphenyltin acetate
glycol ether 2-methoxyethanol	triphenyltin chloride
glyphosate	Triphenyltin derivatives
Gold	Tris(1,3-dichloroisopropyl) phosphate
Gold nanoparticles	tropolone
gold-salt	tungsten
graphene nanoplatelets	Tungsten substances
graphene oxides	Type A trichothecenes
guaiacol	type B trichothecenes
halocarbons	UHF electrical field
halogenated aromatic hydrocarbons	ultrafine particulate matter
halogenated benzo-1,4-quinone	Ultraviolet B
Hapten	ultraviolet light
hardwood smoke	ultraviolet radiation
HCC-230fa (1,1,1,3,3,3-hexachloropropane)	unsymmetrical dimethylhydrazine
head-down tilt bed rest	untreated urban wastewaters
heat	Urban air pollution
heat stress	urban waste water
heavy crude oil	urethane
heavy mechanical trauma	Ustad-10 EC

Heavy Metals	UVA
heavy oil	UVR
hepatotoxic chemicals	Vanadium
Hepatotoxics	Vanadium pentoxide
heptachlor	vehicle emissions
Hexachlorobenzene	vinclozolin
hexachlorocyclohexanes	vinyl chloride
hexachloroethan	vinylidene chloride
hexahydrophthalic anhydride	volatile organic compounds
High altitude	volcano mud
high frequency electromagnetic radiation	VX
high temperature	washing soda
high water temperature	waterborne selenium
high-dose sunlight	welding aerosols
highly chlorinated biphenyls	welding fumes
Hospital wastewaters	whole-body vibration
hydrazine	whole-smoke conditioned medium
hydrocarbon oil	wide-band modulated electromagnetic field
Hydrogen peroxide	xenobiotics
hydrogen sulfide	xylene
Hydrophobic sodium fluoride-based nanocrystals doped with lanthanide ions	zinc
Hydrophobicity	zinc chloride
hydroquinone	Zinc oxide nanoparticles
Hydroxylated fullerenes	Zinc pyrithione
Hypergravity	ziram
hyperoxia	
<b>PSYCHOSOCIAL/SOCIOECONOMIC</b>	
accumulated life stress	male partner violence
acute psychological stressors	mild stress
adverse life events	negative stressful life
anxiety	Posttraumatic stress
anxious women	psycho-emotional stress
Childhood adversity	psychological stress
Chronic life stress	restraint stress
chronic mild stress	social behaviour deficits
chronic restraint stress	social disruption
Chronic social stress	Social isolation
chronic stress	Social stress
Combined Stress	stress
Depression	Stressful life events
Disaster	stress-related behavior
Distress	traumatic life events
intense intellectual stress	vulnerability
life stress	Work stress

“

"A4-B. Bibliography – Contributing Factors to Weakened Immune System – Titles Only

As stated in the previous section, approximately 4,700 records included factors that contributed to weakening the immune system, and these records formed the basis of the analysis. The titles of these records follow. To retrieve the abstract, or in some cases the full-text of these records, insert the title shown below into Pubmed.

18beta-Glycyrrhetic acid from licorice root impairs dendritic cells maturation and Th1 immune responses

192IgG-saporin-induced immunotoxic lesions of cholinergic basal forebrain system differentially affect glutamatergic and GABAergic markers in cortical rat brain regions

2,2',4,6,6'-Pentachlorobiphenyl-induced apoptosis is limited by cyclooxygenase-2 induction

2,3,7,8-TCDD neurotoxicity in neuroblastoma cells is caused by increased oxidative stress, intracellular calcium levels, and tau phosphorylation

2,3,7,8-Tetrachlorodibenzo-p-dioxin affects the differentiation of CD4 helper T cell

2,3,7,8-Tetrachlorodibenzo-p-dioxin (TCDD) and long term immunologic memory

2,3,7,8-Tetrachlorodibenzo-p-dioxin (TCDD) induced suppression of the local immune response

2,3,7,8-Tetrachlorodibenzo-p-dioxin (TCDD)-induced immunotoxicity

2,3,7,8-Tetrachlorodibenzo-p-dioxin impairs stable establishment of oral tolerance in mice

2,3,7,8-Tetrachlorodibenzo-p-dioxin-mediated impairment of B cell differentiation involves dysregulation of paired box 5 (Pax5) isoform, Pax5a

21 Days head-down bed rest induces weakening of cell-mediated immunity - Some spaceflight findings confirmed in a ground-based analog

2'3'-Dideoxyinosine inhibits the humoral immune response in female B6C3F1 mice by targeting the B lymphocyte

25 years of UV-induced immunosuppression mediated by T cells-from disregarded T suppressor cells to highly respected regulatory T cells

2-Methoxyestradiol protects against IgG immune complex-induced acute lung injury by blocking NF-kappaB and CCAAT/enhancer-binding protein beta activities

3,3'-Diindolylmethane induces immunotoxicity via splenocyte apoptosis in neonatal mice

3,4-Dichloropropionanilide-induced atrophy of the thymus: mechanisms of toxicity and recovery

3-Methylcholanthrene induces lymphocyte and phagocyte apoptosis in common carp (*Cyprinus carpio* L) in vitro

3-Methylindole-induced splenotoxicity: functional analysis of immune parameters and lymphocyte phenotyping by flow cytometry

51Cr release assay of clozapine-induced cytotoxicity: evidence for immunogenic mechanism

5-azacytidine promotes an inhibitory T-cell phenotype and impairs immune mediated antileukemic activity

6:2 fluorotelomer sulfonamide alkylbetaine (6:2 FTAB), a novel perfluorooctane sulfonate alternative, induced developmental toxicity in zebrafish embryos

6-Methylprednisolone does not impair anti-thymocyte globulin (ATG) immunosuppressive activity in non-human primates

7,12-Dimethylbenz[a]anthracene-induced bone marrow toxicity is p53-dependent

8:2 Fluorotelomer alcohol causes immunotoxicity and liver injury in adult male C57BL/6 mice

954 MHz microwaves enhance the mutagenic properties of mitomycin C.

A 33-year-old patient with human immunodeficiency virus on antiretroviral therapy with efavirenz-induced complex partial seizures: a case report

A case of daptomycin-induced immune thrombocytopenia

A case of drug-induced hypersensitivity syndrome induced by icotinib managed by intravenous immunoglobulin and systemic corticosteroids

A Case of Heparin-Induced Thrombocytopenia with Discrepancy in the Results of Anti-Platelet Factor 4/Heparin Antibodies between Latex-Particle-Enhanced Immunoturbidimetric Assay and Enzyme Immunoassay

A case of inflammatory pseudotumor of the lung with suppressed immune response in the patient with renal cell carcinoma after nephrectomy and administration of interferon

A case of isoniazid-induced thrombocytopenia: recovery with immunoglobulin therapy

A case of linear immunoglobulin A bullous dermatosis in a patient exposed to sun and an analgesic

A case of pembrolizumab-induced type-1 diabetes mellitus and discussion of immune checkpoint inhibitor-induced type 1 diabetes

A case of profound and prolonged tirofiban-induced thrombocytopenia and its correction by intravenous immunoglobulin G

A case of thrombocytopenia induced by digitoxin--antibody immune complex

A clinical and immunologic study of workers with trimellitic-anhydride-induced immunologic lung disease after transfer to low exposure jobs

A commercial sunscreen's protection against ultraviolet radiation-induced immunosuppression is more than 50% lower than protection against sunburn in humans

A comparative evaluation of immunotoxicity of DDT and its metabolites in rats

A comparative evaluation of immunotoxicity of malathion after subchronic exposure in experimental animals

A comparative evaluation of the immunotoxicity and immunomodulatory effects on macrophages exposed to aromatic trihalogenated DBPs

A Comparative Study on the Effects of Different Parts of Panax ginseng on the Immune Activity of Cyclophosphamide-Induced Immunosuppressed Mice

A comparison of immunotoxic effects of nanomedicinal products with regulatory immunotoxicity testing requirements

A Comparison of Two Types of Rabbit Antithymocyte Globulin Induction Therapy in Immunological High-Risk Kidney Recipients: A Prospective Randomized Control Study

A comprehensive review of pesticides and the immune dysregulation: mechanisms, evidence and consequences

A cooperative effect of surgical stress-induced immunomodulation with the cessation of immunosuppression on hepatitis C virus elimination

A critical reappraisal of the current data on drug-induced linear immunoglobulin A bullous dermatosis: a real and separate nosological entity

A critical review of perfluorooctanoate and perfluorooctanesulfonate exposure and immunological health conditions in humans

A cross-sectional study of changes in markers of immunological effects and lung health due to exposure to multi-walled carbon nanotubes

A cytokine cascade including prostaglandin E2, IL-4, and IL-10 is responsible for UV-induced systemic immune suppression

A detailed study of developmental immunotoxicity of imidacloprid in Wistar rats

A dose-response study of the effects of prenatal and lactational exposure to TCDD on the immune response to influenza A virus

A double-edged sword: does highly active antiretroviral therapy contribute to syphilis incidence by impairing immunity to *Treponema pallidum*

A flow immunoassay for studies of human exposure and toxicity in biological samples

A health food high-peptide meal alleviates immunosuppression induced by hydrocortisone and cyclophosphamide in mice

A high SPF sunscreen's effects on UVB-induced immunosuppression of DNCB contact hypersensitivity

A higher anxiety state in old rats after social isolation is associated to an impairment of the immune response

A high-throughput screening method based on stably transformed human cells was used to determine the immunotoxic effects of fluoranthene and other PAHs

A meta-analysis and morphological review of cyclosporine-induced nephrotoxicity in auto-immune diseases

A method of assessing impairment of immunity after trauma

A model of immune-mediated lung disease in rats sensitized to house dust mite and upregulation of immunity following exposure to nitrogen dioxide

A model of immunologic lung injury induced by trimellitic anhydride inhalation: antibody response

A monoclonal antibody to hippuric acid: an improved enzyme-linked immunosorbent assay for biological monitoring of toluene exposure

A multivariate assessment of innate immune-related gene expressions due to exposure to low concentration individual and mixtures of four kinds of heavy metals on zebrafish (*Danio rerio*) embryos

A murine model of the effects of inhaled CuO nanoparticles on cells of innate and adaptive immunity - a kinetic study of a continuous three-month exposure

A neurotoxic regimen of methamphetamine exacerbates the febrile and neuroinflammatory response to a subsequent peripheral immune stimulus

A new assay system for evaluation of developmental immunotoxicity of chemical compounds using respiratory syncytial virus infection to offspring mice

A new model for an etiology of rheumatoid arthritis: smoking may trigger HLA-DR (shared epitope)-restricted immune reactions to autoantigens modified by citrullination

A new pathologic pathway for pulmonary fibrosis induced by silica: involvement of immunosuppressive responses

A new side effect of immunosuppression: high incidence of hearing impairment after liver transplantation

A nontoxic *Pseudomonas* exotoxin A induces active immunity and passive protective antibody against *Pseudomonas* exotoxin A intoxication

A novel immunochromatographic electrochemical biosensor for highly sensitive and selective detection of trichloropyridinol, a biomarker of exposure to chlorpyrifos

A novel immunosuppressive drug, FTY720, prevents the cancer progression induced by cyclosporine

A novel mouse model of chronic prostatitis/chronic pelvic pain syndrome induced by immunization of special peptide fragment with aluminum hydroxide adjuvant

A novel T(H)17-type cell is rapidly increased in the liver in response to acetaminophen-induced liver injury: T(H)17 cells and the innate immune response



A phase-I study of repeated therapy with radiolabelled antibody to carcinoembryonic antigen using intermittent or continuous administration of cyclosporin A to suppress the immune response

A Potential Mechanism for Immune Suppression by Beta-Adrenergic Receptor Stimulation following Traumatic Injury

A predictive F344 rat immunotoxicology model: cellular parameters combined with humoral response to NP-CgammaG and KLH

A preliminary evaluation of immune stimulation following exposure to metal particles and ions using the mouse popliteal lymph node assay

A prospective, double-blind, randomized study of high-versus low-dose OKT3 induction immunosuppression in cadaveric renal transplantation

A protective effect of vitamin E and selenium in ameliorating the immunotoxicity of malathion in chicks

A proteomic approach for the identification of immunotoxic properties of Tulipalin A

A randomized controlled trial of late conversion from calcineurin inhibitor (CNI)-based to sirolimus-based immunosuppression in liver transplant recipients with impaired renal function

A randomized, double-blinded comparison of Thymoglobulin versus Atgam for induction immunosuppressive therapy in adult renal transplant recipients

A rare case report of acyclovir-induced immune thrombocytopenia with tongue hematomas as the first sign, and a literature review

A review of the immunotoxicity of the pesticide 3,4-dichloropropionanilide

A role for associated transition metals in the immunotoxicity of inhaled ambient particulate matter

A role for HO-1 in renal function impairment in animals subjected to ischemic and reperfusion injury and treated with immunosuppressive drugs

A role for neuropeptides in UVB-induced systemic immunosuppression

A role for oxidative stress in suppressing serum immunoglobulin levels in lead-exposed Fisher 344 rats

A role for reactive oxygen species in endotoxin-induced elevation of MOR expression in the nervous and immune systems

A semiquinone glucoside derivative isolated from *Bacillus* sp. INM-1 provides protection against 5-fluorouracil-induced immunotoxicity

A single dose of trichloroethylene given during development does not substantially alter markers of neuroinflammation in brains of adult mice

A site involving the "hybrid" and PSI homology domains of GPIIIa (beta 3-integrin subunit) is a common target for antibodies associated with quinine-induced immune thrombocytopenia

A soy-derived immunostimulating peptide inhibits etoposide-induced alopecia in neonatal rats

A statement on the developmental immunotoxicity of bisphenol A (BPA): answer to the question from the Dutch Ministry of Health, Welfare and Sport

A study of autoantibodies and circulating immune complexes in mercury-exposed chloralkali workers

A study on behavioral, neurotoxicological, and immunotoxicological effects of subchronic arsenic treatment in rats

A study on geno- and immunotoxicological effects of subacute propoxur and pirimicarb exposure in rats

A study on immunotoxicological effects of subacute amitraz exposure in rats

A study on the effects of exposure to benzene on the activity of immunoglobulin E

A toxicokinetic study of nickel-induced immunosuppression in rats

A UVB wavelength dependency for local suppression of recall immunity in humans demonstrates a peak at 300 nm

Ability of *Lactobacillus plantarum* MON03 to mitigate aflatoxins (B1 and M1) immunotoxicities in mice

Ability of *Lactobacillus rhamnosus* GAF01 to remove AFM1 in vitro and to counteract AFM1 immunotoxicity in vivo

Abnormal immunostaining for dystrophin in isoproterenol-induced acute myocardial injury in rats: evidence for change in dystrophin in the absence of genetic defect

Abnormalities of cellular immune response in arthritis induced by rubella vaccination

Absorption of PCB126 by upper airways impairs G protein-coupled receptor-mediated immune response

Accelerated induction of skin cancers by ultraviolet radiation in hairless mice treated with immunosuppressive agents

Accumulation and immunotoxicity of microplastics in the estuarine worm *Hediste diversicolor* in environmentally relevant conditions of exposure

Accumulation of polychlorinated biphenyls (PCBs) and evaluation of hematological and immunological effects of PCB exposure on turtles

Accumulation, histopathology and immunotoxicological effects of waterborne cadmium on gilthead seabream (*Sparus aurata*)

Acephate immunotoxicity in White Leghorn cockerel chicks upon experimental exposure

Acetaldehyde accelerates HCV-induced impairment of innate immunity by suppressing methylation reactions in liver cells

Acetaldehyde-serum protein adducts inhibit interleukin-2 secretion in concanavalin A-stimulated murine splenocytes: a potential common pathway for ethanol-induced immunomodulation

Acetaminophen toxicity revisited: is drug-induced hepatotoxicity immune mediated

Acetaminophen-induced immunosuppression associated with hepatotoxicity in mice

Acetylcholinesterase inhibition as a biomarker of adverse effect. A study of *Mytilus edulis* exposed to the priority pollutant chlorfenvinphos

Acetylcholinesterase monoclonal antibody-induced sympathectomy: effects on immune status and acute morphine-induced immunomodulation

Acrylamide induced immunosuppression in rats and its modulation by 6-MFA, an interferon inducer

Acrylamide induces immunotoxicity through reactive oxygen species production and caspase-dependent apoptosis in mice splenocytes via the mitochondria-dependent signaling pathways

Actions of alcohol on immunity and neoplasia in fetal alcohol exposed and adult rats

Activating transcription factor 4 underlies the pathogenesis of arsenic trioxide-mediated impairment of macrophage innate immune functions

Activation-induced T-cell death and immune dysfunction after implantation of left-ventricular assist device

Active immunization with amyloid-beta 1-42 impairs memory performance through TLR2/4-dependent activation of the innate immune system

Acute 3,4-methylenedioxymethamphetamine(MDMA) administration produces a rapid and sustained suppression of immune function in the rat

Acute and subchronic immunotoxicity of p-chloronitrobenzene in mice. I. Effect on natural killer, cytotoxic T-lymphocyte activities and mitogen-stimulated lymphocyte proliferation

Acute and subchronic toxic effects of atrazine and chlorpyrifos on common carp (*Cyprinus carpio* L.): Immunotoxicity assessments

Acute effects of 2-bromopropane and 1,2-dibromopropane on hepatotoxic and immunotoxic parameters in female BALB/c mice

Acute exposure to crotonaldehyde induces dysfunction of immune system in male Wistar rats

Acute exposure to perfluorononanoic acid in prepubertal mice: Effect on germ cell dynamics and an insight into the possible mechanisms of its inhibitory action on testicular functions

Acute exposure to waterborne cadmium induced oxidative stress and immunotoxicity in the brain, ovary and liver of zebrafish (*Danio rerio*)

Acute exposure to ZnO nanoparticles induces autophagic immune cell death

Acute immunoallergic hemolytic anemia induced by sulindac

Acute Immunotoxic Effects of Perfluorononanoic Acid (PFNA) in C57BL/6 Mice

Acute immunotoxicity of gallium to carp (*Cyprinus carpio* L)

Acute immunotoxicity of p-chloronitrobenzene in mice: II. Effect of p-chloronitrobenzene on the immunophenotype of murine splenocytes determined by flow cytometry

Acute infection of mice with highly virulent group B streptococci as a host resistance model for immunotoxicity assessment

Acute inflammation and immunoresponses induced by ortho-phthalaldehyde in mice

Acute kidney injury as the presenting complaint of ceftazidime-induced immune-mediated haemolysis

Acute microplastic exposure raises stress response and suppresses detoxification and immune capacities in the scleractinian coral *Pocillopora damicornis*

Acute myocardial infarction caused by delayed heparin-induced thrombocytopenia and acute immunoreaction due to re-exposure to heparin in a systemic lupus erythematosus patient with HIT antibodies

Acute rejection risk in kidney transplant recipients on steroid-avoidance immunosuppression receiving induction with either antithymocyte globulin or basiliximab

Acute renal failure induced by intravenous immune globulin

Acute symptomatic hypocalcemia from immune checkpoint therapy-induced hypoparathyroidism

Acute systemic DNA damage in youth does not impair immune defense with aging

Acute toxicity of cypermethrin on the juvenile of red claw crayfish *Cherax quadricarinatus*

Adaptive immunity is severely impaired by open-heart surgery

Adiponectin ameliorates dyslipidemia induced by the human immunodeficiency virus protease inhibitor ritonavir in mice

Adipose-Derived Mesenchymal Stem Cells Restore Impaired Mucosal Immune Responses in Aged Mice

Adjuvant and immuno-suppressive effect of six monophthalates in a subcutaneous injection model with BALB/c mice

Adjuvant immunization induces high levels of pathogenic antiphospholipid antibodies in genetically prone mice: another facet of the ASIA syndrome

Adrenal suppression in children with the human immunodeficiency virus treated with megestrol acetate

ADTP vaccine induces temporary immunosuppression correctable by using the immunomodulator purified staphylococcal anatoxin

Advances in immunologic mechanisms of lung inflammation induced by airborne particulate matter

Advances in mechanisms and signaling pathways of carbon nanotube toxicity

Advances in non-carcinogenic toxicity of trichloroethylene

Advances on the immunotoxicity of airborne fine particulate matter (PM<sub>2.5</sub>)

Adverse effects and immune dysfunction in response to oral administration of weathered Iranian heavy crude oil in the rockfish *Sebastes schlegeli*

Adverse experience in childhood as a developmental risk factor for altered immune status in adulthood

Adverse immunological effects and autoimmunity induced by dental amalgam and alloy in mice

Aflatoxin B interferes with the antigen-presenting capacity of porcine dendritic cells

Aflatoxin B1 Induces Immunotoxicity through the DNA Methyltransferase-Mediated JAK2/STAT3 Pathway in 3D4/21 Cells

Aflatoxin B1 invokes apoptosis via death receptor pathway in hepatocytes

Aflatoxin B1 Suppressed T-Cell Response to Anti-pig-CD3 Monoclonal Antibody Stimulation in Primary Porcine Splenocytes: A Role for the Extracellular Regulated Protein Kinase (ERK1/2) MAPK Signaling Pathway

Aflatoxin-related immune dysfunction in health and in human immunodeficiency virus disease

Aflatoxins: characteristics and impact on human health

After TGN1412: recent developments in cytokine release assays

Age and Early Graft Function Relate With Risk-Benefit Ratio of Allogenic Islet Transplantation Under Antithymocyte Globulin-Mycophenolate Mofetil-Tacrolimus Immune Suppression

Age and low levels of circulating vitamin D are associated with impaired innate immune function

Age-associated impairment of antitumor immunity in carcinoma-bearing mice and restoration by oral administration of Lentinula edodes mycelia extract

Age-Dependent Effects of Prenatal Dexamethasone Exposure on Immune Responses in Male Rats

Age-related differences in the sensitivity of the fish immune response to a coplanar PCB

Age-related differences in the toxicity of ochratoxin A in female rats

Age-related susceptibility to immunotoxicants: animal data and human parallels

Age-specific immunocompetence of the earthworm Eisenia andrei: exposure to methylmercury chloride

Aggravating effects of Asian sand dust on lung eosinophilia in mice immunized beforehand by ovalbumin

Aggravating impact of nanoparticles on immune-mediated pulmonary inflammation

AIDS, drugs of abuse and the immune system: a complex immunotoxicological network

Airborne fungus exposure prior to hospitalisation as risk factor for mould infections in immunocompromised patients

Airway inflammatory and immunological events in a rat model exposed to toluene diisocyanate

Alcohol abuse and Streptococcus pneumoniae infections: consideration of virulence factors and impaired immune responses

Alcohol abuse enhances neuroinflammation and impairs immune responses in an animal model of human immunodeficiency virus-1 encephalitis

Alcohol administration during adulthood induces alterations of parvalbumin and glial fibrillary acidic protein immunoreactivity in rat hippocampus and cingulate cortex

Alcohol and hepatitis C virus--interactions in immune dysfunctions and liver damage

Alcohol and HIV decrease proteasome and immunoproteasome function in macrophages: implications for impaired immune function during disease

Alcohol-induced alterations in serum immunoglobulin e (IgE) levels in human subjects

Alcohol-induced changes in the immune response: immunological effects of chronic ethanol intake are genetically regulated

Alcohol-induced suppression of tumor necrosis factor--a potential risk factor for secondary infection in the acquired immunodeficiency syndrome

Alcoholism causes alveolar macrophage zinc deficiency and immune dysfunction

Alemtuzumab induction and sirolimus plus mycophenolate mofetil maintenance for CN1 and steroid-free kidney transplant immunosuppression

Alginate oligosaccharide (AOS) improves immuno-metabolic systems by inhibiting STOML2 overexpression in high-fat-diet-induced obese zebrafish

Alkaline ceramidase 3 deficiency aggravates colitis and colitis-associated tumorigenesis in mice by hyperactivating the innate immune system

Allergic disease, immunoglobulins, exposure to mercury and dental amalgam in Swedish adolescents

Allergic Potential and Immunotoxicity Induced by Topical Application of 1-Chloro-4-(Trifluoromethyl)Benzene (PCBTF) in a Murine Model

Allergy and sensitization during childhood associated with prenatal and lactational exposure to marine pollutants

Allergy-Inducing Chromium Compounds Trigger Potent Innate Immune Stimulation Via ROS-Dependent Inflammasome Activation

Alleviation of lead-induced oxidative stress and immune damage by selenium in chicken bursa of Fabricius

Alleviative effect of licorice on copper chloride-induced oxidative stress in the brain: biochemical, histopathological, immunohistochemical, and genotoxic study

Allogeneic compact bone-derived mesenchymal stem cell transplantation increases survival of mice exposed to lethal total body irradiation: a potential immunological mechanism

Alpha tocopherol protects against immunosuppressive and immunotoxic effects of lead

alpha-Lipoic acid ameliorates oral mucositis and oxidative stress induced by methotrexate in rats. Histological and immunohistochemical study

Alpha-momorcharin possessing high immunogenicity, immunotoxicity and hepatotoxicity in SD rats

Alteration of adipose tissue immune cell milieu towards the suppression of inflammation in high fat diet fed mice by flaxseed oil supplementation

Alteration of basic fibroblast growth factor concentration and immunoreactivity in healing of ethanol-induced gastric mucosal damage: effect of sofalcone

Alterations in neuronal development in the substantia nigra pars compacta following in utero ethanol exposure: immunohistochemical and Golgi studies

Alterations in regulatory T-cells: rediscovered pathways in immunotoxicology

Alterations in the pulmonary and systemic immune response in rats exposed to coal fly ash

Alterations of cytokines and MAPK signaling pathways are related to the immunotoxic effect of perfluorononanoic acid

Alterations of immune functions in barrier disrupted skin by UVB irradiation

Alterations of immunoreactivity in chemical production workers in dependence on dose exposure to toxicants

Alterations of seizure-induced c-fos immunolabelling and gene expression in the rat cerebral cortex following dexamethasone treatment

Altered cytokine production in mice exposed to lead acetate

Altered epithelial density and expansion of bulbar projections of a discrete HSP70 immunoreactive subpopulation of rat olfactory receptor neurons in reconstituting olfactory epithelium following exposure to methyl bromide

Altered helper and suppressor lymphocyte populations in surgical patients. A measure of postoperative immunosuppression

Altered hematological and immunological parameters in silver catfish (*Rhamdia quelen*) following short term exposure to sublethal concentration of glyphosate

Altered hepatic mRNA expression of immune response-associated DNA damage in mice liver induced by potassium bromate: Protective role of vanillin

Altered immunohistochemical localization of basic fibroblast growth factor after bleomycin-induced lung injury

Altered MARCH1 ubiquitination-regulated dendritic cell immune functions during the early stage of zymosan-induced multiple organ dysfunction syndrome (MODS) in mice

Altered or Impaired Immune Response to Hepatitis B Vaccine in WNIN/GR-Ob Rat: An Obese Rat Model with Impaired Glucose Tolerance

Altered serum cytokine and immunoglobulin levels in the workers exposed to antimony

Aluminum adjuvant potentiates gilthead seabream immune responses but induces toxicity in splenic melanomacrophage centers

Aluminum chloride- and norepinephrine-induced immunotoxicity on splenic lymphocytes by activating beta2-AR/cAMP/PKA/NF-kappaB signal pathway in rats

Ambient air pollution associated with suppressed serologic responses to *Pneumocystis jirovecii* in a prospective cohort of HIV-infected patients with *Pneumocystis pneumonia*

Amelioration of 2,4,6-trinitrobenzene sulfonic acid-induced colitis in mice by immunoregulatory dendritic cells

Amelioration of doxorubicin-induced myocardial oxidative stress and immunosuppression by grape seed proanthocyanidins in tumour-bearing mice

Amelioration of Ochratoxin A-induced immunotoxic effects by silymarin and Vitamin E in White Leghorn cockerels

Ameliorative action of *Vernonia cinerea* L. on cyclophosphamide-induced immunosuppression and oxidative stress in mice

Ameliorative effect of nanoencapsulated flavonoid against chlorpyrifos-induced hepatic oxidative damage and immunotoxicity in Wistar rats

Ameliorative Effects of Grape Seed Proanthocyanidin Extract on Growth Performance, Immune Function, Antioxidant Capacity, Biochemical Constituents, Liver Histopathology and Aflatoxin Residues in Broilers Exposed to Aflatoxin B

Amiodarone-induced thyrotoxicosis and thyroid cancer: clinical, immunohistochemical, and molecular genetic studies of a case and review of the literature

Amlodipine-induced immune thrombocytopenia

Ammonia exposure alters the expression of immune-related and antioxidant enzymes-related genes and the gut microbial community of crucian carp (*Carassius auratus*)

Ammonia inhalation impaired immune function and mitochondrial integrity in the broilers bursa of fabricius: Implication of oxidative stress and apoptosis

Ammonia toxicity induces glutamine accumulation, oxidative stress and immunosuppression in juvenile yellow catfish *Pelteobagrus fulvidraco*

Amoxicillin-clavulanic acid-induced linear immunoglobulin A bullous dermatosis: case report and review of the literature

AMP-Conjugated Quantum Dots: Low Immunotoxicity Both In Vitro and In Vivo

Amphotericin-induced stridor: a review of stridor, amphotericin preparations, and their immunoregulatory effects

AMPK agonist downregulates innate and adaptive immune responses in TNBS-induced murine acute and relapsing colitis

Amprenavir-induced maculopapular exanthem followed by desensitization in a patient with late-stage human immunodeficiency virus



Amyloid beta protein in rat soleus muscle in chloroquine-induced myopathy using end-specific antibodies for A beta 40 and A beta 42: immunohistochemical evidence for amyloid beta protein

An action spectrum for ultraviolet radiation-induced immunosuppression in humans

An antibody to morphine as an indicator of chronic morphine intoxication and disrupted immunologic reactivity

An anti-murine CD3 monoclonal antibody with a low affinity for Fc gamma receptors suppresses transplantation responses while minimizing acute toxicity and immunogenicity

An aryl hydrocarbon receptor independent mechanism of JP-8 jet fuel immunotoxicity in Ah-responsive and Ah-nonresponsive mice

An atypical immune-inflammatory disorder secondary to breast implant exposure

An enzyme-linked immunosorbent assay (ELISA) specific for antibodies to TNP-LPS detects alterations in serum immunoglobulins and isotype switching in C57BL/6 and DBA/2 mice exposed to 2,3,7,8-tetrachlorodibenzo-p-dioxin and related compounds

An epidemic ignored. Endometriosis linked to dioxin and immunologic dysfunction

An evaluation of the immunotoxic potential of isobutyl nitrite

An examination of the effect of aerosolized Permanone insecticide on zebra finch susceptibility to West Nile virus

An experimental approach to studying the effectiveness and safety of meglumine antimoniate formulations

An F1-extended one-generation reproductive toxicity study in CrI:CD(SD) rats with 2,4-dichlorophenoxyacetic acid

An Immune Suppression-associated EBV-positive Anaplastic Large Cell Lymphoma With a BRAF V600E Mutation

An immunohistochemical study of dibutyltin-induced thymus atrophy

An immunohistochemical study on mild skin irritation induced by a single application of a low-molarity sodium dodecyl sulfate solution: keys to the prevention of irritant contact dermatitis

An immunological study of workers exposed to diphenylmethane diisocyanate (MDI)

An immunotoxicity assessment of food flavouring ingredients

An immunotoxicological evaluation of 4,4'-thiobis-(6-t-butyl-m-cresol) in female B6C3F1 mice. 1. Body and organ weights, hematology, serum chemistries, bone marrow cellularity, and hepatic microsomal parameters

An immunotoxicological evaluation of 4,4'-thiobis-(6-t-butyl-m-cresol) in female B6C3F1 mice. 2. Humoral and cell-mediated immunity, macrophage function, and host resistance

An improved definition of immune heparin-induced thrombocytopenia in postoperative orthopedic patients

An in vitro model of human acute ethanol exposure that incorporates CXCR3- and CXCR4-dependent recruitment of immune cells

An induction versus no-induction protocol in anticalcineurin-based immunosuppression using very low-dose steroids

An injectable, low-toxicity phospholipid-based phase separation gel that induces strong and persistent immune responses in mice

An international validation study of the IL-2 Luc assay for evaluating the potential immunotoxic effects of chemicals on T cells and a proposal for reference data for immunotoxic chemicals

An investigation of the immunotoxicity of oil sands processed water and leachates in trout leukocytes

An outline of chloro-organic compound toxicology

An update to the toxicological profile for water-soluble and sparingly soluble tungsten substances

Anaemia, hypothyroidism and immune suppression associated with polychlorinated biphenyl exposure in bottlenose dolphins (*Tursiops truncatus*)

Analysis of adverse reactions induced by subcutaneous immunotherapy against dust mite allergy in 234 cases with allergic rhinitis and asthma

Analysis of allogeneic hematopoietic stem cell transplantation with high-dose cyclophosphamide-induced immune tolerance for severe aplastic anemia

Analysis of apoptosis of lymphoid cells in fish exposed to immunotoxic compounds

Analysis of glycerophospholipid metabolism after exposure to PCB153 in PC12 cells through targeted lipidomics by UHPLC-MS/MS

Analysis of heavy metal immunotoxicity by multiparameter flow cytometry: correlation of flow cytometry and immune function data in B6CF1 mice

Analysis of hematologic alterations, immune responses and metallothionein gene expression in Nile tilapia (*Oreochromis niloticus*) exposed to silver nanoparticles

Analysis of immune privilege in eyes with *Mycobacteria tuberculosis* adjuvant-induced uveitis

Analysis of immunomodulating nitric oxide, iNOS and cytokines mRNA in mouse macrophages induced by microcystin-LR

Analysis of proliferative activity in oral gingival epithelium in immunosuppressive medication induced gingival overgrowth

Analysis of the allergic reaction types of Chinese medicine injection based on immunotoxicity

Anatoxin-a induces apoptosis of leukocytes and decreases the proliferative ability of lymphocytes of common carp (*Cyprinus carpio* L.) in vitro

Energy, immunosuppressive serum, and impaired lymphocyte blastogenesis in burn patients

ANN multiplexing model of drugs effect on macrophages; theoretical and flow cytometry study on the cytotoxicity of the anti-microbial drug G1 in spleen

Anthrax lethal toxin impairs innate immune functions of alveolar macrophages and facilitates *Bacillus anthracis* survival

Anthrax lethal toxin-mediated killing of human and murine dendritic cells impairs the adaptive immune response

Anti PF4/heparin antibodies detection by immunoturbidimetry in the diagnosis of heparin-induced thrombocytopenia

Anti-apoptotic, anti-inflammatory, and immunomodulatory activities of 3,3'-diselenodipropionic acid in mice exposed to whole body gamma-radiation

Antibodies in sulfonamide-induced immune thrombocytopenia recognize calcium-dependent epitopes on the glycoprotein IIb/IIIa complex

Antibodies to morphine as indicators of chronic morphine intoxication and impaired immune reactivity

Antibodies to the costimulatory molecule CD86 interfere with ultraviolet radiation-induced immune suppression

Antibody-mediated immunotoxicity in American kestrels (*Falco sparverius*) exposed to polychlorinated biphenyls

Anti-CD3 immunotoxin prevents low-dose STZ/interferon-induced autoimmune diabetes in mouse

Anti-CD40 Treatment of 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD)-exposed C57Bl/6 mice induces activation of antigen presenting cells yet fails to overcome TCDD-induced suppression of allograft immunity

Anticholinesterase mechanism as a factor of immunotoxicity of various chemical compounds

Anticonvulsant-induced marrow suppression and immune thrombocytopenia

Anticonvulsant-induced toxic epidermal necrolysis: monitoring the immunologic response

Anti-D immunoglobulin-induced prolonged intravascular hemolysis and neutropenia

Anti-dopamine beta-hydroxylase immunotoxin-induced sympathectomy in adult rats

Antifouling biocides: Impairment of bivalve immune system by chlorothalonil

Antigen-Specific Gut Inflammation and Systemic Immune Responses Induced by Prolonging Wheat Gluten Sensitization in BALB/c Murine Model

Anti-glucocorticoid-induced Tumor Necrosis Factor-Related Protein (GITR) Therapy Overcomes Radiation-Induced Treg Immunosuppression and Drives Abscopal Effects

Antigranulocyte antibodies and deranged immune function associated with phenytoin-induced serum sickness

Antihapten antibodies in workers exposed to trimellitic anhydride fumes: a potential immunopathogenetic mechanism for the trimellitic anhydride pulmonary disease--anemia syndrome

Anti-HMGCR antibodies demonstrate high diagnostic value in the diagnosis of immune-mediated necrotizing myopathy following statin exposure

Anti-inflammatory and immunomodulatory effects of *Ulomoides dermestoides* on induced pleurisy in rats and lymphoproliferation in vitro

Antilymphocyte globulin with a small dose of cyclosporine A and prednisone as the induction of immunosuppression in renal allograft recipients

Antimicrobial activity against oral pathogens and immunomodulatory effects and toxicity of geopropolis produced by the stingless bee *Melipona fasciculata* Smith

Antioxidant and anti-inflammatory effects of N-acetylcysteine against malathion-induced liver damages and immunotoxicity in rats

Antioxidant compounds in the seaweed *Gelidiella acerosa* protects human Peripheral Blood Mononuclear Cells against TCDD induced toxicity

Antioxidant eugenosedin-A protects against lipopolysaccharide-induced hypotension, hyperglycaemia and cytokine immunoreactivity in rats and mice

Antioxidant therapy partially blocks immune-induced lung fibrosis

Anti-oxidant vitamins reduce normal tissue toxicity induced by radio-immunotherapy

Anti-oxidative and anti-inflammatory activities of placental extracts in benzo[a]pyrene-exposed rats

Anti-PF4-heparin immunoglobulin G is the major class of heparin-induced thrombocytopenia antibody: findings of an enzyme-linked immunofiltration assay using membrane-bound hPF4-heparin

Anti-Programmed Cell Death Protein-1 Immunotherapy for Glioblastoma is Impaired by Systemic Chemotherapy but Enhanced in Combination With Locally Delivered Chemotherapy

Antiretroviral therapy, immune suppression and renal impairment in HIV-positive persons

Antithymocyte globulin or interleukin 2 receptor antibody for immunosuppressive induction therapy after orthotopic liver transplantation: a follow-up study

Antituberculosis drug-induced hepatotoxicity. The role of hepatitis C virus and the human immunodeficiency virus

Antitumor effects of doxorubicin against a virally-induced rat osteosarcoma with minimal immunosuppression

Antiviral activity of recombinant rat interferon gamma in immunologically impaired and immunosuppressed rats

Antiviral Activity, Safety, and Exposure-Response Relationships of GSK3532795, a Second-Generation Human Immunodeficiency Virus Type 1 Maturation Inhibitor, Administered as Monotherapy or in Combination With Atazanavir With or Without Ritonavir in a Phase 2a Randomized, Dose-Ranging, Controlled Trial (A1468002)

Apigenin reverses lung injury and immunotoxicity in paraquat-treated mice

Apoptosis and cell proliferation in short-term and long-term effects of radioiodine-131-induced kidney damage: an experimental and immunohistochemical study

Apoptosis as a mechanism of tributyltin cytotoxicity to thymocytes: relationship of apoptotic markers to biochemical and cellular effects

Apoptosis in immunocytes induced by several types of pesticides

Apoptosis-mediated immunotoxicity of polychlorinated biphenyls (PCBs) in murine splenocytes

Apparent immunotoxic response to phenolic compounds

Application of a biologically-based RFD estimation method to tetrachlorodibenzo-p-dioxin (TCDD) mediated immune suppression and enzyme induction

Application of a novel integrated toxicity testing strategy incorporating "3R" principles of animal research to evaluate the safety of a new agrochemical sulfoxaflor

Application of an immunoperoxidase staining method for detection of 7,8-dihydro-8-oxodeoxyguanosine as a biomarker of chemical-induced oxidative stress in marine organisms

Application of benchmark analysis for mixed contaminant exposures: Mutual adjustment of perfluoroalkylate substances associated with immunotoxicity

Application of cellular biosensors for detection of atypical toxic bioactivity in microcystin-containing cyanobacterial extracts

Application of immunosuppressants for therapy of patients with vasculitis

Arabinosylated lipoarabinomannan modulates the impaired cell mediated immune response in Mycobacterium tuberculosis H37Rv infected C57BL/6 mice

Are iron oxide nanoparticles safe? Current knowledge and future perspectives

Are serum levels of immunoglobulin classes and IgG subclasses involved in delayed pulmonary complications induced by sulfur mustard? Sardasht-Iran Cohort Study

Are suppressor T-cells the primary target cells of lead immunotoxicity

Argemone oil, an edible oil adulterant, induces systemic immunosuppression in Balb/c mice in an oral 28 days repeated dose toxicity study

Aroclor 1254 exposure reduces disease resistance and innate immune responses in fasted Arctic charr

Aroclor 1254 inhibits the chemiluminescence response of peritoneal cavity cells from sharpsnout sea bream (*Diplodus puntazzo*)

Aromatic retinoid Ro 40-8757 reduces immunotoxicities of cyclophosphamide as revealed by immunohistochemical staining of lymphoid tissues and general pathologic examinations

Arsenic (III) or/and copper (II) exposure induce immunotoxicity through trigger oxidative stress, inflammation and immune imbalance in the bursa of chicken

Arsenic and Immune Response to Infection During Pregnancy and Early Life

Arsenic decreases RXRalpha-dependent transcription of CYP3A and suppresses immune regulators in hepatocytes

Arsenic exposure and cell-mediated immunity in pre-school children in rural Bangladesh

Arsenic Exposure and Immunotoxicity: a Review Including the Possible Influence of Age and Sex

Arsenic immunotoxicity and immunomodulation by phytochemicals: potential relations to develop chemopreventive approaches

Arsenic immunotoxicity: a review

Arsenic-associated oxidative stress, inflammation, and immune disruption in human placenta and cord blood

Arsenic-Induced Carcinogenesis and Immune Dysregulation

Arsenic-induced dose-dependent modulation of the NF-kappaB/IL-6 axis in thymocytes triggers differential immune responses

Aryl hydrocarbon receptor (AhR) agonists suppress interleukin-6 expression by bone marrow stromal cells: an immunotoxicology study

Asbestos induces reduction of tumor immunity

Asbestos-induced cellular and molecular alteration of immunocompetent cells and their relationship with chronic inflammation and carcinogenesis

Aseptic meningitis induced by intravenous immunoglobulins in a child with acute Epstein-Barr virus infection and thrombocytopenia

Aspects of the immunotoxicity of chronic tobacco smoke exposure of the rat

Assessing a theoretical risk of dolutegravir-induced developmental immunotoxicity in juvenile rats

Assessing environmental exposure in children: immunotoxicology screening

Assessing the clinical and cost impact of on-demand immunoassay testing for the diagnosis of heparin induced thrombocytopenia

Assessment by c-Fos immunostaining of changes in brain neural activity induced by 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD) and leptin in rats

Assessment of apoptosis in thymocytes and splenocytes from mice exposed to arsenate in drinking water: cytotoxic effects of arsenate on the cells in vitro

Assessment of apoptosis in xenobiotic-induced immunotoxicity

Assessment of dietary fat intake and innate immune activation as risk factors for impaired lung function

Assessment of immune status of yellowfin seabream (*Acanthopagrus latus*) during short term exposure to phenanthrene

Assessment of immunotoxicity and genotoxicity in workers exposed to low concentrations of formaldehyde

Assessment of immunotoxicity in female Fischer 344/N and Sprague Dawley rats and female B6C3F1 mice exposed to hexavalent chromium via the drinking water

Assessment of immunotoxicity induced by chemicals in human precision-cut lung slices (PCLS)

Assessment of immunotoxicity of buprenorphine

Assessment of immunotoxicity of dibutyl phthalate using live zebrafish embryos

Assessment of immunotoxicity of irinotecan determined by the novel method, by which productivity of TNF-alpha from whole blood is stimulated by lipopolysaccharide

Assessment of immunotoxicity parameters in individuals occupationally exposed to lead

Assessment of lymphocyte subpopulations and cytokine secretion in children exposed to arsenic

Assessment of myelotoxicity caused by environmental chemicals

Assessment of possible immunotoxicity of the antipsychotic drug clozapine

Assessment of potential immunotoxic effects caused by cypermethrin, fluoxetine, and thiabendazole using heat shock protein 70 and interleukin-1beta mRNA expression in the anuran *Xenopus laevis*

Assessment of recent developmental immunotoxicity studies with bisphenol A in the context of the 2015 EFSA t-TDI

Assessment of the immunotoxic potential of the fungicide dinocap in mice

Assessment of the immunotoxic potential of trichloroethylene and perchloroethylene in rats following inhalation exposure

Assessment of the mechanism of astrocyte swelling induced by the macrolide immunosuppressant sirolimus using multinuclear nuclear magnetic resonance spectroscopy

Assessment of the Status of Effector and Regulatory Components of Immune System in Chelyabinsk Region Residents Exposed to Radiation Due to Residence in the Area Contaminated as a Result of the Radiation Accident at Mayak PA and in Their Offspring

Assessment of the usefulness of the murine cytotoxic T cell line CTLL-2 for immunotoxicity screening by transcriptomics

Assessment of trichloroethylene (TCE) exposure in murine strains genetically-prone and non-prone to develop autoimmune disease

Association between abacavir exposure and increased risk for cardiovascular disease in patients with human immunodeficiency virus

Association between chronic organochlorine exposure and immunotoxicity in the round stingray (*Uroobatis halleri*)

Association between lymphocyte proliferation and polychlorinated biphenyls in free-ranging harbor seal (*Phoca vitulina*) pups from British Columbia, Canada

Association between prenatal exposure to perfluorinated compounds and symptoms of infections at age 1-4years among 359 children in the Odense Child Cohort

Association Between Renal Dysfunction and Major Adverse Cardiac Events After Liver Transplantation: Evidence from an International Randomized Trial of Everolimus-Based Immunosuppression

Association of antibody induction immunosuppression with cancer after kidney transplantation

Association of life style with high risk of hyperimmunity and of immunosuppression mediated by IgE

Association of mitochondrial dysfunction with oxidative stress and immune suppression in Blunt Snout Bream *Megalobrama amblycephala* fed a high-fat diet

Association of serum immunoglobulins levels and eye injuries in sulfur mustard exposed: Sardasht-Iran Cohort Study

Association of transcription-coupled repair but not global genome repair with ultraviolet-B-induced Langerhans cell depletion and local immunosuppression

Ataxin-1 oligomers induce local spread of pathology and decreasing them by passive immunization slows Spinocerebellar ataxia type 1 phenotypes

Atezolizumab after Nivolumab-Induced Inflammatory Polyarthritis: Can Anti-PD-L1 Immunotherapy Be Administered after Anti-PD-1-Related Immune Toxicities

Atmospheric H<sub>2</sub>S triggers immune damage by activating the TLR-7/MyD88/NF-kappaB pathway and NLRP3 inflammasome in broiler thymus

Atomic layer deposition coating of carbon nanotubes with zinc oxide causes acute phase immune responses in human monocytes in vitro and in mice after pulmonary exposure

Atrazine hinders PMA-induced neutrophil extracellular traps in carp via the promotion of apoptosis and inhibition of ROS burst, autophagy and glycolysis

Atrazine induces endoplasmic reticulum stress-mediated apoptosis of T lymphocytes via the caspase-8-dependent pathway

Atrazine induces necroptosis by miR-181-5p targeting inflammation and glycometabolism in carp lymphocytes

Atrazine promotes immunomodulation by melanomacrophage centre alterations in spleen and vascular disorders in gills from *Oreochromis niloticus*



Atrazine-induced apoptosis of splenocytes in BALB/C mice

Attenuated Bordetella pertussis vaccine strain BPZE1 modulates allergen-induced immunity and prevents allergic pulmonary pathology in a murine model

Attenuating immune pathology using a microbial-based intervention in a mouse model of cigarette smoke-induced lung inflammation

Attenuation of catecholamine-induced immunosuppression in whole blood from patients with sepsis

Attenuation of inducible respiratory immune responses by oseltamivir treatment in mice infected with influenza A virus

Attenuation of phosphamidon-induced oxidative stress and immune dysfunction in rats treated with N-acetylcysteine

Attenuation of verapamil-induced myocardial toxicity in an ex-vivo rat model using a verapamil-specific ovine immunoglobulin

Attractive and immunosuppressive properties of volatile secretions induced in mice separately and combine influence of ionizing radiation and cyclophosphamide

Author Correction: Immunotoxicological impact and biodistribution assessment of bismuth selenide (Bi<sub>2</sub>Se<sub>3</sub>) nanoparticles following intratracheal instillation in mice

Autoantibodies and drug- or metabolite-dependent antibodies in patients with diclofenac-induced immune haemolysis

Autoimmune diseases after hepatitis B immunization in adults: Literature review and meta-analysis, with reference to 'autoimmune/autoinflammatory syndrome induced by adjuvants' (ASIA)

Autoimmune diseases after unusual exposure to silica or silicones. 3 cases

Autoimmune dysthyroidism induced by alpha interferon in two female patients with chronic non-A, non-B hepatitis

Autoimmune hemolytic anemia induced by alpha-methyldopa (Aldomet)

Autoimmune hemolytic anemia induced by mefenamic acid

Autoimmune Hepatitis (Immune-Mediated Liver Injury) Induced By Rosuvastatin

Autoimmune hepatitis induced by fibrates

Autoimmune hepatitis triggered by nitrofurantoin: A rare drug-induced toxicity

Autoimmune Neuromuscular Diseases Induced by Immunomodulating Drugs

Auto-immune polyradiculoneuropathy and a novel IgG biomarker in workers exposed to aerosolized porcine brain

Auto-immune thyroid dysfunction induced by tyrosine kinase inhibitors in a patient with recurrent chordoma

Autoimmune/inflammatory syndrome induced by adjuvants (Shoenfeld's syndrome): clinical and immunological spectrum

Autoimmunity by pesticides: a critical review of the state of the science

Autoimmunity induced by adjuvant hydrocarbon oil components of vaccine

Autoimmunity induced by drugs. Immunological characteristics and etiopathogenic hypotheses

Autoimmunity: Impaired mucosal immunity in patients with SLE

Autoimmunization induced by interferon alpha therapy in chronic hepatitis C

Autonomic dysreflexia causes chronic immune suppression after spinal cord injury

Autophagy protects murine macrophages from beta-cypermethrin-induced mitochondrial dysfunction and cytotoxicity via the reduction of oxidation stress

Avermectin induced autophagy in pigeon spleen tissues

Avermectin induces the oxidative stress, genotoxicity, and immunological responses in the Chinese Mitten Crab, *Eriocheir sinensis*

B and T lymphocyte attenuator inhibits LPS-induced endotoxic shock by suppressing Toll-like receptor 4 signaling in innate immune cells

B7-1-transfected tumor vaccine counteracts chemotherapy-induced immunosuppression and prolongs the survival of rats bearing highly metastatic osteosarcoma cells

Bacillus Calmette-Guerin (BCG) and immunoglobulins synergistically enhance mineral dust-induced production of reactive oxygen metabolites by human monocytes

Bacterial lipopolysaccharides induce peripheral nerve disturbances in rats that mimic human immune-mediated polyneuropathies

Bacterial Pneumonia-induced Persistent Remission of Severe Immune Thrombocytopenia after Allogeneic Hematopoietic Stem Cell Transplantation

Bacteroides uniformis CECT 7771 ameliorates metabolic and immunological dysfunction in mice with high-fat-diet induced obesity

Bariatric Surgery Induces Disruption in Inflammatory Signaling Pathways Mediated by Immune Cells in Adipose Tissue: A RNA-Seq Study

Baseline Immunoglobulin E Levels as a Marker of Doxorubicin- and Trastuzumab-Associated Cardiac Dysfunction

BDE-47 exposure changed the immune function of haemocytes in *Mytilus edulis*: An explanation based on ROS-mediated pathway

Beauvericin and enniatin B effects on a human lymphoblastoid Jurkat T-cell model

Behavior of various factors of non-specific immunity and immunoglobulins in the oral cavity and blood of subjects exposed to chemical stress

Behavioral and immunotoxic effects of Prograf (tacrolimus) in the male Siamese fighting fish

Behavioral changes with alterations of choline acetyltransferase immunoreactivities induced by N-butyl benzenesulfonamide

Behavioural endocrine immune-conditioned response is induced by taste and superantigen pairing

Belatacept as maintenance immunosuppression for postrenal transplant de novo drug-induced thrombotic microangiopathy

Below background levels of blood lead impact cytokine levels in male and female mice

Bendamustine associated immune suppression and infections during therapy of hematological malignancies

Bendamustine can severely impair T-cell immunity against cytomegalovirus

Beneficial effect of *Sepia esculenta* ink polysaccharide on cyclophosphamide-induced immunosuppression and ovarian failure in mice

Beneficial effects of vitamin C treatment on pregnant rats exposed to formaldehyde: Reversal of immunosuppression in the offspring

Benzanthrone induced immunotoxicity via oxidative stress and inflammatory mediators in Balb/c mice

Benzo[a]pyrene exposure under future ocean acidification scenarios weakens the immune responses of blood clam, *Tegillarca granosa*

Benzo[a]pyrene Induces Autophagic and Pyroptotic Death Simultaneously in HL-7702 Human Normal Liver Cells

Benzo[a]pyrene-induced immunotoxicity in Japanese medaka (*Oryzias latipes*): relationship between lymphoid CYP1A activity and humoral immune suppression

Benzo[a]pyrene-induced immunotoxicity: comparison to DNA adduct formation in vivo, in cultured splenocytes, and in microsomal systems

Benzylacetylcholine reverses azidothymidine-induced marrow suppression without impairment of anti-human immunodeficiency virus activity

Berberine is a dopamine D1- and D2-like receptor antagonist and ameliorates experimentally induced colitis by suppressing innate and adaptive immune responses

Beryllium-induced disturbances of the murine immune system reflect some phenomena observed in sarcoidosis

beta-(1,3)-Glucan exposure assessment by passive airborne dust sampling and new sensitive immunoassays

beta5i subunit deficiency of the immunoproteasome leads to reduced Th2 response in OVA induced acute asthma

beta-Cypermethrin and its metabolite 3-phenoxybenzoic acid exhibit immunotoxicity in murine macrophages

Beta-glucan ameliorates methotrexate-induced oxidative organ injury via its antioxidant and immunomodulatory effects

Beta-trace Protein as a new non-invasive immunological Marker for Quinolinic Acid-induced impaired Blood-Brain Barrier Integrity

Bevacizumab-induced immunoglobulin A vasculitis with nephritis: A case report

Bifenthrin induces developmental immunotoxicity and vascular malformation during zebrafish embryogenesis

Bilirubin augments radiation injury and leads to increased infection and mortality in mice: molecular mechanisms

Bioaccumulation of lead and non-specific immune responses in white shrimp (*Litopenaeus vannamei*) to Pb exposure

Bioavailability and immunotoxicity of silver nanoparticles to the freshwater mussel *Elliptio complanata*

Biochemical and immunological basis of silymarin effect, a milk thistle (*Silybum marianum*) against ethanol-induced oxidative damage

Biochemical and immunological characterisation of mutants induced in V79 Chinese hamster cells by a benzo[a]pyrene dilepoxide

Biochemical and immunotoxicological alterations following repeated gallium arsenide exposure and their recoveries by meso-2,3-dimercaptosuccinic acid and 2,3-dimercaptopropane 1-sulfonate administration in rats

Bioconcentration and immunotoxicity of an experimental oil spill in European sea bass (*Dicentrarchus labrax* L

Biological and immunotoxicity evaluation of antimicrobial peptide-loaded coatings using a layer-by-layer process on titanium

Biological effects of gestational and lactational PCB exposure in neonatal and juvenile C57BL/6 mice

Biology of diesel exhaust effects on allergic pulmonary inflammation

Biomarkers for responses to heavy metals

Biomarkers of immune dysfunction following combination antiretroviral therapy for HIV infection

Biomarkers of immunotoxicity in fish and other non-mammalian sentinel species: predictive value for mammals

Biomarkers of immunotoxicity in fish: from the lab to the ocean

Biomarkers of methylmercury exposure immunotoxicity among fish consumers in Amazonian Brazil

Biomonitoring human exposure to organohalogenated substances by measuring urinary chlorophenols using a high-throughput screening (HTS) immunochemical method

Biomonitoring of a population of Portuguese workers exposed to lead

Birth defects in juvenile Wistar rats after exposure to immunosuppressive drugs during pregnancy

Bisphenol A (BPA) Exposure In Utero Leads to Immunoregulatory Cytokine Dysregulation in the Mouse Mammary Gland: A Potential Mechanism Programming Breast Cancer Risk

Bisphenol A and immunotoxic potential: A commentary

Bisphenol A causes hyperactivity in the rat concomitantly with impairment of tyrosine hydroxylase immunoreactivity

Bisphenol A Disrupts HNF4alpha-Regulated Gene Networks Linking to Prostate Preneoplasia and Immune Disruption in Noble Rats

Bisphenol A exposure alters release of immune and developmental modulators and expression of estrogen receptors in human fetal lung fibroblasts

Bleomycin-induced lung injury in baboons: alteration of cells and immunoglobulins recoverable by bronchoalveolar lavage

Blocking Sympathetic Nervous System Reverses Partially Stroke-Induced Immunosuppression but does not Aggravate Functional Outcome After Experimental Stroke in Rats

Blood transfusion-induced immunomodulation--is storage time important

Bone marrow changes induced by recombinant granulocyte colony-stimulating factor resembling metastatic carcinoma: distinction with cytochemical and immunohistochemical studies

Boric acid-induced immunotoxicity and genotoxicity in model insect *Galleria mellonella* L. (Lepidoptera: Pyralidae)

Bortezomib significantly impairs the immunostimulatory capacity of human myeloid blood dendritic cells

Brain met-enkephalin immunostaining after subacute and subchronic exposure to benzene

Brentuximab vedotin combined with donor lymphocyte infusions for early relapse of Hodgkin lymphoma after allogeneic stem-cell transplantation induces tumor-specific immunity and sustained clinical remission

Brief butyltin exposure induces irreversible inhibition of the cytotoxic function on human natural killer cells, in vitro

Broad-spectrum sunscreens provide better protection from solar ultraviolet-simulated radiation and natural sunlight-induced immunosuppression in human beings

Bucillamine induced pulmonary injury occurs with immunoglobulin decrease

Bursopentin (BP5) protects dendritic cells from lipopolysaccharide-induced oxidative stress for immunosuppression

Cadmium and chlorpyrifos inhibit cellular immune response in spleen of rats

Cadmium and T cell differentiation: limited impact in vivo but significant toxicity in fetal thymus organ culture

Cadmium-induced endoplasmic reticulum stress in chicken neutrophils is alleviated by selenium

Cadmium-induced immune abnormality is a key pathogenic event in human and rat models of preeclampsia

Cadmium-induced immunotoxicity

Cadmium-induced suppression of the primary immune response and acute toxicity in mice: differential interaction of zinc

Calcineurin inhibitor Tacrolimus impairs host immune response against urinary tract infection

Calcineurin inhibitor-induced chronic nephrotoxicity in liver transplant patients is reversible using rapamycin as the primary immunosuppressive agent

Calcitonin gene-related peptide and nitric oxide are involved in ultraviolet radiation-induced immunosuppression

Calcium-binding proteins and GFAP immunoreactivity alterations in murine hippocampus after 1 month of exposure to 835 MHz radiofrequency at SAR values of 1.6 and 4.0 W/kg

Calcium-mediated activation of c-Jun NH2-terminal kinase (JNK) and apoptosis in response to cadmium in murine macrophages

Calretinin immunoreactivity in normal and carbon tetrachloride-induced nephrotoxic rats

Calretinin-containing neurons in trimethyltin-induced neurodegeneration in the rat hippocampus: an immunocytochemical study

Camel milk exosomes modulate cyclophosphamide-induced oxidative stress and immuno-toxicity in rats

Can prophylactic application of immunoglobulin decrease radiotherapy-induced oral mucositis

Cancer Immunotherapy with Anti-CTLA-4 Monoclonal Antibodies Induces an Inflammatory Bowel Disease

Cancer induction by immunosuppression in psoriasis after heavy PUVA treatment

Candida albicans derived fungal PAMPS, CAWS, water soluble mannoprotein-beta-glucan complex shows similar immunotoxicological activity with bacterial endotoxin from Escherichia coli O9

Candida albicans infection enhances immunosuppression induced by cyclophosphamide by selective priming of suppressive myeloid progenitors for NO production

Capture ELISA and flow cytometry methods for toxicologic assessment following immunization and cyclophosphamide challenges in beagles

Carbamazepine induced transient monoclonal gammopathy and immunodeficiency

Carbamazepine-induced immune thrombocytopenia

Carbamazepine-induced pseudolymphoma and immune dysregulation

Carbendazim has the potential to induce oxidative stress, apoptosis, immunotoxicity and endocrine disruption during zebrafish larvae development

Carbofuran suppresses T-cell-mediated immune responses by the suppression of T-cell responsiveness, the differential inhibition of cytokine production, and NO production in macrophages

Carbon fullerenes (C60s) can induce inflammatory responses in the lung of mice

Carbon tetrachloride is immunosuppressive and decreases host resistance to *Listeria monocytogenes* and *Streptococcus pneumoniae* in female B6C3F1 mice

Carbon tetrachloride suppresses T-cell-dependent immune responses by induction of transforming growth factor-beta 1

Carboplatin-induced immune hemolytic anemia

Carcinogen-induced inflammation and immunosuppression are enhanced in xeroderma pigmentosum group A model mice associated with hyperproduction of prostaglandin E2

Cardiac allograft tolerance induction with limited immunosuppression

Cardioprotective effect of green tea extract and vitamin E on Cisplatin-induced cardiotoxicity in mice: Toxicological, histological and immunohistochemical studies

Case of hyper-IgM immunodeficiency caused by B cell dysfunction following long-term administration of diphenylhydantoin

Case of vancomycin-induced linear immunoglobulin A bullous dermatosis

Case-control study on concentrations of organohalogen compounds and titers of antibodies to Epstein-Barr virus antigens in the etiology of non-Hodgkin lymphoma

Cecal ligation and puncture-induced impairment of innate immune function does not occur in the absence of caspase-1

Cefotaxime-induced immune hemolytic anemia due to antibodies reacting in vitro by more than one mechanism

Cefotetan-induced immune hemolytic anemia

Cefotetan-induced immune hemolytic anemia following prophylaxis for cesarean delivery

Ceftaroline modulates the innate immune and host defense responses of immunocompetent cells exposed to cigarette smoke

Ceftazidime-induced hemolysis in a patient with drug-dependent antibodies reactive by immune complex and drug adsorption mechanisms

Ceftriaxone induced immune hemolytic anemia: detection of drug-dependent antibody by ex-vivo antigen in urine

Ceftriaxone-induced hemolytic anemia in a child successfully managed with intravenous immunoglobulin

Ceftriaxone-induced immune hemolytic anemia as a life-threatening complication of antibiotic treatment of 'chronic Lyme disease

Cefuroxime-induced immune hemolysis

Celecoxib aggravates cardiac apoptosis in L-NAME-induced pressure overload model in rats: Immunohistochemical determination of cardiac caspase-3, Mcl-1, Bax and Bcl-2

Cell to organ: physiological, immunotoxic and oxidative stress responses of *Lamellidens marginalis* to inorganic arsenite

Cell-mediated immune suppression due to alfathesin in short-term anaesthesia

Cell-mediated immunosuppressive mechanisms induced by UV radiation

Cellphone electromagnetic radiation damages the testicular ultrastructure of male rats.

Cellular immune response and immunotoxicity induced by DON (deoxynivalenol) in piglets

Cellular immune responses and phagocytic activity of fishes exposed to pollution of volcano mud

Cellular immunity impaired among patients on left ventricular assist device for 6 months

Cellular immunity impaired among patients on left ventricular assist device for 6 months. Invited commentary

Cellular immunity in depressed, conduct disorder, and normal adolescents: role of adverse life events

Cellular immunoreaction in alcohol-induced liver damage

Cellular immunotoxicity of amyl nitrite

Cellular immunotoxicity of rAAV gene medicine and possible solutions

Central venous catheters and upper-extremity deep-vein thrombosis complicating immune heparin-induced thrombocytopenia

Cerebrospinal fluid inflammation induced by intravenous immunoglobulins

Challenges for nanoparticle characterization

Change in plasma membrane potential of rat thymocytes by tert-butylhydroquinone, a food additive: Possible risk on lymphocytes



Changed distribution and immune effects of nickel augment viral-induced inflammatory heart lesions in mice

Changes in ageing pace and major immune parameters among individuals with long exposure to hydrogen sulfide

Changes in biochemical and immunological parameters in humans exposed to organosilicon compounds

Changes in biologic and/or immunologic parameters induced by intratracheal instillation of pregnant mice with benzo(a)pyrene are potentially confounded by anesthesia

Changes in brain enkephalin immunostaining after acute carbon disulfide exposure in rats

Changes in DNA methylation and gene expression during 2,3,7,8-tetrachlorodibenzo-p-dioxin-induced suppression of the lipopolysaccharide-stimulated IgM response in splenocytes

Changes in enteroendocrine and immune cells following colitis induction by TNBS in rats

Changes in galanin immunoreactivity in rat micturition reflex pathways after cyclophosphamide-induced cystitis

Changes in HPBMC markers of immune function following controlled short-term inhalation exposures of humans to hardwood smoke

Changes in human peripheral blood mononuclear cell (HPBMC) populations and T-cell subsets associated with arsenic and polycyclic aromatic hydrocarbon exposures in a Bangladesh cohort

Changes in humoral immunity in sensitized guinea pigs exposed to trichloroethylene

Changes in immunological and hematological parameters of female residents exposed to volatile organic compounds in the city of Kaohsiung, Taiwan

Changes in selected aspects of immune function in the leopard frog, *Rana pipiens*, associated with exposure to cold

Changes in serum immunoglobulin levels in patients with chronic bronchitis exposed to nitrous oxide

Changes in Somatostatin-Like Immunoreactivity in the Sympathetic Neurons Projecting to the Prepyloric Area of the Porcine Stomach Induced by Selected Pathological Conditions

Changes in the toxicity and immunotoxicity of tetrachloromethane and carbophos under the action of 2,4,6-triphenyl-4H-selenopyran and their connection with cytochrome P-450 dependent monooxygenase system

Changes of CD8 T cells in dextran sulfate sodium-induced colitis mice pretreated with oral immune regulation

Changes of cerebral vasoactive intestinal polypeptide- and somatostatin-like immunoreactivity induced by noise and whole-body vibration in the rat

Changes of hematological and biochemical parameters revealed genotoxicity and immunotoxicity of neonicotinoids on Chinese rare minnows (*Gobiocypris rarus*)

Changes of the immunological and haematological parameters in rabbits after bendiocarbamate application

Characterisation of Immune and Neuroinflammatory Changes Associated with Chemotherapy-Induced Peripheral Neuropathy

Characterisation of immune-related gene expression in clam (*Venerupis philippinarum*) under exposure to di(2-ethylhexyl) phthalate

Characterisation of transcriptional responses to dioxins and dioxin-like contaminants in roach (*Rutilus rutilus*) using whole transcriptome analysis

Characterising vancomycin's immunotoxic profile using Swiss and CFW mice as an experimental model

Characteristics of the immunosuppression induced by cutaneous photodynamic therapy: persistence, antigen specificity and cell type involved

Characterization and protective effect of *Polygonatum sibiricum* polysaccharide against cyclophosphamide-induced immunosuppression in Balb/c mice

Characterization of antibody and selection of alternative drug therapy in hydrochlorothiazide-induced immune hemolytic anemia

Characterization of G12 sandwich ELISA, a next-generation immunoassay for gluten toxicity

Characterization of HCV-specific CD4+Th17 immunity in recurrent hepatitis C-induced liver allograft fibrosis

Characterization of human lymphoblastoid cell lines as a novel in vitro test system to predict the immunotoxicity of xenobiotics

Characterization of phenotypes of immune cells and cytokines associated with chronic exposure to *Premolis semirufa* caterpillar bristles extract

Characterization of protein phosphorylation by 2,3,7,8-tetrachlorodibenzo-p-dioxin in murine lymphocytes: indirect evidence for a role in the suppression of humoral immunity

Characterization of the humoral immune response and hepatotoxicity after multiple halothane exposures in guinea pigs

Characterization of the humoral immune response in heparin-induced thrombocytopenia

Characterization of the modes of action of deoxynivalenol (DON) in the human Jurkat T-cell line

Characterization of the role played by antigen challenge in the suppression of in vivo humoral immunity by 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD)

Characterization of the structural requirements for a carbohydrate based anticoagulant with a reduced risk of inducing the immunological type of heparin-associated thrombocytopenia

Characterization of vascular leak syndrome induced by the toxin component of *Pseudomonas* exotoxin-based immunotoxins and its potential inhibition with nonsteroidal anti-inflammatory drugs

Characterization, biomarkers, and reversibility of a monoclonal antibody-induced immune complex disease in cynomolgus monkeys (*Macaca fascicularis*)

Checkpoint inhibitors-induced hypophysitis

Chemical nature and immunotoxicological properties of arachidonic acid degradation products formed by exposure to ozone

Chemical xenogenization of murine lymphoma cells with triazene derivatives: immunotoxicological studies

Chemical-induced autoimmune reactions and Spanish toxic oil syndrome. Focus on hydantoins and related compounds

Chemical-induced inflammation and inflammatory diseases

Chemically induced immunotoxicity in a medium-term multiorgan bioassay for carcinogenesis with Wistar rats

Chemo- and radiotherapy and the risk of immunological suppression

Chemopreventive anti-cancer agent acyclic retinoid suppresses allogeneic immune responses in rats

Chemoprotective and immunomodulatory effect of *Acacia nilotica* during cyclophosphamide induced toxicity

Chemotactic activity of human polymorphonuclear leukocytes and industrial xenobiotics: a brief review

Chemotherapy-induced immunosuppression and reconstitution of immune function

Chemotherapy-induced peripheral neurotoxicity in immune-deficient mice: new useful ready-to-use animal models

Childhood adversity increases vulnerability for behavioral symptoms and immune dysregulation in women with breast cancer

Childhood exposure to ambient polycyclic aromatic hydrocarbons is linked to epigenetic modifications and impaired systemic immunity in T cells

Children's immunology, what can we learn from animal studies (3): Impaired mucosal immunity in the gut by 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD): a possible role for allergic sensitization

Children's white blood cell counts in relation to developmental exposures to methylmercury and persistent organic pollutants

Chimeric DNA vaccine reverses morphine-induced immunosuppression and tumorigenesis

Chitosan nanoparticles act as an adjuvant to promote both Th1 and Th2 immune responses induced by ovalbumin in mice

Chitosan Nanoparticles: Shedding Light on Immunotoxicity and Hemocompatibility

Chlamydia pneumoniae impairs the innate immune response in infected epithelial cells by targeting TRAF3

Chlorpyrifos exposure in common carp (*Cyprinus carpio* L.) leads to oxidative stress and immune responses

Chlorpyrifos Suppresses Neutrophil Extracellular Traps in Carp by Promoting Necroptosis and Inhibiting Respiratory Burst Caused by the PKC/MAPK Pathway

Chromium (VI)-induced immunotoxicity and intracellular accumulation in human primary dendritic cells

Chromium alters lipopolysaccharide-induced inflammatory responses both in vivo and in vitro

Chromophore for UV-induced immunosuppression: DNA

Chromophore for UV-induced immunosuppression: urocanic acid

Chromosomal aberrations in UVB-induced tumors of immunosuppressed mice

Chronic alcohol intoxication attenuates human immunodeficiency virus-1 glycoprotein 120-induced superoxide anion release by isolated Kupffer cells

Chronic allograft dysfunction: role of immunosuppressive treatment

Chronic aluminum intoxication in rat induced both serotonin changes in the dorsal raphe nucleus and alteration of glycoprotein secretion in the subcommissural organ: Immunohistochemical study

Chronic Binge Alcohol-Induced Dysregulation of Mitochondrial-Related Genes in Skeletal Muscle of Simian Immunodeficiency Virus-Infected Rhesus Macaques at End-Stage Disease

Chronic dietary toxicity study on 2,4-dichlorophenoxybutyric acid in the dog

Chronic effects of perfluorooctanesulfonate exposure on immunotoxicity in adult male C57BL/6 mice

Chronic endosulfan exposure impairs immune response rendering *Clarias gariepinus* susceptible to microbial infection

Chronic ethanol consumption impairs cellular immune responses against HCV NS5 protein due to dendritic cell dysfunction

Chronic exposure of tilapia (*Oreochromis niloticus*) to iron oxide nanoparticles: Effects of particle morphology on accumulation, elimination, hematology and immune responses

Chronic exposure to a trichloroethylene metabolite in autoimmune-prone MRL+/+ mice promotes immune modulation and alopecia

Chronic exposure to dim light at night suppresses immune responses in Siberian hamsters

Chronic exposure to low concentration of arsenic is immunotoxic to fish: role of head kidney macrophages as biomarkers of arsenic toxicity to *Clarias batrachus*

Chronic exposure to low concentrations of beryllium on the level of human serum immunoglobulins

Chronic Exposure to the Fusarium Mycotoxin Deoxynivalenol: Impact on Performance, Immune Organ, and Intestinal Integrity of Slow-Growing Chickens

Chronic exposure to trichloroethylene increases DNA methylation of the Ifng promoter in CD4+ T cells

Chronic exposure to water pollutant trichloroethylene increased epigenetic drift in CD4(+) T cells

Chronic fatigue syndrome and fibromyalgia following immunization with the hepatitis B vaccine: another angle of the 'autoimmune (auto-inflammatory) syndrome induced by adjuvants' (ASIA)

Chronic fetal exposure to Ureaplasma parvum suppresses innate immune responses in sheep

Chronic fluoride exposure exacerbates headkidney pathology and causes immune commotion in Clarias gariepinus

Chronic ingestion of deoxynivalenol and fumonisin, alone or in interaction, induces morphological and immunological changes in the intestine of piglets

Chronic lead (Pb) exposure results in diminished hemocyte count and increased susceptibility to bacterial infection in Drosophila melanogaster

Chronic lead exposure induces histopathological damage, microbiota dysbiosis and immune disorder in the cecum of female Japanese quails (Coturnix japonica)

Chronic life stress alters sympathetic, neuroendocrine, and immune responsivity to an acute psychological stressor in humans

Chronic orthostatic and antiorthostatic restraint induce neuroendocrine, immune and neurophysiological disorders in rats

Chronic pulmonary accumulation of iron oxide nanoparticles induced Th1-type immune response stimulating the function of antigen-presenting cells

Chronic restraint stress impairs T-cell immunity and promotes tumor progression in mice

Chronic Sleep Restriction Impairs the Antitumor Immune Response in Mice

Chronic social stress impairs virus specific adaptive immunity during acute Theiler's virus infection

Chronic Stress Contributes to Osteosarcopenic Adiposity via Inflammation and Immune Modulation: The Case for More Precise Nutritional Investigation

Chronic stress impairs the local immune response during cutaneous repair in gilthead sea bream (Sparus aurata, L

Chronic toxicity of azathioprine and the effect of this immunosuppressant on liver tumor induction by the carcinogen N-hydroxy-N-2-fluorenylacetamide

Chronic unpredictable mild stress impairs erythrocyte immune function and changes T-lymphocyte subsets in a rat model of stress-induced depression

Cigarette smoke condensate may disturb immune function with apoptotic cell death by impairing function of organelles in alveolar macrophages

Cigarette smoke extract suppresses the RIG-I-initiated innate immune response to influenza virus in the human lung

Cigarette smoke impairs NK cell-dependent tumor immune surveillance

Cigarette Smoke Induces Immune Responses to Vimentin in both, Arthritis-Susceptible and -Resistant Humanized Mice

Cigarette smoke suppresses type I interferon-mediated antiviral immunity in lung fibroblast and epithelial cells

Cigarette smoke-induced airway hyperresponsiveness is not dependent on elevated immunoglobulin and eosinophilic inflammation in a mouse model of allergic airway disease

Cigarette smoke-induced pulmonary emphysema in scid-mice. Is the acquired immune system required

Cigarette smoking impairs human pulmonary immunity to *Mycobacterium tuberculosis*

Cigarette Smoking Impairs the Bioenergetic Immune Response to *Mycobacterium tuberculosis* Infection

Cimetidine effects on the immunosuppression induced by burn injury

Cimetidine-induced Leydig cell apoptosis and reduced EG-VEGF (PK-1) immunoexpression in rats: Evidence for the testicular vasculature atrophy

Ciprofloxacin-induced immunoglobulin A disease

Circadian time-dependent antioxidant and inflammatory responses to acute cadmium exposure in the brain of zebrafish

Circadian variation of cytotoxicity and genotoxicity induced by an immunosuppressive agent "Mycophenolate Mofetil" in rats

Cis-bifenthrin causes immunotoxicity in murine macrophages

Cis-bifenthrin induces immunotoxicity in adolescent male C57BL/6 mice

Cis-urocanic acid increases immunotoxicity and lethality of dermally administered permethrin in C57BL/6N mice

cis-urocanic acid induces mast cell degranulation and release of preformed TNF- $\alpha$ : A possible mechanism linking UVB and cis-urocanic acid to immunosuppression of contact hypersensitivity

Citalopram enhances B cell numbers in a murine model of morphine-induced immunosuppression

Clarification of molecular targets of dioxin toxicity

CLARITY-BPA: Effects of chronic Bisphenol A exposure on the immune system: Part 1 - Quantification of the relative number and proportion of leukocyte populations in the spleen and thymus

Clcn5 knockout mice exhibit novel immunomodulatory effects and are more susceptible to dextran sulfate sodium-induced colitis

Clemastine causes immune suppression through inhibition of extracellular signal-regulated kinase-dependent proinflammatory cytokines

Clethodim exposure induces developmental immunotoxicity and neurobehavioral dysfunction in zebrafish embryos

Clinical and economic outcomes of rabbit antithymocyte globulin induction in adults who received kidney transplants from living unrelated donors and received cyclosporine-based immunosuppression

Clinical and immunological characteristics of a pediatric population with food protein-induced enterocolitis syndrome (FPIES) to fish

Clinical and immunological study of 7 patients with minocycline-induced autoimmune phenomena

Clinical aspects of endothelial dysfunction associated with human immunodeficiency virus infection and antiretroviral agents

Clinical case: idelalisib-induced immunoglobulin flare

Clinical immunological disorders in children from various observation cohorts exposed to radiation factor during various stages of oncogenesis

Clinical immunotoxicity of antirheumatic drugs

Clinical results of steroid-free induction immunosuppression after heart transplantation

Clinical study on changes of serum levels of immunoglobulin and cytokines in anaphylaxis patients induced by Qingkailing injection

Clinically relevant immunosuppressants influence UVB-induced tumor size through effects on inflammation and angiogenesis

Clinically Relevant Radiation Exposure Differentially Impacts Forms of Cell Death in Human Cells of the Innate and Adaptive Immune System

Clonal Epstein-Barr virus-positive mucocutaneous ulcer mimicking a mature B-cell lymphoma in a patient with mycophenolate-induced immune suppression

Clostridial gas gangrene: evidence that alpha and theta toxins differentially modulate the immune response and induce acute tissue necrosis

Clostridium perfringens alpha-Toxin Impairs Innate Immunity via Inhibition of Neutrophil Differentiation

Clozapine-induced agranulocytosis: Evidence for an immune-mediated mechanism from a patient-specific in-vitro approach

Clozapine-induced agranulocytosis: genetic risk factors and an immunologic explanatory model

Clozapine-induced parotitis: an immunological cause

CLP-induced impairment of innate immune function is caused by exposure to the cecal luminal contents and not the tissue trauma or tissue ischemia/necrosis components

CM-Dil labeled mesenchymal stem cells homed to thymus inducing immune recovery of mice after haploidentical bone marrow transplantation

CO<sub>2</sub>-induced ocean acidification impairs the immune function of the Pacific oyster against *Vibrio splendidus* challenge: An integrated study from a cellular and proteomic perspective

Coarse(PM<sub>2.5-10</sub>), fine(PM<sub>2.5</sub>), and ultrafine air pollution particles induce/increase immune costimulatory receptors on human blood-derived monocytes but not on alveolar macrophages

Cocaine immunotoxicity: abnormal cytokine production in Hispanic drug users

Cocaine/levamisole-induced systemic vasculitis with retiform purpura and pauci-immune glomerulonephritis

Cocaine-induced vasculitis: clinical and immunological spectrum

Co-cultivated damp building related microbes *Streptomyces californicus* and *Stachybotrys chartarum* induce immunotoxic and genotoxic responses via oxidative stress

Co-cultivation of *Streptomyces californicus* and *Stachybotrys chartarum* stimulates the production of cytostatic compound(s) with immunotoxic properties

Coexisting *Bacillus Calmette-Guerin*-Induced Lupus Vulgaris Involving the Vaccination Site and *Lichen Scrofulosorum* in an Immunocompetent Boy

Coexposure to mercury increases immunotoxicity of trichloroethylene

Cognitive and behavioral consequences of impaired immunoregulation in aging

Coinfection with an intestinal helminth impairs host innate immunity against *Salmonella enterica* serovar Typhimurium and exacerbates intestinal inflammation in mice

Cold exposure and absorption of colostral immunoglobulins by neonatal pigs

Cold-stress induced the modulation of catecholamines, cortisol, immunoglobulin M, and leukocyte phagocytosis in tilapia

Colocalization of taurine and glial fibrillary acidic protein immunoreactivity in mouse hippocampus induced by short-term ethanol exposure

Colonic adenocarcinomas rapidly induced by the combined treatment with 2-amino-1-methyl-6-phenylimidazo[4,5-b]pyridine and dextran sodium sulfate in male ICR mice possess beta-catenin gene mutations and increases immunoreactivity for beta-catenin, cyclooxygenase-2 and inducible nitric oxide synthase

Combating chronic renal allograft dysfunction : optimal immunosuppressive regimens

Combination exposure to zidovudine plus sulfamethoxazole-trimethoprim diminishes B-lymphocyte immune responses to *Pneumocystis murina* infection in healthy mice

Combination of chick embryo and nutrient mixture prevent D-galactose-induced cognitive deficits, immune impairment and oxidative stress in aging rat model



Combination of Selenomethionine and N-Acetylcysteine Alleviates the Joint Toxicities of Aflatoxin B1 and Ochratoxin A by ERK MAPK Signal Pathway in Porcine Alveolar Macrophages

Combined Immunosuppressive Therapy Induces Remission in Patients With Severe Type B Insulin Resistance: A Prospective Cohort Study

Combining Growth Factor and Bone Marrow Cell Therapy Induces Bleeding and Alters Immune Response After Stroke in Mice

Comfortable lifestyle-induced imbalance of neuro-endocrine-immunity network: a possible mechanism of vascular endothelial dysfunction

Comment on "5-azacytidine promotes an inhibitory T-cell phenotype and impairs immune mediated antileukemic activity

Comment on "Misoprostol impairs female reproductive tract innate immunity against *Clostridium sordellii*

Comments on: "a case of isoniazid-induced thrombocytopenia: recovery with immunoglobulin therapy

Commercial naphthenic acids and the organic fraction of oil sands process water downregulate pro-inflammatory gene expression and macrophage antimicrobial responses

Common classification schemes for PCB congeners and the gene expression of CYP17, CYP19, ESR1 and ESR2

Comorbidity is a better predictor of impaired immunity than chronological age in older adults

Comparative effect of ochratoxin A on inflammation and oxidative stress parameters in gut and kidney of piglets

Comparative effects of cadmium, copper, paraquat and benzo[a]pyrene on the actin cytoskeleton and production of reactive oxygen species (ROS) in mussel haemocytes

Comparative effects of immunotoxic chemicals on in vitro proliferative responses of human and rodent lymphocytes

Comparative efficacy of piperine, curcumin and picroliv against Cd immunotoxicity in mice

Comparative haemato-immunotoxic impacts of long-term exposure to tartrazine and chlorophyll in rats

Comparative immunosuppression of various glycol ethers orally administered to Fischer 344 rats

Comparative immunotoxicity assessment of N4-Trimethoxybenzoyl-5'-deoxy-5- fluorocytidine (Ro 09-1390) and 5'-deoxy-5-fluorouridine (5'-DFUR) in BDF1 mice

Comparative immunotoxicity of 2,2'-dichlorodiethyl sulfide and cyclophosphamide: evaluation of L1210 tumor cell resistance, cell-mediated immunity, and humoral immunity

Comparative immunotoxicity of free doxorubicin and doxorubicin encapsulated in cardiolipin liposomes

Comparative immunotoxicology of ultraviolet B exposure I. Effects of in vitro and in situ ultraviolet B exposure on the functional activity and morphology of Langerhans cells in the skin of different species

Comparative in vitro study of various titanium compounds on the immune system

Comparative investigation of behavioral, neurotoxicological, and immunotoxicological indices in detection of subacute combined exposure with methyl parathion and propoxur in rats

Comparative lung immunotoxicity of inhaled quartz and coal combustion fly ash

Comparative proteomics of inhaled silver nanoparticles in healthy and allergen provoked mice

Comparative studies on the antitumor and immunosuppressive effects of the new fluorouracil derivative N4-trimethoxybenzoyl-5'-deoxy-5-fluorocytidine and its parent drug 5'-deoxy-5-fluorouridine

Comparative study of respiratory tract immune toxicity induced by three sterilisation nanoparticles: silver, zinc oxide and titanium dioxide

Comparative toxicity of Corexit 9500, oil, and a Corexit/oil mixture on the eastern oyster, *Crassostrea virginica* (Gmelin)

Comparative toxicity of dithiocarbamates and butadiene metabolites in human lymphoid and bone marrow cells

Comparing immune activation (lipopolysaccharide) and toxin (lithium chloride)-induced gustatory conditioning: lipopolysaccharide produces conditioned taste avoidance but not aversion

Comparison between spousal donor transplantation treated with anti-thymocyte globulin induction therapy and, living related donor transplantation treated with standard immunosuppression

Comparison between tacrolimus and cyclosporine as immunosuppressive agents compatible with tolerance induction by CD4/CD8 blockade

Comparison of an IgG-Specific Enzyme-Linked Immunosorbent Assay Cutoff of 0.4 Versus 0.8 and 1.0 Optical Density Units for Heparin-Induced Thrombocytopenia

Comparison of antilymphocyte globulin and continuous i.v. cyclosporine A as induction immunosuppression for cadaver kidney transplants: a prospective randomized study

Comparison of cytotoxic and inflammatory responses of pristine and functionalized multi-walled carbon nanotubes in RAW 264.7 mouse macrophages

Comparison of detection sensitivity of immuno- and genotoxicological effects of subacute cypermethrin and permethrin exposure in rats

Comparison of experimental murine immunodeficiency models induced by cyclophosphamide and hydrocortisone

Comparison of five cyclosporine-prednisone regimens for induction of immunosuppression in cadaveric kidney recipients: a retrospective analysis of 245 cases

Comparison of immune status and 1,2-dimethylhydrazine induced tumorigenesis in brown--Norway and Fischer rats. Emphasis on splenic and colonic lymphocyte function

Comparison of immunotoxic effects induced by the extracts from methanol and gasoline engine exhausts in vitro

Comparison of immunotoxicity among tetrachloro-, pentachloro-, tetrabromo- and pentabromo-dibenzo-p-dioxins in mice

Comparison of inflammatory responses in mouse lungs exposed to atranones A and C from *Stachybotrys chartarum*

Comparison of OKT3 and antithymocyte globulin as induction immunosuppressive agents in renal transplantation

Comparison of polychlorinated biphenyl (PCB) induced effects on innate immune functions in harbour and grey seals

Comparison of sensitizing protocols for ultraviolet B-induced immunosuppression in C3H mice

Comparison of single bolus ATG and Basiliximab as induction therapy in presensitized renal allograft recipients receiving tacrolimus-based immunosuppressive regimen

Comparison of subchronic immunotoxicity of four different types of aluminum-based nanoparticles

Comparison of the Acute Immunotoxicity of Nonfractionated and Fractionated Oil Sands Process-Affected Water Using Mammalian Macrophages

Comparison of the D1-dopamine agonists SKF-38393 and A-68930 in neonatal 6-hydroxydopamine-lesioned rats: behavioral effects and induction of c-fos-like immunoreactivity

Comparison of the enteric mucosal immunomodulatory activity of combinations of *Coptis chinensis* Franch. Rhizomes and *Evodia rutaecarpa* (Juss.) Benth. Fruits in mice with dextran sulphate sodium-induced ulcerative colitis

Comparison of the immunotoxicity of propanil and its metabolite, 3,4-dichloroaniline, in C57Bl/6 mice

Comparisons of PAH-induced immunomodulation in three bivalve molluscs

Compensatory changes in cortical cholinergic innervation in the rat following an immunotoxic lesion

Complement activation-related pseudoallergy: a new class of drug-induced acute immune toxicity

Complement system and immunological mediators: Their involvements in the induced inflammatory process by *Androctonus australis* hector venom and its toxic components

Complement, circulating immunocomplexes and autoantibodies after ionizing radiation exposure

Complete Freund's adjuvant induces experimental autoimmune myocarditis by enhancing IL-6 production during initiation of the immune response

Complete reversal of FK 506 induced diabetes in a liver transplant recipient by change of immunosuppression to cyclosporine A

Complex Changes in the Innate and Adaptive Immunity Accompany Progressive Degeneration of the Nigrostriatal Pathway Induced by Intrastriatal Injection of 6-Hydroxydopamine in the Rat

Complications of using OKT3 for induction of immunosuppression in recipients of kidneys from nonheart beating donors

Comprehensive review on toxicity of persistent organic pollutants from petroleum refinery waste and their degradation by microorganisms

Computational tool for immunotoxic assessment of pyrethroids toward adaptive immune cell receptors

Condensed tannins decreased the growth performance and impaired intestinal immune function in on-growing grass carp (*Ctenopharyngodon idella*)

Conjunctiva-associated lymphoid tissue (CALT) reactions to antiglaucoma prostaglandins with or without BAK-preservative in rabbit acute toxicity study

Consecutive evaluation of graphene oxide and reduced graphene oxide nanoplatelets immunotoxicity on monocytes

Constant illumination reduces circulating melatonin and impairs immune function in the cricket *Teleogryllus commodus*

Constant light exposure impairs immune tolerance development in mice

Constant light suppresses production of Met-enkephalin-containing peptides in cultured splenic macrophages and impairs primary immune response in rats

Consumption of home-made spirits is one of the main source of exposure to higher alcohols and there may be a link to immunotoxicity

Contact dermatitis: from pathomechanisms to immunotoxicology

Contaminant exposure and effects in pinnipeds: implications for Steller sea lion declines in Alaska

Contaminant-induced immunosuppression and mass mortalities among harbor seals

Contaminant-induced immunotoxicity in harbour seals: wildlife at risk

Continuous and discontinuous cigarette smoke exposure differentially affects protective Th1 immunity against pulmonary tuberculosis

Contrasting effects of immunosuppression on herpes simplex virus type 1 (HSV 1) induced central nervous system (CNS) demyelination in mice

Contributions of nonhematopoietic cells and mediators to immune responses: implications for immunotoxicology

Conventional and novel "omics"-based approaches to the study of carbon nanotubes pulmonary toxicity

Coplanar and non-coplanar congener-specificity of PCB bioaccumulation and immunotoxicity in sea stars

Copper hypersensitivity: dermatologic aspects

Copper hypersensitivity: dermatologic aspects--an overview

Copper induced immunotoxicity promote differential apoptotic pathways in spleen and thymus

Copper nanoparticles induce early fibrotic changes in the liver via TGF-beta/Smad signaling and cause immunosuppressive effects in rats

Copper nanoparticles induced oxidation stress, cell apoptosis and immune response in the liver of juvenile Takifugu fasciatus

Copper-induced immunotoxicity involves cell cycle arrest and cell death in the liver

Copper-induced immunotoxicity involves cell cycle arrest and cell death in the spleen and thymus

Cord blood gene expression supports that prenatal exposure to perfluoroalkyl substances causes depressed immune functionality in early childhood

Coronary vasomotor dysfunction in the cardiac allograft: impact of different immunosuppressive regimens

Correlation between change of immunotoxicology indexes of cerum and the occurrence of anaphylaxis in patients induced by qingkailing injection: a clinical study

Correlation between HLA antigens and immune status in dust-induced lung diseases in workers of machine-building industry

Correlation between mast cell-mediated allergic inflammation and length of perfluorinated compounds

Correlation between maternal milk and infant serum levels of chlorinated pesticides (CP) and the impact of elevated CP on bleeding tendency and immune status in some infants in Egypt

Correlations between polychlorinated biphenyl immunotoxicity, the aromatic hydrocarbon locus, and liver microsomal enzyme induction in C57BL/6 and DBA/2 mice

Corrigendum to "Benzanthrone induced immunotoxicity via oxidative stress and inflammatory mediators in Balb/c mice" [Immunobiology 220, (March) (2015) 369-381

Corrigendum to "Immunotoxicity and allergenic potential induced by topical application of perfluorooctanoic acid (PFOA) in a murine model" [Food Chem. Toxicol. 136 (2020) 111114

Corrigendum to "Immunotoxicity and biodistribution analysis of arsenic trioxide in C57B1/6 mice following a 2-week inhalation exposure": [Toxicol. Appl. Pharmacol. 241 (2009) 253-259

Corrigendum to "Insights into the heavy metal-induced immunotoxic and genotoxic alterations as health indicators of Clarias gariepinus inhabiting a rivulet" [Ecotoxicol. Environ. Saf. 183 (2019

Corticosteroid-free immunosuppression with tacrolimus following induction with daclizumab: a large randomized clinical study

Corticosteroid-free immunosuppression with tacrolimus, mycophenolate mofetil, and daclizumab induction in renal transplantation

Corticosteroid-induced lymphopenia, immunosuppression, and body defense

Corticosteroids, immune suppression, and psychosis

Cortisol plays a role in the high environmental ammonia associated suppression of the immune response in zebrafish

Cost-effectiveness of postexposure prophylaxis after sexual or injection-drug exposure to human immunodeficiency virus

Could gosereline acetate induce autoimmune-like hepatitis

Coxsackievirus-induced myocarditis in mice: a model of autoimmune disease for studying immunotoxicity

CpG-based immunotherapy impairs antitumor activity of BRAF inhibitors in a B-cell-dependent manner

Creation of lung-targeted dexamethasone immunoliposome and its therapeutic effect on bleomycin-induced lung injury in rats

Critical periods for the teratogenicity of immune-suppressant Leflunomide in mice

Critical role for CCAAT/enhancer-binding protein beta in immune complex-induced acute lung injury

Crosstalk Among UV-Induced Inflammatory Mediators, DNA Damage and Epigenetic Regulators Facilitates Suppression of the Immune System

Crotonaldehyde induces apoptosis and immunosuppression in alveolar macrophages

Crude oil impairs immune function and increases susceptibility to pathogenic bacteria in southern flounder

Cryopreserved human whole blood: a human-based in vitro immunotoxicological system

CT20126, a novel immunosuppressant, prevents collagen-induced arthritis through the down-regulation of inflammatory gene expression by inhibiting NF-kappaB activation

Cumulative exposure to stimulants and immune function outcomes among HIV-positive and HIV-negative men in the Multicenter AIDS Cohort Study

Cumulative immunosuppressant exposure is associated with diversified cancer risk among 14 832 patients with systemic lupus erythematosus: a nested case-control study

Curcumin modulates the immune response associated with LPS-induced periodontal disease in rats

Curcumin protects against nicotine-induced stress during protein malnutrition in female rat through immunomodulation with cellular amelioration

Current developments in reproductive toxicity testing of pesticides

Current understanding of interactions between nanoparticles and the immune system

Cutaneous and Visceral Chronic Granulomatous Disease Triggered by a Rubella Virus Vaccine Strain in Children With Primary Immunodeficiencies

Cutaneous immunopathology of cyclosporin-A-induced autoimmunity in the rat

Cutaneous manifestations of *Paecilomyces lilacinus* infection induced by a contaminated skin lotion in patients who are severely immunosuppressed

CXCR3 deficiency exacerbates liver disease and abrogates tolerance in a mouse model of immune-mediated hepatitis

Cyclin D1/cdk4, estrogen receptors alpha and beta, in N-methyl-N'-nitro-N-nitrosoguanidine-induced rat gastric carcinogenesis: immunohistochemical study

Cyclooxygenase-2 mediates interleukin-6 upregulation by vomitoxin (deoxynivalenol) in vitro and in vivo

Cyclophosphamide and the Teratology Society: an awkward marriage

Cyclophosphamide effects on immune function of European starlings

Cyclophosphamide-induced neutrophilic disease in a patient with angioimmunoblastic lymphadenopathy and myelodysplastic syndrome

Cyclophosphamide-induced tolerance in kidney transplantation avoids long-term immunosuppressive therapy

Cyclophosphamide-like cellular immunostimulation induced by 6-thioguanine (Lanvis

Cyclosporin A, a powerful immunosuppressant in vivo and in vitro in the dog, fails to induce tolerance

Cyclosporin A: a new immune suppressive agent

Cyclosporine A-induced hyperactivity in rats: is it mediated by immunosuppression, neurotrophism, or both

Cyclosporine and rapamycin act in a synergistic and dose-dependent manner in a model of immunosuppressant-induced kidney damage

Cyclosporine effect on immunoregulatory cells in kidney transplant recipients: suppression of gamma-interferon and interleukin production

Cyclosporine induced autoimmunity in newborns prevented by early immunization

Cyclosporine, an immunosuppressant, attenuates phorbol-induced lung injury in rats

Cyclosporine-induced autoimmunity and immune hyperreactivity

CYP1A2 is not required for 2,3,7,8-tetrachlorodibenzo-p-dioxin-induced immunosuppression

Cypermethrin Formulation (Ustad-10 EC) Induces Genotoxicity via Apoptosis, Affects Nutritional Physiology, and Modulates Immune Response in Silkworm *Philosamia ricini* (Lepidoptera: Saturniidae)

Cytobiochemical characteristics of the function of pulmonary alveolar macrophages during development of immune reactions after exposure to chemical air pollutants

Cytochrome P450 1B1 is required for 7,12-dimethylbenz(a)-anthracene (DMBA) induced spleen cell immunotoxicity

Cytochrome P4501B1 mediates induction of bone marrow cytotoxicity and preleukemia cells in mice treated with 7,12-dimethylbenz[a]anthracene

Cytogenetic and immunological changes after dermal exposure to polycyclic aromatic hydrocarbons and UV radiation

Cytokines and cytokine receptors in mouth mucosa of immune suppressed patients

Cytokines as biomarkers of nanoparticle immunotoxicity

Cytokines related to three major types of cell-mediated immunity in short- and long-term exposures to lead compounds

Cytomegalovirus Immunity After Alemtuzumab Induction in Desensitized Kidney Transplant Patients

Cytomegalovirus infection impairs immune responses and accentuates T-cell pool changes observed in mice with aging

Cytomegalovirus-induced thrombocytopenia in an immunocompetent adult effectively treated with intravenous immunoglobulin: a case report and review

Cytostatic therapy reduces the immune defense. Children treated for leukemia have impaired immunity against measles and rubella

Cytotoxic action of triphenyltin on mouse thymocytes: a flow-cytometric study using fluorescent dyes for membrane potential and intracellular  $Ca^{2+}$

Cytotoxic and immunotoxic effects of Fusarium mycotoxins using a rapid colorimetric bioassay

Cytotoxic effects and changes in cytokine gene expression induced by microcystin-containing extract in fish immune cells--an in vitro and in vivo study

Cytotoxic effects of exposure to the human immunodeficiency virus type 1 protein Tat in the hippocampus are enhanced by prior ethanol treatment

Cytotoxicity and immunotoxicity of cyclopiazonic acid on human cells

Cytotoxicity of lambda-cyhalothrin on the macrophage cell line RAW 264.7

Cytotoxicity of nitroaromatic explosives and their biodegradation products in mice splenocytes: implications for their immunotoxicity

Cytotoxicity, inhibition of DNA and protein syntheses and oxidative damage in cultured cells exposed to zearalenone

Daclizumab induction and maintenance steroid-free immunosuppression with mycophenolate mofetil and tacrolimus to prevent acute rejection of hepatic allografts

Daclizumab induction, tacrolimus, mycophenolate mofetil and steroids as an immunosuppression regimen for primary kidney transplant recipients

Daily exposure to dust alters innate immunity



Danish bacille Calmette-Guerin vaccine-induced disease in human immunodeficiency virus-infected children

Data sieving analysis as a novel method to assess immunotoxic exposure to dioxins retrospectively

DDT inhibits the functional activation of murine macrophages and decreases resistance to infection by *Mycobacterium microti*

Decreased blood level of beta 2-microglobulin in the employees of a factory which produced polychlorinated biphenyls

Deep sequencing of blood and gut T-cell receptor beta-chains reveals gluten-induced immune signatures in celiac disease

Deficiency of dietary niacin impaired gill immunity and antioxidant capacity, and changes its tight junction proteins via regulating NF-kappaB, TOR, Nrf2 and MLCK signaling pathways in young grass carp (*Ctenopharyngodon idella*)

Deficiency of dietary niacin impaired intestinal mucosal immune function via regulating intestinal NF-kappaB, Nrf2 and MLCK signaling pathways in young grass carp (*Ctenopharyngodon idella*)

Deficient Th1-type immune responses via impaired CD28 signaling in ultraviolet B-induced systemic immunosuppression and the restorative effect of IL-12

Deficit in selective and divided attention associated with cholinergic basal forebrain immunotoxic lesion produced by 192-saporin; motoric/sensory deficit associated with Purkinje cell immunotoxic lesion produced by OX7-saporin

Defining pollen exposure times for clinical trials of allergen immunotherapy for pollen-induced rhinoconjunctivitis - an EAACI position paper

Deflazacort induced stronger immunosuppression than expected

Degradation of superparamagnetic iron oxide nanoparticle-induced ferritin by lysosomal cathepsins and related immune response

DEHP induces immunosuppression through disturbing inflammatory factors and CYPs system homeostasis in common carp neutrophils

Delayed complications of sulfur mustard poisoning in the skin and the immune system of Iranian veterans 16-20 years after exposure

Delayed developmental immunotoxicity of prenatal benzodiazepines

Delayed developmental neuro- and immunotoxicity of benzodiazepines

Delayed discovery, dissemination, and decisions on intervention in environmental health: a case study on immunotoxicity of perfluorinated alkylate substances

Delayed immune-mediated thrombocytopenia after re-exposure to abciximab therapy

Delayed immunologic thrombocytopenia induced by abciximab

Delayed redistribution of CD27, CD40 and CD80 positive B cells and the impaired in vitro immunoglobulin production in patients with non-Hodgkin lymphoma after rituximab treatment as an adjuvant to autologous stem cell transplantation

Deleterious effects in reproduction and developmental immunity elicited by pulmonary iron oxide nanoparticles

Delta-9-tetrahydrocannabinol augments murine retroviral induced immunosuppression and infection

Delta-9-tetrahydrocannabinol enhances breast cancer growth and metastasis by suppression of the antitumor immune response

Deltamethrin Increases *Candida albicans* infection susceptibility in mice

Deltamethrin-Induced Immunotoxicity and its Protection by Quercetin: An Experimental Study

Deltamethrin-induced oxidative stress and mitochondrial caspase-dependent signaling pathways in murine splenocytes

Dendrobium sonia polysaccharide regulates immunity and restores the dysbiosis of the gut microbiota of the cyclophosphamide-induced immunosuppressed mice

Density of CD1a-labeled Langerhans' cells in normal human gingiva and in nifedipine- and immunosuppressive medication-induced gingival overgrowth

Dental Fluorosis and Catalase Immunoreactivity of the Brain Tissues in Rats Exposed to High Fluoride Pre- and Postnatally

Deoxynivalenol (DON) naturally contaminated feed impairs the immune response induced by porcine reproductive and respiratory syndrome virus (PRRSV) live attenuated vaccine

Deoxynivalenol impairs the immune functions of neutrophils

Deoxynivalenol suppresses circulating and splenic leukocyte subpopulations in BALB/c mice: dose response, time course and sex differences

Deoxynivalenol, gut microbiota and immunotoxicity: A potential approach

Deoxynivalenol: mechanisms of action, human exposure, and toxicological relevance

Deoxynivalenol-induced IgA production and IgA nephropathy-aberrant mucosal immune response with systemic repercussions

Depressed immunity and impaired proliferation of hematopoietic progenitor cells in patients with complete spinal cord injury

Depression in caregivers of demented patients is associated with altered immunity: impaired proliferative capacity, increased CD8+, and a decline in lymphocytes with surface signal transduction molecules (CD38+) and a cytotoxicity marker (CD56+ CD8

Depressive symptoms following interferon-alpha therapy: mediated by immune-induced reductions in brain-derived neurotrophic factor

Depressive symptoms lead to impaired cellular immune response

Dermal application of jet fuel suppresses secondary immune reactions

Dermal application of JP-8 jet fuel induces immune suppression

Dermal exposure to jet fuel suppresses delayed-type hypersensitivity: a critical role for aromatic hydrocarbons

Dermatomyositis-like syndrome revealing statin-induced necrotizing autoimmune myopathy with anti-HMGCR antibodies

Detailed immunohistologic evaluation of a methotrexate-induced nodule

Detection of 1,2,4-benzenetriol induced aneuploidy and microtubule disruption by fluorescence in situ hybridization and immunocytochemistry

Detection of benzo[a]pyrene-induced immunotoxicity in orange spotted grouper (*Epinephelus coioides*)

Detection of immunotoxic effects of estrogenic and androgenic endocrine disrupting compounds using splenic immune cells of the female three-spined stickleback, *Gasterosteus aculeatus* (L)

Detection of immunotoxicity of benzo[a]pyrene in a subacute toxicity study after oral exposure in rats

Detection of immunotoxicity using T-cell based cytokine reporter cell lines ("Cell Chip")

Detection of the effects of repeated dose combined propoxur and heavy metal exposure by measurement of certain toxicological, haematological and immune function parameters in rats

Detection of the mechanism of immunotoxicity of cyclosporine A in murine in vitro and in vivo models

Determination of enantiomeric fractions of cypermethrin and cis-bifenthrin in Chinese teas by GC/ECD

Determination of N7- and O6-methylguanine in rat liver DNA after oral exposure to hydrazine by use of immunochemical and electrochemical detection methods

Determination of the immunotoxic potential of heavy metals on the functional activity of bottlenose dolphin leukocytes in vitro

Determination of the immunotoxic potential of pesticides on functional activity of sheep leukocytes in vitro

Determining the relationship between nanoparticle characteristics and immunotoxicity: key challenges and approaches

Detrimental interactions of neonicotinoid pesticide exposure and bumblebee immunity

Development of a lymphocytic lymphoma during immunosuppressive therapy with azathioprine for systemic lupus erythematosus with renal involvement induced by phenylbutazone

Development of a squamous cell carcinoma mouse model for immunotoxicity testing

Development of non-antibiotic macrolide that corrects inflammation-driven immune dysfunction in models of inflammatory bowel diseases and arthritis

Development of the "Cell Chip": a new in vitro alternative technique for immunotoxicity testing

Development of the murine and human immune system: differential effects of immunotoxicants depend on time of exposure

Development of vigabatrin-induced lesions in the rat brain studied by magnetic resonance imaging, histology, and immunocytochemistry

Developmental and reproductive toxicity testing of vaccines

Developmental atrazine exposure suppresses immune function in male, but not female Sprague-Dawley rats

Developmental exposure to 2,3,7,8 tetrachlorodibenzo-p-dioxin attenuates later-life Notch1-mediated T cell development and leukemogenesis

Developmental exposure to 2,3,7,8 tetrachlorodibenzo-p-dioxin attenuates capacity of hematopoietic stem cells to undergo lymphocyte differentiation

Developmental exposure to bisphenol A modulates innate but not adaptive immune responses to influenza A virus infection

Developmental exposure to lead causes persistent immunotoxicity in Fischer 344 rats

Developmental exposure to the potent aryl hydrocarbon receptor agonist 2,3,7,8-tetrachlorodibenzo-p-dioxin Impairs the cell-mediated immune response to infection with influenza a virus, but enhances elements of innate immunity

Developmental immunotoxicity (DIT): assays for evaluating effects of exogenous agents on development of the immune system

Developmental immunotoxicity in male rats after juvenile exposure to ethanol

Developmental immunotoxicity in male rats after juvenile exposure to di-n-octyltin dichloride (DOTC

Developmental immunotoxicity investigations in the SD rat following pre- and post-natal exposure to cyclosporin

Developmental immunotoxicity of atrazine in rodents

Developmental immunotoxicity of chemicals in rodents and its possible regulatory impact

Developmental immunotoxicity of cocaine and ethanol in postnatal Lewis rats

Developmental immunotoxicity of cyclosporin-A in rats: age-associated differential effects

Developmental immunotoxicity of dexamethasone: comparison of fetal versus adult exposures

Developmental immunotoxicity of di-n-octyltin dichloride (DOTC) in an extended one-generation reproductive toxicity study

Developmental immunotoxicity of ethanol in an extended one-generation reproductive toxicity study

Developmental immunotoxicity of lead in the rat: influence of maternal diet

Developmental immunotoxicity of lead: impact on thymic function

Developmental immunotoxicity of methylmercury: the relative sensitivity of developmental and immune parameters

Developmental immunotoxicity of trichloroethylene (TCE): studies in B6C3F1 mice

Developmental immunotoxicity testing of 4-methyl anisole

Developmental Immunotoxicity, Perinatal Programming, and Noncommunicable Diseases: Focus on Human Studies

Developmental Immunotoxicology of Di-(2-Ethylhexyl)phthalate (DEHP): Age-Based Assessment in the Female Rat

Developmental immunotoxicology of lead

Developmental immunotoxicology: emerging issues

Developmental neurotoxicity and immunotoxicity induced by graphene oxide in zebrafish embryos

Developmental stage-specific changes in immunological biomarkers in male C3H/HeN mice after early life toluene exposure

Developmental toxicity of cadmium in mice. II. Immunotoxic effects

Developmental toxicity of cesium in the mouse

Developmental toxicity of N,N1-bis(dichloroacetyl)-1,8-octamethylene diamine: effects of in utero exposure on the postnatal murine immune system

Developmental toxicity, oxidative stress and immunotoxicity induced by three strobilurins (pyraclostrobin, trifloxystrobin and picoxystrobin) in zebrafish embryos

Developmental windows of differential lead-induced immunotoxicity in chickens

Developmental, neuro and immunotoxic effects of perinatal diazepam treatment in rats

Developments of immunotoxicology methods in the rat and applications to the study of environmental pollutants

Dextran sulfate sodium-induced colitis occurs in severe combined immunodeficient mice

Di-(2-ethylhexyl) phthalate adjuvantly induces imbalanced humoral immunity in ovalbumin-sensitized BALB/c mice ascribing to T follicular helper cells hyperfunction

Di(2-ethylhexyl) phthalate inhibits B cell proliferation and reduces the abundance of IgM-secreting cells in cultured immune tissues of the rainbow trout

Diagnosis and treatment of heparin-induced thrombocytopenia (HIT) based on its atypical immunological features

Diagnosis and Treatment of Ventilator-Associated Infection: Review of the Critical Illness Stress-Induced Immune Suppression Prevention Trial Data

Diagnostic accuracy of IgG-specific versus polyspecific enzyme-linked immunoassays in heparin-induced thrombocytopenia: a systematic review and meta-analysis

Diagnostic accuracy of rapid immunoassays for heparin-induced thrombocytopenia. A systematic review and meta-analysis

Diagnostic Performance of a New Rapid Lateral Flow Immunoassay in Patients Suspected of Heparin-Induced Thrombocytopenia and Its Clinical Consequences

Diagnostic value of immunoassays for heparin-induced thrombocytopenia: a systematic review and meta-analysis

Diagnostic value of two immunoassays for detecting heparin/PF4 complex antibodies in heparin-induced thrombocytopenia

Diazepam leads to enhanced severity of orthopoxvirus infection and immune suppression

Diazinon immunotoxicity in mice: modulation of cytokines level and their gene expression

Dibenzo[def,p]chrysene (DBC) suppresses antibody formation in spleen cells following oral exposures of mice

Dibromoacetic acid induced Cl.Ly1+2/-9 T-cell apoptosis and activation of MAPKs signaling cascades

Dibromoacetic Acid Induces Thymocyte Apoptosis by Blocking Cell Cycle Progression, Increasing Intracellular Calcium, and the Fas/FasL Pathway in Vitro

Dibutyl phthalate-mediated oxidative stress induces splenic injury in mice and the attenuating effects of vitamin E and curcumin

Dibutyltin and tributyltin compounds induce thymus atrophy in rats due to a selective action on thymic lymphoblasts

Dibutyltin-induced alterations of interleukin 1beta secretion from human immune cells

Dichloroacetate improves immune dysfunction caused by tumor-secreted lactic acid and increases antitumor immunoreactivity

Dicyclohexylphthalate causes hyperactivity in the rat concomitantly with impairment of tyrosine hydroxylase immunoreactivity

Dietary biotin deficiency decreased growth performance and impaired the immune function of the head kidney, spleen and skin in on-growing grass carp (*Ctenopharyngodon idella*)

Dietary *Chlorella vulgaris* Ameliorates Altered Immunomodulatory Functions in Cyclophosphamide-Induced Immunosuppressive Mice

Dietary chromium methionine supplementation could alleviate immunosuppressive effects of heat stress in broiler chicks

Dietary eicosapentaenoic acid prevents systemic immunosuppression in mice induced by UVB radiation

Dietary exposure to low pesticide doses causes long-term immunosuppression in the leopard frog (*Rana pipiens*)

Dietary fish oil suppresses experimental immunoglobulin A nephropathy in mice

Dietary grape-seed proanthocyanidin inhibition of ultraviolet B-induced immune suppression is associated with induction of IL-12

Dietary iron deficiency impaired intestinal immune function of on-growing grass carp under the infection of *Aeromonas hydrophila*: Regulation of NF- $\kappa$ B and TOR signaling

Dietary Iron Deficiency Impaired Peripheral Immunity but Did Not Alter Brain Microglia in PRRSV-Infected Neonatal Piglets

Dietary L-carnitine and vitamin-E; a strategy to combat ochratoxin-A induced immunosuppression

Dietary low or excess levels of lipids reduced growth performance, and impaired immune function and structure of head kidney, spleen and skin in young grass carp (*Ctenopharyngodon idella*) under the infection of *Aeromonas hydrophila*

Dietary lutein reduces ultraviolet radiation-induced inflammation and immunosuppression

Dietary mustard seeds (*Sinapis alba* Linn) suppress 1,2-dimethylhydrazine-induced immuno-imbalance and colonic carcinogenesis in rats

Dietary protein restriction impairs growth, immunity, and disease resistance in southern leopard frog tadpoles

Dietary pyridoxine deficiency reduced growth performance and impaired intestinal immune function associated with TOR and NF- $\kappa$ B signalling of young grass carp (*Ctenopharyngodon idella*)

Dietary resveratrol supplementation inhibits heat stress-induced high-activated innate immunity and inflammatory response in spleen of yellow-feather broilers

Dietary selenium protect against redox-mediated immune suppression induced by methylmercury exposure

Dietary selenium supplementation alleviates immune toxicity in the hearts of chickens with lead-added drinking water

Dietary zinc deficiency impairs humoral and cellular immune responses to BCG and ESAT-6/CFP-10 vaccination in offspring and adult rats

Differences in patient and transplant professional perceptions of immunosuppression-induced cosmetic side effects

Differences in the incidence and clinical evolution of early neurotoxicity after liver transplantation based on tacrolimus formulation used in the immunosuppressive induction protocol

Differential changes in rat cholinergic parameters subsequent to immunotoxic lesion of the basal forebrain nuclei

Differential effects of 2,3,7,8-tetrachlorodibenzo-p-dioxin, bis(tri-n-butyltin) oxide and cyclosporine on thymus histophysiology

Differential effects of coadministration of aminoacetonitrile on immunosuppression and hepatotoxicity produced by dimethylnitrosamine

Differential effects of selective immunotoxic lesions of medial septal cholinergic cells on spatial working and reference memory

Differential immunogenic and neurogenic inflammatory responses in an allergic mouse model exposed to low levels of formaldehyde

Differential immunotoxic effects of ethanol on murine EL-4 lymphoma and normal lymphocytes is mediated through increased ROS production and activation of p38MAPK

Differential immunotoxic effects of inorganic and organic mercury species in vitro

Differential immunotoxic effects of the environmental chemical benzo[a]pyrene in young and aged mice

Differential immunotoxicities of poly(ethylene glycol)- vs. poly(carboxybetaine)-coated nanoparticles

Differential immunotoxicity induced by two different windows of developmental trichloroethylene exposure

Differential immunotoxicity of histone deacetylase inhibitors on malignant and naive hepatocytes

Differential induction of rat neuronal excitotoxic cell death by human immunodeficiency virus type 1 clade B and C tat proteins

Differential inhibition of IL-1 alpha and TNF-alpha generation by ammonium metavanadate in murine macrophages

Differential modification of inflammatory enzymes in J774A.1 macrophages by ochratoxin A alone or in combination with lipopolysaccharide

Differential mRNA expression of neuroimmune markers in the hippocampus of infant mice following toluene exposure during brain developmental period

Differential protease, innate immunity, and NF-kappaB induction profiles during lung inflammation induced by subchronic cigarette smoke exposure in mice

Differential protection by two sunscreens from UV radiation-induced immunosuppression

Differential role of serotonin projections from the dorsal and median raphe nuclei in phencyclidine-induced hyperlocomotion and fos-like immunoreactivity in rats

Differential sensitivities of bone marrow, spleen and thymus to genotoxicity induced by environmentally relevant concentrations of arsenite

Differential sensitivity to cadmium of immunomarkers measured in hemocyte subpopulations of zebra mussel *Dreissena polymorpha*



Differential, histochemical and immunohistochemical changes in rat hepatocytes after isoflurane or sevoflurane exposure

Digoxin immune fab protects endothelial cells from ouabain-induced barrier injury

Dim light at night interferes with the development of the short-day phenotype and impairs cell-mediated immunity in Siberian hamsters (*Phodopus sungorus*)

Diminished expression of CYP1A1 in urethane-induced lung tumors in strain A/J mice: analysis by in situ hybridization and immunohistochemical methods

Diminished neutrophil extracellular trap (NET) formation is a novel innate immune deficiency induced by acute ethanol exposure in polymicrobial sepsis, which can be rescued by CXCL1

Diminished protection? Early childhood PCB exposure and reduced immune response to vaccinations

Dioxin strongly immunotoxic

Dioxin toxicity, aryl hydrocarbon receptor signaling, and apoptosis-persistent pollutants affect programmed cell death

Dioxin-like (DL-) polychlorinated biphenyls induced immunotoxicity through apoptosis in mice splenocytes via the AhR mediated mitochondria dependent signaling pathways

Dioxins and the immune system: mechanisms of interference. A meeting report

Diphenyl diselenide dietary supplementation protects against methylmercury-chloride-induced immunotoxicity in the head kidney and spleen of grass carp (*Ctenopharyngodon idella*) via regulation of purinergic signaling and the NLRP3 inflammasome

Direct effects of carbon nanotubes on dendritic cells induce immune suppression upon pulmonary exposure

Direct suppression of cultured spleen cell responses by chlordane and the basis for differential effects on in vivo and in vitro immunocompetence

Direct thrombin inhibitors in the treatment of immune-mediated heparin-induced thrombocytopenia

Direct-Acting Antiviral Therapy Restores Immune Tolerance to Patients With Hepatitis C Virus-Induced Cryoglobulinemia Vasculitis

Disaster: stress, immunologic function, and health behavior

Disorders in the murine chromosome apparatus induced by immunization with a complex of antiviral vaccines

Disorders of humoral immunity in workers exposed to organic solvents

Disruption of gut immune system caused by damage of intestinal stem cells and their niche in graft-versus-host disease after allogeneic hematopoietic stem cell transplantation

Disruption of human plasma cell differentiation by an environmental polycyclic aromatic hydrocarbon: a mechanistic immunotoxicological study

Disruption of the interaction of T cells with antigen-presenting cells by the active leflunomide metabolite teriflunomide: involvement of impaired integrin activation and immunologic synapse formation

Disseminated superficial porokeratosis secondary to immunosuppression induced by etanercept for extensive psoriasis

Distribution and behavior of the Ah receptor in murine T lymphocytes

Distribution and immunotoxicity by intravenous injection of iron nanoparticles in a murine model

Distribution, elimination, and toxicity of silver nanoparticles and silver ions in rats after 28-day oral exposure

Divergent immunological responses following glutaraldehyde exposure

Diverse Roads to Relapse: A Discriminative Cue Signaling Cocaine Availability Is More Effective in Renewing Cocaine Seeking in Goal Trackers Than Sign Trackers and Depends on Basal Forebrain Cholinergic Activity

DNA damage and adduct formation in immune organs of developing chicks by polycyclic aromatic hydrocarbons

DNA double strand breaks in epidermal cells cause immune suppression in vivo and cytokine production in vitro

DNA repair and cyclin D1 polymorphisms and styrene-induced genotoxicity and immunotoxicity

Do cryopreserved mesenchymal stromal cells display impaired immunomodulatory and therapeutic properties

Do tobacco stimulate the production of nitric oxide by up regulation of inducible nitric oxide synthesis in cancer: Immunohistochemical determination of inducible nitric oxide synthesis in oral squamous cell carcinoma--a comparative study in tobacco habitués and non-habitués

Docosahexaenoic acid attenuates mycotoxin-induced immunoglobulin a nephropathy, interleukin-6 transcription, and mitogen-activated protein kinase phosphorylation in mice

Does allergen-specific immunotherapy induce contact allergy to aluminium

Does developmental exposure to perfluorooctanoic acid (PFOA) induce immunopathologies commonly observed in neurodevelopmental disorders

Does exposure to immunosuppressive therapy increase the 10 year malignancy and mortality risks in rheumatoid arthritis? A matched cohort study

Does exposure to UV radiation induce a shift to a Th-2-like immune reaction

Does high organochlorine (OC) exposure impair the resistance to infection in polar bears (*Ursus maritimus*)? Part I: Effect of OCs on the humoral immunity

Does major surgery induce immune suppression and increase the risk of postoperative infection

Does negative heparin-platelet factor 4 enzyme-linked immunosorbent assay effectively exclude heparin-induced thrombocytopenia

DON shares a similar mode of action as the ribotoxic stress inducer anisomycin while TBTO shares ER stress patterns with the ER stress inducer thapsigargin based on comparative gene expression profiling in Jurkat T cells

Dose response for UV-induced immune suppression in people of color: differences based on erythral reactivity rather than skin pigmentation

Dose-response studies in murine mercury-induced autoimmunity and immune-complex disease

D-penicillamine: mechanism of cellular action and induced autoimmune diseases

D-penicillamine-induced pancreatic islet autoantibody production is independent of the immunogenetic background: a lesson from patients with Wilson's disease

Drinking water disinfection byproducts: review and approach to toxicity evaluation

Drug fever induced by piperacillin/tazobactam in an elderly patient with underlying human immunodeficiency virus (HIV) infection

Drug induced aseptic meningitis caused by intravenous immunoglobulin therapy

Drug induced immune haemolytic anaemia in the Berlin Case-Control Surveillance Study

Drug interaction effects on antitumour drugs (XV): Disulfiram as protective agent against cyclophosphamide-induced urotoxicity without compromising antitumour activity in mice

Drug-dependent clearance of human platelets in the NOD/scid mouse by antibodies from patients with drug-induced immune thrombocytopenia

Drug-induced cutaneous lupus erythematosus after immunoglobulin treatment in chronic inflammatory demyelinating polyneuropathy: a case series

Drug-induced immune hemolytic anemia associated with anti-vancomycin complicated by a paraben antibody

Drug-induced immune thrombocytopaenia: results from the Berlin Case-Control Surveillance Study

Drug-induced immune thrombocytopenia due to moxifloxacin

Drug-induced immune thrombocytopenia: a descriptive survey in the French Pharmacovigilance database

Drug-induced immunotoxicity

Drug-induced linear immunoglobulin A bullous dermatosis mimicking Stevens-Johnson syndrome: a case report

Drug-induced linear immunoglobulin A bullous disease that clinically mimics toxic epidermal necrolysis

Drugs of abuse and dysfunction of neuroendocrine and immune systems: the importance of animal research

Dualistic immunomodulation of sub-chronic microcystin-LR exposure on the innate-immune defense system in male zebrafish

Dysbiosis-induced IL-33 contributes to impaired antiviral immunity in the genital mucosa

Dysregulation of immune responses in an allergic mouse model following low-level toluene exposure

Dysregulation of the immune system caused by silica and asbestos

Early acute necrosis and delayed apoptosis induced by methyl mercury in murine peritoneal neutrophils

Early and late effects of the immunosuppressants rapamycin and mycophenolate mofetil on UV carcinogenesis

Early and Long-Term Impaired T Lymphocyte Immune Reconstitution after Cord Blood Transplantation with Antithymocyte Globulin

Early changes of lymphocyte RNA and serum immunoglobulins following chronic exposure to benzene

Early experience with FK 506 induction immunosuppression--suggestion for using oral FK 506

Early exposure to thirdhand cigarette smoke affects body mass and the development of immunity in mice

Early Immunosuppressive Exposure of Enteric-Coated-Mycophenolate Sodium Plus Tacrolimus Associated with Acute Rejection in Expanded Criteria Donor Kidney Transplantation

Early initiating and promoting effects in 2-AAF-induced rat liver carcinogenesis: an immunohistochemical study

Early innate immune events induced by prolonged cold ischemia exacerbate allograft vasculopathy

Early life allergen and air pollutant exposures alter longitudinal blood immune profiles in infant rhesus monkeys

Early life environment and developmental immunotoxicity in inflammatory dysfunction and disease

Early lifestage exposure and potential developmental susceptibility to tetrachloroethylene

Early onset of immunological heparin-induced thrombocytopenia in acute myocardial infarction

Early onset of virus infection and up-regulation of cytokines in mice treated with cadmium and manganese

Early onset steroid induced posterior subcapsular cataract in a patient with common variable immunodeficiency: case reports and review of literature

Early phosphoproteomic changes in the mouse spleen during deoxynivalenol-induced ribotoxic stress

Early Postnatal Secondhand Smoke Exposure Disrupts Bacterial Clearance and Abolishes Immune Responses in Muco-Obstructive Lung Disease

Early suppression of B cell immune responses by low doses of chloroquine and pyrimethamine: implications for studying immunity in malaria

Early testicular effects in rats perinatally exposed to DEHP in combination with DEHA--apoptosis assessment and immunohistochemical studies

Early use of steroids affects immune cells and impairs immunotherapy efficacy

Early weaning stress in pigs impairs innate mucosal immune responses to enterotoxigenic *E. coli* challenge and exacerbates intestinal injury and clinical disease

Early withdrawal of calcineurin inhibitors and rescue immunosuppression with sirolimus-based therapy in renal transplant recipients with moderate to severe renal dysfunction

Early-life environment, developmental immunotoxicology, and the risk of pediatric allergic disease including asthma

Early-life exposure to combustion-derived particulate matter causes pulmonary immunosuppression

Early-life Exposure to Widespread Environmental Toxicants and Health Risk: A Focus on the Immune and Respiratory Systems

Early-life long-term exposure to ZnO nanoparticles suppresses innate immunity regulated by SKN-1/Nrf and the p38 MAPK signaling pathway in *Caenorhabditis elegans*

*Echinacea purpurea* Protects Against Restraint Stress-Induced Immunosuppression in BALB/c Mice

Echinoderm reactive oxygen species (ROS) production measured by peroxidase, luminol-enhanced chemiluminescence (PLCL) as an immunotoxicological tool

E-cigarette use results in suppression of immune and inflammatory-response genes in nasal epithelial cells similar to cigarette smoke

Ecological impacts of the deepwater horizon oil spill: implications for immunotoxicity

Ecotoxicity of CdTe quantum dots to freshwater mussels: impacts on immune system, oxidative stress and genotoxicity

Ecotoxicological biomarkers as investigating tools to evaluate the impact of acrylamide on *Theba pisana* snails

Ecotoxicological impact of engineered nanomaterials in bivalve molluscs: An overview

Ecstasy" induced immunosuppression and herpes zoster ophthalmicus

Efavirenz induces depressive-like behaviour, increased stress response and changes in the immune response in rats

Effect of 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD) on influenza virus host resistance in mice

Effect of 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD) on maternal immune response during pregnancy

Effect of acetochlor on transcription of genes associated with oxidative stress, apoptosis, immunotoxicity and endocrine disruption in the early life stage of zebrafish

Effect of Acupuncture and Moxibustion Intervention on Contents of Serum Interleukin-7 and Interleukin-18 in Cyclophosphamide-induced Immune Suppression in Tumor-bearing Mice

Effect of anesthesia on the immune system: suppression of the immunogenic capacity of macrophages and of lymphocyte transformation

Effect of atrazine on immunocompetence of red-eared slider turtle(*Trachemys scripta*)

Effect of *Bacillus cereus* against cadmium induced hematological disturbances and immunosuppression in *Carassius auratus gibelio*

Effect of benzo[a]pyrene on the immune status of mice with anxious-depressive syndrome

Effect of beryllium exposure on PPD cellular immunity in guinea pigs

Effect of biphenyl dimethyl dicarboxylate on the cellular and nonspecific immunotoxicity by ethanol in mice

Effect of Bisphenol A on non-specific immunodefenses against non-pathogenic *Escherichia coli*

Effect of cadmium on the beta-adrenoreceptor of splenic cell of rats and its immunotoxicity

Effect of cannabinoid ingestion (in the form of bhang) on the immune system of high school and university students

Effect of Chinese medicine therapy for strengthening-Pi and nourishing-Shen in preventing lamivudine induced YMDD mutation and its immunologic mechanism

Effect of chronic developmental lead exposure on cell-mediated immune functions

Effect of chronic toluene exposure on tyrosine hydroxylase-positive nerve elements in the rat forebrain: an immunohistochemical study combined with semiquantitative morphometric analysis

Effect of Cyclooxygenase(COX)-1 and COX-2 inhibition on furosemide-induced renal responses and isoform immunolocalization in the healthy cat kidney

Effect of CYP2E1 induction by ethanol on the immunotoxicity and genotoxicity of extended low-level benzene exposure

Effect of diazinon, an organophosphate pesticide, on signal transduction and death induction in mononuclear cells of Nile tilapia fish (*Oreochromis niloticus*)

Effect of dietary glutamate on chemotherapy-induced immunosuppression

Effect of Different Selenium Supplementation Levels on Oxidative Stress, Cytokines, and Immunotoxicity in Chicken Thymus

Effect of DTC on humoral response of SRBC-immunized mice exposed to restraint stress. Comparison with calf thymus extract

Effect of electromagnetic radiation on T-lymphocyte subpopulations and immunoglobulin level in human blood serum after occupational exposure.

Effect of electromagnetic waves from mobile phone on immune status of male rats: possible protective role of vitamin D.

Effect of ethanol and 4-methylpyrazole on the immunotoxicity of ethylene glycol

Effect of ethanol on methanol immunotoxicity

Effect of experimental desynchronosis on immunotoxicity of benz(a)pyrene in (CBAXC57Bl)F1 mice

Effect of exposure time, particle size and uptake pathways in immune cell lysosomal cytotoxicity of mussels exposed to silver nanoparticles

Effect of exposure to nitrogen dioxide on alveolar macrophage-mediated immunosuppressive activity in rats

Effect of extremely high frequency electromagnetic radiation of low intensity on parameters of humoral immunity in healthy mice.

Effect of fluoride exposure on different immune parameters in humans

Effect of heavy oil exposure on antibacterial activity and expression of immune-related genes in Japanese flounder *Paralichthys olivaceus*

Effect of IL-12 on immune suppression and suppressor cell induction by ultraviolet radiation

Effect of immunocyte therapy on benzene-induced bone marrow haemopoietic dysfunction

Effect of immunomodulators pyrimethamine and cimetidine on immunosuppression induced by sulfur mustard in mice

Effect of immunosuppression on neonatally diethylstilbestrol-induced genital tract lesion and tumor development in female mice

Effect of in vitro exposure to zinc on immunological parameters of haemocytes from the marine gastropod *Haliotis tuberculata*

Effect of inhaled industrial chemicals on systemic and local immune response

Effect of inhibitors of oxygen radical and nitric oxide formation on UV radiation-induced erythema, immunosuppression and carcinogenesis

Effect of ketamine on cocaine-induced immunotoxicity in rats

Effect of lactational exposure to tributyltin chloride on innate immunodefenses in the F1 generation in mice

Effect of lead and arsenic on murine macrophage response

Effect of lead exposure on serum immunoglobulins and reactive nitrogen and oxygen intermediate

Effect of lead exposure on the immune function of lymphocytes and erythrocytes in preschool children

Effect of lemon verbena powder and vitamin C on performance and immunity of heat-stressed broilers

Effect of licorice roots on carrageenan-induced decrease in immune complexes clearance in mice

Effect of lindane exposure on rainbow trout (*Oncorhynchus mykiss*) immunity. IV. Prevention of nonspecific and specific immunosuppression by dietary vitamin C (ascorbate-2-polyphosphate

Effect of low copper exposure on the antioxidant system and some immune parameters

Effect of low intensity and very high frequency electromagnetic radiation on occupationally exposed personnel.

Effect of low-intensity microwave of on mitomycin C-induced genotoxicity in vitro.

Effect of maternal exposure to ozone on reproductive outcome and immune, inflammatory, and allergic responses in the offspring

Effect of microwave radiation on cellular immunity indices in conditions of chronic exposure.

Effect of moisture-damage intervention on the immunotoxic potential and microbial content of airborne particles and on occupants' upper airway inflammatory responses

Effect of N-acetylcysteine on some aspects of cyclophosphamide-induced toxicity and immunosuppression

Effect of nickel exposure on peripheral tissues: role of oxidative stress in toxicity and possible protection by ascorbic acid

Effect of orally administered zinc oxide nanoparticles on albino rat thymus and spleen

Effect of ozone exposure on alveolar macrophage-mediated immunosuppressive activity in rats

Effect of passive smoking on frequency of respiratory illnesses and serum immunoglobulin-E (IgE) and interleukin-4 (IL-4) concentrations in exposed children

Effect of polyethylene glycol surface charge functionalization of SWCNT on the in vitro and in vivo nanotoxicity and biodistribution monitored noninvasively using MRI

Effect of polysaccharide from cultured *Cordyceps sinensis* on immune function and anti-oxidation activity of mice exposed to <sup>60</sup>Co

Effect of prenatal exposure to combined immunosuppressive agrochemicals in a mouse model of allergic airway inflammation

Effect of prenatal exposure to ethanol on glutamate and GABA immunoreactivity in macaque somatosensory and motor cortices: critical timing of exposure

Effect of progestins on immunity: medroxyprogesterone but not norethisterone or levonorgestrel suppresses the function of T cells and pDCs

Effect of PYCNOGENOL on the toxicity of heart, bone marrow and immune organs as induced by antitumor drugs



Effect of quercetin on impaired immune function in mice exposed to irradiation

Effect of selenomethionine supplementation in food on the excretion and toxicity of arsenic exposure in female mice

Effect of some mycotoxins on superoxide anion production of isolated human neutrophils and in whole blood

Effect of subchronic 2,3,7,8-tetrachlorodibenzo-p-dioxin exposure on immune system and target gene responses in mice: calculation of benchmark doses for CYP1A1 and CYP1A2 related enzyme activities

Effect of sulfur dioxide inhalation on cytokine levels in lungs and serum of mice

Effect of synthetic thymic hormones on the cocaine-induced inhibition of the primary immune response in mice

Effect of tetrachloromethane on the immune system

Effect of the Aqueous Extract of *Lantana grisebachii* Stuck Against Bioaccumulated Arsenic-Induced Oxidative and Lipid Dysfunction in Rat Splenocytes

Effect of the *Fusarium* toxin deoxynivalenol (DON) on IgA, IgM and IgG concentrations and proliferation of porcine blood lymphocytes

Effect of the H-2 and Igh complexes on the susceptibility to ultraviolet B-induced immunosuppression in murine contact sensitivity and contact photosensitivity

Effect of the immobilized microcystin-LR-degrading enzyme MlrA on nodularin degradation and its immunotoxicity study

Effect of the immunomodulator AS101 on chemotherapy-induced multilineage myelosuppression, thrombocytopenia, and anemia in mice

Effect of the protein corona on nanoparticles for modulating cytotoxicity and immunotoxicity

Effect of the thiol group on experimental gold-induced autoimmunity

Effect of thyroidectomy and thyroxine on 2,3,7,8-tetrachlorodibenzo-p-dioxin-induced immunotoxicity

Effect of topiramate on the kainate-induced status epilepticus, lipid peroxidation and immunoreactivity of rats

Effect of tributyltin on antioxidant ability and immune responses of zebrafish (*Danio rerio*)

Effect of ultraviolet radiation on the immune system and the effect of exogenous photochemoprotective agents on the ultraviolet radiation induced immunosuppression

Effect of wide-band modulated electromagnetic fields on the workers of high-frequency telephone exchanges.

Effect on the immune system of mice exposed chronically to 50 Hz amplitude-modulated 2.45 GHz microwaves

Effective treatment of SIVcpz-induced immunodeficiency in a captive western chimpanzee

Effectiveness of concomitant immunosuppressive therapy in suppressing the formation of antibodies to infliximab in Crohn's disease

Effects of (Anti) Androgenic Endocrine Disruptors (DEHP and Butachlor) on Immunoglobulin M (IgM) and Leukocytes Counts of Male Rainbow Trout (*Oncorhynchus mykiss*)

Effects of 2000  $\mu\text{W}/\text{cm}^2$ ; electromagnetic radiation on expression of immunoreactive protein and mRNA of NMDA receptor 2A subunit in rats hippocampus.

Effects of 4-nonylphenol and/or diisononylphthalate on THP-1 cells: impact of endocrine disruptors on human immune system parameters

Effects of a new immunotherapeutic agent (CG5601) on endotoxin-induced uveitis

Effects of acid water exposure on plasma cortisol, ion balance, and immune functions in the "cobalt" variant of rainbow trout

Effects of Acute Low-Dose Exposure to the Chlorinated Flame Retardant Dechlorane 602 and Th1 and Th2 Immune Responses in Adult Male Mice

Effects of adolescent methamphetamine and nicotine exposure on behavioral performance and MAP-2 immunoreactivity in the nucleus accumbens of adolescent mice

Effects of agricultural pesticides on the immune system of *Rana pipiens* and on its resistance to parasitic infection

Effects of aluminum exposure on the allergic responses and humoral immune function in rats

Effects of ammonia-N exposure on the concentrations of neurotransmitters, hemocyte intracellular signaling pathways and immune responses in white shrimp *Litopenaeus vannamei*

Effects of androgen disruption by DDE on the development and functioning of the immune system in Japanese quail

Effects of antiretroviral therapy on immunity in patients infected with HIV

Effects of arsenic on porcine dendritic cells in vitro

Effects of atrazine on the proliferation and cytotoxicity of murine lymphocytes with the use of carboxyfluorescein succinimidyl ester-based flow cytometric approaches

Effects of benzene on splenic, thymic, and femoral lymphocytes in mice

Effects of brevetoxin exposure on the immune system of loggerhead sea turtles

Effects of Caramel Colour III on the number of blood lymphocytes: a human study on Caramel Colour III immunotoxicity and a comparison of the results with data from rat studies

Effects of ceftaroline on the innate immune and on the inflammatory responses of bronchial epithelial cells exposed to cigarette smoke

Effects of chromium on lymphocyte subsets and immunoglobulins from normal population and exposed workers

Effects of chronic exposure to the aqueous fraction of produced water on growth, detoxification and immune factors of Atlantic cod

Effects of cigarette smoke on immune response: chronic exposure to cigarette smoke impairs antigen-mediated signaling in T cells and depletes IP3-sensitive Ca(2+) stores

Effects of cis-nonachlor, trans-nonachlor and chlordane on the immune system of Sprague-Dawley rats following a 28-day oral (gavage) treatment

Effects of combined exposure to formaldehyde and benzene on immune cells in the blood and spleen in Balb/c mice

Effects of Corticosterone on Immune Functions of Cultured Rat Splenic Lymphocytes Exposed to Aluminum Trichloride

Effects of cyclophosphamide on the phenotypes and functions of THP-1 cells

Effects of decabrominated diphenyl ether (DBDE) on developmental immunotoxicity in offspring mice

Effects of diazinon on the lymphocytic cholinergic system of Nile tilapia fish (*Oreochromis niloticus*)

Effects of dietary exposure to chlorpyrifos on immune cell populations and inflammatory responses in mice with dextran sulfate sodium-induced colitis

Effects of dietary toxic cyanobacteria and ammonia exposure on immune function of blunt snout bream (*Megalabrama amblycephala*)

Effects of electromagnetic radiation on health and immune function of operators.

Effects of epidural anaesthesia on surgical stress-induced immunosuppression during upper abdominal surgery

Effects of exposure to multiple heavy metals on biochemical and histopathological alterations in common carp, *Cyprinus carpio* L

Effects of exposure to oil spills on human health: Updated review

Effects of fetal exposure to urban particulate matter on the immune system of male mouse offspring

Effects of four Indian medicinal herbs on Isoniazid-, Rifampicin- and Pyrazinamide-induced hepatic injury and immunosuppression in guinea pigs

Effects of fusariotoxin co-exposure on THP-1 human immune cells

Effects of glucan on immunosuppressive actions of mercury

Effects of GSM 1800 MHz radiofrequency electromagnetic fields on DNA damage in Chinese hamster lung cells.

Effects of haemofiltration and mannitol treatment on cardiopulmonary-bypass induced immunosuppression

Effects of heavy alcohol consumption on the cardiovascular system may be mediated in part by the influence of alcohol-induced depression on the immune system

Effects of high-dose intravenous immunoglobulin on lipopolysaccharide-induced acute lung injury

Effects of hydrazyl group containing drugs on leucocyte functions: an immunoregulatory model for the hydralazine-induced lupus-like syndrome

Effects of immunoglobulin on the neuronal expression of IL-1 $\beta$  and IL-1 $\alpha$  and the neuronal death at hippocampus in rats with convulsion induced by pentylenetetrazol

Effects of immunological and hematological parameter in mice exposed to mixture of volatile organic compounds

Effects of immunological challenge induced by lipopolysaccharide on skeletal muscle fiber type conversion of piglets

Effects of immunosuppressants on concanavalin A-induced interleukin-2 mRNA expression in mouse liver

Effects of immunosuppressive chemicals on lymphoid development in foetal thymus organ cultures

Effects of in vitro nickel exposure on the macrophage-mediated immune functions of rainbow trout (*Oncorhynchus mykiss*)

Effects of in vivo chronic exposure to pendimethalin/Prowl 400 on sanitary status and the immune system in rainbow trout (*Oncorhynchus mykiss*)

Effects of levamisole, DTC and low-dose mechlorethamine on humoral response of SRBC-immunized rabbits exposed to cold stress

Effects of lindane exposure on rainbow trout (*Oncorhynchus mykiss*) immunity. II. In vitro restoration of antibody-secreting cells and lymphocyte proliferation activity by nitrogranulogen after in vivo immunosuppression due to lindane

Effects of lindane exposure on rainbow trout (*Oncorhynchus mykiss*) immunity. III. Effect on nonspecific immunity and B lymphocyte functions

Effects of low concentrations of cadmium on immunoglobulin E production by human B lymphocytes in vitro

Effects of manganese and hypoxia on coelomocyte renewal in the echinoderm, *Asterias rubens* (L)

Effects of maternal silver acetate exposure on immune biomarkers in rodent model

Effects of melatonin and green-wavelength LED light on the physiological stress and immunity of goldfish, *Carassius auratus*, exposed to high water temperature

Effects of metal compounds with distinct physicochemical properties on iron homeostasis and antibacterial activity in the lungs: chromium and vanadium

Effects of methyl substitutions on benz[a]anthracene derivatives-induced immunosuppression

Effects of Methylmercury on Harbour Seal Peripheral Blood Leucocytes In Vitro Studied by Electron Microscopy

Effects of microbial cocultivation on inflammatory and cytotoxic potential of spores

Effects of Moderate Prenatal Alcohol Exposure during Early Gestation in Rats on Inflammation across the Maternal-Fetal-Immune Interface and Later-Life Immune Function in the Offspring

Effects of nanosized titanium dioxide on innate immune system of fathead minnow (*Pimephales promelas* Rafinesque, 1820

Effects of nanostructured zeolite and aflatoxin B1 in growth performance, immune parameters and pathological conditions of rainbow trout *Oncorhynchus mykiss*

Effects of negative life events on immune suppression in children and youth infected with human immunodeficiency virus type 1

Effects of neurotoxic insecticides on heat-shock proteins and cytokine transcription in Chinook salmon (*Oncorhynchus tshawytscha*

Effects of norfloxacin nicotinate on the early life stage of zebrafish (*Danio rerio*): Developmental toxicity, oxidative stress and immunotoxicity

Effects of ochratoxin A on mouse oocyte maturation and fertilization, and apoptosis during fetal development

Effects of ochratoxin A on the mouse immune system after subchronic exposure

Effects of ochratoxin A on the rat immune system after perinatal exposure

Effects of organochlorine contaminants on loggerhead sea turtle immunity: comparison of a correlative field study and in vitro exposure experiments

Effects of ozone on the defense to a respiratory *Listeria monocytogenes* infection in the rat. Suppression of macrophage function and cellular immunity and aggravation of histopathology in lung and liver during infection

Effects of PCB126 and PCB153 on telomerase activity and telomere length in undifferentiated and differentiated HL-60 cells

Effects of perinatal exposure to low doses of PCB 153 and PCB 126 on lymphocyte proliferation and hematology in goat kids

Effects of persistent organic pollutants on the developing respiratory and immune systems: a systematic review

Effects of pesticide exposure on serum immunoglobulin and complement levels

Effects of PM2.5 exposure on the Notch signaling pathway and immune imbalance in chronic obstructive pulmonary disease

Effects of polychlorinated biphenyls with Ah receptor affinity on lymphoid development in the thymus and the bursa of Fabricius of chick embryos in ovo and in mouse thymus anlagen in vitro

Effects of prenatal exposure to cigarette smoke on offspring tumor susceptibility and associated immune mechanisms

Effects of prenatal exposure to perfluoroalkyl acids on prevalence of allergic diseases among 4-year-old children

Effects of prenatal immune activation on amphetamine-induced addictive behaviors: Contributions from animal models

Effects of pretreatment with SDZ MRL 953, a novel immunostimulatory lipid A analog, on endotoxin-induced acute lung injury in guinea pigs

Effects of prior oral exposure to combinations of environmental immunosuppressive agents on ovalbumin allergen-induced allergic airway inflammation in Balb/c mice

Effects of pyrene exposure on immune response and oxidative stress in the pearl oyster, *Pinctada martensii*

Effects of red wine on ochratoxin A toxicity in intestinal Caco-2/TC7 cells

Effects of selected pharmaceutical products on phagocytic activity in *Elliptio complanata* mussels

Effects of selenium exposure on the hematology, innate immunity, and metabolic rate of yellow-bellied sliders (*Trachemys scripta scripta*)

Effects of short-term oral combined exposure to environmental immunotoxic chemicals in mice

Effects of sodium selenite on aflatoxin B1-induced decrease of ileal IgA+ cell numbers and immunoglobulin contents in broilers

Effects of spinetoram on the developmental toxicity and immunotoxicity of zebrafish

Effects of subchronic exposure of rats to 2-methoxyethanol or 2-butoxyethanol: thymic atrophy and immunotoxicity

Effects of subchronic exposure to Caramel Colour III on the immune system in mice

Effects of Th1 and Th2 cells balance in pulmonary injury induced by nano titanium dioxide

Effects of the endoparasitoid *Cotesia chilonis* (Hymenoptera: Braconidae) parasitism, venom, and calyx fluid on cellular and humoral immunity of its host *Chilo suppressalis* (Lepidoptera: Crambidae) larvae

Effects of the ethanol extract of *Cichorium intybus* on the immunotoxicity by ethanol in mice

Effects of transplacental and trans-breast milk exposure to the organophosphate compound chlorpyrifos on the developing immune system of mice

Effects of trophic exposure to diclofenac and dexamethasone on hematological parameters and immune response in freshwater fish

Effects of various LED light spectra on antioxidant and immune response in juvenile rock bream, *Oplegnathus fasciatus* exposed to bisphenol A

Effects on the amount of total hemolytic complement levels(CH50) and immunoglobulin in serum induced by the implantation of biomaterials into rats

Efficacy and safety of immunotherapy with interferon-gamma in the management of chronic sulfur mustard-induced cutaneous complications: comparison with topical betamethasone 1

Efficacy and safety of the SQ house dust mite sublingual immunotherapy tablet in Japanese adults and adolescents with house dust mite-induced allergic rhinitis

Efficacy of dendritic cell-cytokine-induced killer immunotherapy plus intensity-modulated radiation therapy in treating elderly patients with esophageal carcinoma

Efficacy of histopathology in detecting petrochemical-induced toxicity in wild cotton rats (*Sigmodon hispidus*)

Efficacy of micronized titanium dioxide-containing compounds in protection against UVB-induced immunosuppression in humans in vivo

Efficacy of *Nigella sativa* in alleviating benzo[a]pyrene-induced immunotoxicity in broilers

Efficacy of polyvinylpolypyrrolidone in reducing the immunotoxicity of aflatoxin in growing broilers

Efficacy of prophylactic application of immunoglobulins in radiation induced mucositis

Efficacy of steroid pulse therapy in combination with mizoribine following tonsillectomy for immunoglobulin A nephropathy in renally impaired patients

Efficacy of various Marek's disease vaccines protocols for prevention of Marek's disease virus-induced immunosuppression

Efficient in vitro model for immunotoxicologic assessment of mammary silicone implants

Efflux Transporters Regulate Arsenite-Induced Genotoxicity in Double Negative and Double Positive T Cells

EGFR-mediated Akt and MAPKs signal pathways play a crucial role in patulin-induced cell proliferation in primary murine keratinocytes via modulation of Cyclin D1 and COX-2 expression

Egg Yolk Immunoglobulin Supplementation Prevents Rat Liver from Aflatoxin B1-Induced Oxidative Damage and Genotoxicity

Egyptian sweet marjoram leaves protect against genotoxicity, immunosuppression and other complications induced by cyclophosphamide in albino rats

Electroacupuncture regulates the DREAM/NF-kappaB signalling pathway and ameliorates cyclophosphamide-induced immunosuppression in mice

Electromagnetic fields may act via calcineurin inhibition to suppress immunity, thereby increasing risk for opportunistic infection: Conceivable mechanisms of action

Electromagnetic fields may act via calcineurin inhibition to suppress immunity, thereby increasing risk for opportunistic infection: Conceivable mechanisms of action.

Elemental diet-induced bacterial translocation associated with systemic and intestinal immune suppression

Elemental diet-induced immune suppression is caused by both bacterial and dietary factors

Elemental imbalance elicited by arsenic and copper exposures leads to oxidative stress and immunotoxicity in chicken gizzard, activating the protective effects of heat shock proteins

Elevated lead levels and adverse effects on natural killer cells in children from an electronic waste recycling area

Elevated serum alpha-fetoprotein and impaired immune response in malnutrition

Elicitation of the immune response to p-phenylenediamine in allergic patients: the role of dose and exposure time

Elucidating mechanisms of immunotoxicity by benzotriazole ultraviolet stabilizers in zebrafish (*Danio rerio*): Implication of the AHR-IL17/IL22 immune pathway

Embolizing pulmonary aspergillosis, mycobacterial & aspergillous splenic abscess and cytomegalovirus co-infection following steroid induced immunosuppression: a case report

Embryo implantation dysfunction via the decreased uterine dendritic cells's non-immune function after exposed to carbon disulfide

Embryonic exposure to aflatoxin-B1: mutagenicity and influence on development and immunity

Embryonic exposure to butachlor in zebrafish (*Danio rerio*): endocrine disruption, developmental toxicity and immunotoxicity

Embryonic exposure to cadmium (II) and chromium (VI) induce behavioral alterations, oxidative stress and immunotoxicity in zebrafish (*Danio rerio*)

Embryonic exposure to carbendazim induces the transcription of genes related to apoptosis, immunotoxicity and endocrine disruption in zebrafish (*Danio rerio*)

Embryonic exposure to cis-bifenthrin enantioselectively induces the transcription of genes related to oxidative stress, apoptosis and immunotoxicity in zebrafish (*Danio rerio*)

Embryonic exposure to cypermethrin induces apoptosis and immunotoxicity in zebrafish (*Danio rerio*)

Embryonic exposure to PFOS induces immunosuppression in the fish larvae of marine medaka

Embryotoxicity Caused by DON-Induced Oxidative Stress Mediated by Nrf2/HO-1 Pathway

Emerging Role of Immunosuppression in Diseases Induced by Micro- and Nano-Particles: Time to Revisit the Exclusive Inflammatory Scenario



Enantioselective developmental toxicity and immunotoxicity of pyraclofos toward zebrafish (*Danio rerio*)

Enantioselectivity in the immunotoxicity of the insecticide acetofenatate in an in vitro model

Endocrine disrupting activities and immunomodulatory effects in lymphoblastoid cell lines of diclofenac, 4-hydroxydiclofenac and paracetamol

Endocrine disrupting compounds: can they target the immune system of fish

Endocrine disruption in adolescence: immunologic, hematologic, and bone effects in monkeys

Endocrine-disrupting chemicals and skin manifestations

Endocrine-disrupting chemicals impair the innate immune prophenoloxidase system in the intertidal mud crab, *Macrophthalmus japonicus*

Endocrinopathies induced by immune-checkpoint inhibitors in advanced non-small cell lung cancer

Endoplasmic Reticulum Stress and Apoptosis Triggered by Sub-Chronic Lead Exposure in Mice Spleen: a Histopathological Study

Endosulfan splenic pathology and amelioration by vitamin C in New Zealand rabbit

Endotoxin Exposure during Sensitization to *Blomia tropicalis* Allergens Shifts TH2 Immunity Towards a TH17-Mediated Airway Neutrophilic Inflammation: Role of TLR4 and TLR2

Energy restriction restores the impaired immune response in overweight (cafeteria) rats

Enhanced activation-induced cell death as a mechanism of 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD)-induced immunotoxicity in peripheral T cells

Enhanced dendritic cell survival attenuates lipopolysaccharide-induced immunosuppression and increases resistance to lethal endotoxic shock

Enhanced expression of single immunoglobulin IL-1 receptor-related molecule ameliorates LPS-induced acute lung injury in mice

Enhanced immunological and detoxification responses in Pacific oysters, *Crassostrea gigas*, exposed to chemically dispersed oil

Enhanced pulmonary arteriopathy in simian immunodeficiency virus-infected macaques exposed to morphine

Enhanced suppression of humoral immunity in DBA/2 mice following subchronic exposure to 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD)

Enhancement of immediate allergic reactions by trichloroethylene ingestion via drinking water in mice

Enhancement of immune tolerance via induction of NK1.1 positive liver-associated-lymphocytes under immunosuppressive conditions

Enhancement of lipopolysaccharide-induced nitric oxide and interleukin-6 production by PEGylated gold nanoparticles in RAW264.7 cells

Enhancement of metastasis of prostate adenocarcinoma cells by immune-suppressive cyclosporine A

Enniatin B1 exerts embryotoxic effects on mouse blastocysts and induces oxidative stress and immunotoxicity during embryo development

Enteric reovirus infection as a probe to study immunotoxicity of the gastrointestinal tract

Enterovirus-induced myocarditis: hemodynamic deterioration with immunosuppressive therapy and successful application of interferon-alpha

Environmental exposure to mercury and its toxicopathologic implications for public health

Environmental health in the Baltic region--toxic metals

Environmentally relevant doses of tetrabromobisphenol A (TBBPA) cause immunotoxicity in murine macrophages

Environmentally Relevant Level of Aflatoxin B1 Dysregulates Human Dendritic Cells Through Signaling on Key Toll-Like Receptors

Environmentally relevant level of aflatoxin B1 elicits toxic pro-inflammatory response in murine CNS-derived cells

Epicutaneous allergen-specific immunotherapy ameliorates grass pollen-induced rhinoconjunctivitis: A double-blind, placebo-controlled dose escalation study

Epicutaneous immunotherapy (EPIT) blocks the allergic esophago-gastro-enteropathy induced by sustained oral exposure to peanuts in sensitized mice

Epigallocatechin gallate, a potential immunomodulatory agent of tea components, diminishes cigarette smoke condensate-induced suppression of anti-Legionella pneumophila activity and cytokine responses of alveolar macrophages

Epigenetic targets for reversing immune defects caused by alcohol exposure

Epigenetic underpinnings of developmental immunotoxicity and autoimmune disease

Epigenotoxicity of environmental pollutants evaluated by a combination of DNA methylation inhibition and capillary electrophoresis-laser-induced fluorescence immunoassay

Epstein-Barr virus strain heterogeneity impairs human T-cell immunity

Epstein-Barr virus-encoded dUTPase modulates immune function and induces sickness behavior in mice

Erionite induces production of autoantibodies and IL-17 in C57BL/6 mice

ERK and p38 MAPK signaling pathways are involved in ochratoxin A-induced G2 phase arrest in human gastric epithelium cells

Eruptive disseminated porokeratosis associated with corticosteroid-induced immunosuppression

Erythrodermic pustular psoriasis triggered by intravesical bacillus Calmette-Guerin immunotherapy

Establishment of pantropic spotted dolphin (*Stenella attenuata*) fibroblast cell line and potential influence of polybrominated diphenyl ethers (PBDEs) on cytokines response

Estimated exposures to perfluorinated compounds in infancy predict attenuated vaccine antibody concentrations at age 5-years

Estradiol impairs the Th17 immune response against *Candida albicans*

Estrogenic xenobiotics affect the intracellular activation signal in mitogen-induced human peripheral blood lymphocytes: immunotoxicological impact

Ethanol enhances immunosuppression induced by cocaine

Ethanol impairs intestinal barrier defense by modulation of immunoglobulin A transport

Ethanol impairs mucosal immunity against *Streptococcus pneumoniae* infection by disrupting interleukin 17 gene expression

Ethanol inhibits lipid raft-mediated TCR signaling and IL-2 expression: potential mechanism of alcohol-induced immune suppression

Ethanol-induced increase of agouti-related protein (AgRP) immunoreactivity in the arcuate nucleus of the hypothalamus of C57BL/6J, but not 129/SvJ, inbred mice

Ethanol-induced suppression of cell-mediated immunity in the mouse

Ethylene dibromide: evidence of systemic and immunologic toxicity without impairment of in vivo host defenses

European medicinal and edible plants associated with subacute and chronic toxicity part II: Plants with hepato-, neuro-, nephro- and immunotoxic effects

Evaluating the effects of immunotoxicants using carbon fiber microelectrode amperometry

Evaluation of a new nanoparticle-based lateral-flow immunoassay for the exclusion of heparin-induced thrombocytopenia (HIT)

Evaluation of Acute Immunotoxicity of Aerosolized Aflatoxin B(1) in Female C57BL/6N Mice

Evaluation of acute immunotoxicity of alachlor in male F344/N rats

Evaluation of cyclosporine-induced nephrotoxicity by immunoperoxidase analysis of renal allograft biopsies

Evaluation of cytokine toxicity induced by vaccinia virus-mediated IL-2 and IL-12 antitumour immunotherapy

Evaluation of fetal skeletal malformations in deoxynivalenol-treated mice using microarray analysis

Evaluation of furfuryl alcohol sensitization potential following dermal and pulmonary exposure: enhancement of airway responsiveness

Evaluation of genotoxic and immunotoxic activities of potential glucose biosensor components: ferrocenes

Evaluation of household dust mite exposure and levels of specific IgE and IgG antibodies in asthmatic patients enrolled in a trial of immunotherapy

Evaluation of immune-mediated idiosyncratic drug toxicity using chimeric HLA transgenic mice

Evaluation of immunomodulatory effects of zearalenone in mice

Evaluation of immunoresponses and cytotoxicity from skin exposure to metallic nanoparticles

Evaluation of immunostimulant activity and chemoprotective effect of mangrove *Rhizophora apiculata* against cyclophosphamide induced toxicity in BALB/c mice

Evaluation of immunotoxic and immunodisruptive effects of inorganic arsenite on human monocytes/macrophages

Evaluation of immunotoxicity induced by diazinon in C57bl/6 mice

Evaluation of immunotoxicity induced by pirimiphos-methyl in male Balb/c mice following exposure to for 28 days

Evaluation of immunotoxicity induced by single or concurrent exposure to N,N-diethyl-m-tolamide (DEET), pyridostigmine bromide (PBR), and JP-8 jet fuel

Evaluation of immunotoxicity testings using azathioprine-treated rats: the International Collaborative Immunotoxicity Study (Azathioprine)

Evaluation of immunotoxicity tests using cyclosporin A-treated rats: the International Collaborative Immunotoxicity Study II (cyclosporin A)

Evaluation of in vivo acute immunotoxicity of a major organic arsenic compound arsenobetaine in seafood

Evaluation of multivariate statistical methods for analysis and modeling of immunotoxicology data

Evaluation of nanoparticle immunotoxicity

Evaluation of neuroprotective effects of insulin on immuno-inflammatory and systemic disorders induced by kalitoxin, a Kv1.3 channel blocker

Evaluation of neurotoxic and immunotoxic effects of trichloroacetic acid on rats

Evaluation of niridazole as a suppressant of cellular immunity in chickens

Evaluation of non-specific immune functions in dab, *Limanda limanda* L., following short-term exposure to sediments contaminated with polyaromatic hydrocarbons and/or polychlorinated biphenyls

Evaluation of potential immunotoxicity of pesticides in human lymphocyte tests

Evaluation of relative potencies of PCB126 and PCB169 for the immunotoxicities in ovalbumin (OVA)-immunized mice

Evaluation of sex- and strain-dependency of cocaine-induced immunosuppression in B6C3F1 and DBA/2 mice

Evaluation of specific immune responses to BoNT/A and tetanus toxoid in patients undergoing treatment for neurologic disorders

Evaluation of suppressive effects of paraquat on innate immunity in Balb/c mice

Evaluation of the adverse effect of low concentration of cadmium on interleukin-4 induced class switch recombination in Burkett's lymphoma Raji cell line

Evaluation of the clinical effectivity and toxicity of the FDN regimen (fludarabin, mitoxantron, dexamethason) in patients with follicular lymphoma

Evaluation of the humoral immune response of CD rats following a 2-week exposure to the pesticide carbaryl by the oral, dermal, or inhalation routes

Evaluation of the immunological cellular response of Cebus apella exposed to the carcinogen N-methyl-N-nitrosourea and treated with CANOVA

Evaluation of the immunotoxic potential of chlordecone with comparison to cyclophosphamide

Evaluation of the immunotoxicity of orally administered 2-methoxyacetic acid in Fischer 344 rats

Evaluation of the Multi-ImmunoTox Assay composed of 3 human cytokine reporter cells by examining immunological effects of drugs

Evaluation of the potential immunotoxicity of 3-monochloro-1,2-propanediol in Balb/c mice II. Effect on thymic subset, delayed-type hypersensitivity, mixed-lymphocyte reaction, and peritoneal macrophage activity

Evaluation of the potential immunotoxicity of 3-monochloro-1,2-propanediol in Balb/c mice. I. Effect on antibody forming cell, mitogen-stimulated lymphocyte proliferation, splenic subset, and natural killer cell activity

Evaluation of the potential immunotoxicity of bromodichloromethane in rats and mice

Evaluation of the utility of popliteal lymph node examination in a cyclophosphamide model of immunotoxicity in the rat

Evaluation of toxicological monitoring markers using proteomic analysis in rats exposed to formaldehyde

Evidence for a novel endocrine disruptor: the pesticide propanil requires the ovaries and steroid synthesis to enhance humoral immunity

Evidence for heterogeneous TCR V beta repertoire expression in mercury-induced immune disorders in rats

Evidence for immunotoxic effects of crude Ginkgo biloba L. leaf extracts using the popliteal lymph node assay in the mouse

Evidence for the induction of apoptosis in thymocytes by 2,3,7,8-tetrachlorodibenzo-p-dioxin in vivo

Evidence implicating rifampin-independent antiplatelet antibodies in the pathogenesis of rifampin-induced immune thrombocytopenia

Evidence of an immune-mediated mechanism for an idiosyncratic nevirapine-induced reaction in the female Brown Norway rat

Evidence of immunocompetence reduction induced by cadmium exposure in honey bees (*Apis mellifera*)

Evidence that the immunopathogenic mechanism of lithium-induced psoriasis differs from that of regular psoriasis

Evidence-based strategies to reduce intravenous immunoglobulin-induced headaches

Ex vivo cytokine release and pattern recognition receptor expression of subjects exposed to dampness: pilot study to assess the outcome of mould exposure to the innate immune system

Ex vivo immunotherapy for patients with benzene-induced aplastic anemia

Exacerbation of allergic contact dermatitis during immunosuppression with cyclosporine A

Exacerbation of an autoimmune thrombocytopenic purpura during treatment with interferon alpha in a woman with chronic viral hepatitis C

Exacerbation of atopic dermatitis symptoms by ustekinumab in psoriatic patients with elevated serum immunoglobulin E levels: Report of two cases

Exacerbation of autoimmune arthritis by copolymer-I through promoting type 1 immune response and autoantibody production

Exacerbation of benzene pneumotoxicity in connexin 32 knockout mice: enhanced proliferation of CYP2E1-immunoreactive alveolar epithelial cells

Exacerbation of invasive aspergillosis by the immunosuppressive fungal metabolite, gliotoxin

Examination of age-dependent effects of fetal ethanol exposure on behavior, hippocampal cell counts, and doublecortin immunoreactivity in rats

Examining the relationship between impaired host resistance and altered immune function in mice treated with TCDD

Excessive Cu<sup>2+</sup> deteriorates arsenite-induced apoptosis in chicken brain and resulting in immunosuppression, not in homeostasis

Exogenous Ca<sup>2+</sup> mitigates the toxic effects of TiO<sub>2</sub> nanoparticles on phagocytosis, cell viability, and apoptosis in haemocytes of a marine bivalve mollusk, *Tegillarca granosa*

Exogenous factors in the immunotoxicity of oral PMN

Exogenous melatonin modulates carbon ion radiation-induced immune dysfunction in mice

Experience of FK506 immune suppression in pediatric heart transplantation: a study of long-term adverse effects

Experimental fowl typhoid in chicks impaired immunologically by treatment with cyclophosphamide

Experimental immunoglobulin A nephropathy induced by gram-negative bacteria

Experimental induction of arthritis in rats immunized with *Escherichia coli* O:14 lipopolysaccharide

Experimental rabies in skunks: effects of immunosuppression induced by cyclophosphamide

Experimental studies on immunosuppression: how do they predict for man

Experimental Studies on Some Immunotoxicological Aspects of Aflatoxins Containing Diet and Protective Effect of Bee Pollen Dietary Supplement

Experimental T-2 toxicosis in swine following inhalation exposure: clinical signs and effects on hematology, serum biochemistry, and immune response

Experimental T-2 toxicosis in swine following inhalation exposure: effects on pulmonary and systemic immunity, and morphologic changes

Exploration of immunomodulatory and protective effect of *Withania somnifera* on trace metal oxide (zinc oxide nanoparticles) induced toxicity in Balb/c mice

Exploratory behavior and recognition memory in medial septal electrolytic, neuro- and immunotoxic lesioned rats

Exploring hazards of acute exposure of Acephate in *Drosophila melanogaster* and search for l-ascorbic acid mediated defense in it

Exploring the immunotoxicity of carbon nanotubes

Exposure and immunological determinants in a murine model for toluene diisocyanate (TDI) asthma

Exposure Cessation During Adulthood Did Not Prevent Immunotoxicity Caused by Developmental Exposure to Low-Level Trichloroethylene in Drinking Water

Exposure of Japanese medaka (*Oryzias latipes*) to benzo[a]pyrene suppresses immune function and host resistance against bacterial challenge

Exposure to 1.8 GHz electromagnetic fields affects morphology, DNA-related Raman spectra and mitochondrial functions in human lympho-monocytes.

Exposure to 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD) suppresses the humoral and cell-mediated immune responses to influenza A virus without affecting cytolytic activity in the lung

Exposure to 900 MHz radiofrequency radiation induces caspase 3 activation in proliferating human lymphocytes.

Exposure to a firefighting overhaul environment without respiratory protection increases immune dysregulation and lung disease risk

Exposure to a Social Stressor Induces Translocation of Commensal Lactobacilli to the Spleen and Priming of the Innate Immune System

Exposure to ambient particulate matter alters the microbial composition and induces immune changes in rat lung

Exposure to benzene induces oxidative stress, alters the immune response and expression of p53 in gasoline filling workers

Exposure to benzo[a]pyrene of Hepatic Cytochrome P450 Reductase Null (HRN) and P450 Reductase Conditional Null (RCN) mice: Detection of benzo[a]pyrene diol epoxide-DNA adducts by immunohistochemistry and 32P-postlabelling

Exposure to bifenthrin causes immunotoxicity and oxidative stress in male mice

Exposure to bisphenol A during gestation and lactation causes loss of sex difference in corticotropin-releasing hormone-immunoreactive neurons in the bed nucleus of the stria terminalis of rats

Exposure to cypermethrin and mancozeb alters the expression profile of THBS1, SPP1, FEZ1 and GPNMB in human peripheral blood mononuclear cells

Exposure to Deepwater Horizon oil and Corexit 9500 at low concentrations induces transcriptional changes and alters immune transcriptional pathways in sheepshead minnows

Exposure to diclofop-methyl induces immunotoxicity and behavioral abnormalities in zebrafish embryos

Exposure to ethanol during the last trimester of pregnancy alters the maturation and immunity of the fetal lung

Exposure to inhibitors of the renin-angiotensin system is a major independent risk factor for acute renal failure induced by sucrose-containing intravenous immunoglobulins: a case-control study

Exposure to inhibitors of the renin-angiotensin system is a major independent risk factor for acute renal failure induced by sucrose containing intravenous immunoglobulins. A case-control study

Exposure to lead during critical windows of embryonic development: differential immunotoxic outcome based on stage of exposure and gender

Exposure to mercuric chloride during the induction phase and after the onset of collagen-induced arthritis enhances immune/autoimmune responses and exacerbates the disease in DBA/1 mice

Exposure to mercuric chloride induces developmental damage, oxidative stress and immunotoxicity in zebrafish embryos-larvae

Exposure to mercury and population health. I. Immunotoxicity of mercury

Exposure to mycotoxins increases the allergic immune response in a murine asthma model

Exposure to negatively charged-particle dominant air-conditions on human lymphocytes in vitro activates immunological responses

Exposure to per-fluoroalkyl and polyfluoroalkyl substances leads to immunotoxicity: epidemiological and toxicological evidence



Exposure to radiation from single or combined radio frequencies provokes macrophage dysfunction in the RAW 264.7 cell line.

Exposure to sodium tungstate and Respiratory Syncytial Virus results in hematological/immunological disease in C57BL/6J mice

Exposure to TiO<sub>2</sub> Nanoparticles Induces Immunological Dysfunction in Mouse Testitis

Exposure to toxic heavy metals--their sequelae for immune competence in the human

Exposures to the environmental toxicants pentachlorophenol (PCP) and dichlorodiphenyltrichloroethane (DDT) modify secretion of interleukin 1-beta (IL-1beta) from human immune cells

Expression analysis of XPhyH-like during development and tail regeneration in *Xenopus* tadpoles: possible role of XPhyH-like expressing immune cells in impaired tail regenerative ability

Expression of CD44 and L-selectin in the innate immune system is required for severe joint inflammation in the proteoglycan-induced murine model of rheumatoid arthritis

Expression of dynamin immunoreactivity in experimental pancreatic tumors induced in rat by mancozeb-nitrosomethylurea

Extension of the protocol of OECD guideline 407 (28-day repeated dose oral toxicity test in the rat) to detect potential immunotoxicity of chemicals

Extensive metastatic Kaposi sarcoma in chronic immune suppressed bronchial asthma

External gamma irradiation-induced effects in early-life stages of zebrafish, *Danio rerio*

Extracellular signal-regulated kinase-signaling-dependent G2/M arrest and cell death in murine macrophages by cadmium

Extractable organic matter of standard reference material 1649a influences immunological response induced by pathogen-associated molecular patterns

Extremely low-level microwaves attenuate immune imbalance induced by inhalation exposure to low-level toluene in mice

Failure of peritoneal exudate macrophages to reverse immunologic impairment by Friend leukemia virus

False hyperglycemia induced by polivalent immunoglobulins

False-negative direct immunofluorescence testing in vancomycin-induced linear IgA bullous dermatosis: a diagnostic pitfall

Fasciola hepatica tegumental coat impairs mast cells' ability to drive Th1 immune responses

Fatal ceftriaxone-induced hemolysis in a child with acquired immunodeficiency syndrome

Fate of silver nanoparticles in wastewater and immunotoxic effects on rainbow trout

Favorable outcome of severe, extensive, granulocyte colony-stimulating factor-induced, corticosteroid-resistant Sweet's syndrome treated with high-dose intravenous immunoglobulin

Fc receptor binding of anti-CD3 monoclonal antibodies is not essential for immunosuppression, but triggers cytokine-related side effects

Feasibility of the extended one-generation reproductive toxicity study (OECD 443)

Features of immune response in chronic exposure to industrial aerosols

Feraheme suppresses immune function of human T lymphocytes through mitochondrial damage and mitoROS production

Fermented camel milk by *Lactococcus lactis* subsp. *cremoris* attenuates erythrocytes oxidative stress-induced hematological and immunological damage in CCl<sub>4</sub>-intoxicated mice

Fermented Red Ginseng Alleviates Cyclophosphamide-Induced Immunosuppression and 2,4,6-Trinitrobenzenesulfonic Acid-Induced Colitis in Mice by Regulating Macrophage Activation and T Cell Differentiation

Fetal and maternal immune responses to methylmercury exposure: a cross-sectional study

Field biomonitoring using the zebra mussel *Dreissena polymorpha* and the quagga mussel *Dreissena bugensis* following immunotoxic responses. Is there a need to separate the two species

Fine Particulate Matter-Induced Exacerbation of Allergic Asthma via Activation of T-cell Immunoglobulin and Mucin Domain 1

First report of a B cell lymphoproliferative disorder arising in a patient treated with immune suppressants for severe aplastic anemia

First studies on the bioconcentration and immunotoxicity of tetrachlorodiarlymethanes in the rat

Fish oil infusion reverses 5-fluorouracil-induced impairments in mucosal immunity in mice

FK506 treatment of graft-versus-host disease developing or exacerbating during prophylaxis and therapy with cyclosporin and/or other immunosuppressants. Japanese FK506 BMT Study Group

Flavored e-cigarette liquids and cinnamaldehyde impair respiratory innate immune cell function

Florfenicol impairs the immune responses to vaccination against foot-and-mouth disease in mice

Florfenicol induces more severe hemotoxicity and immunotoxicity than equal doses of chloramphenicol and thiamphenicol in Kunming mice

Flow cytometric analysis of in vitro cytotoxicity of cadmium in haemocytes from the tiger shrimp, *Penaeus monodon*

Flow cytometric analysis of lymphocyte subpopulations in the spleen and thymus of mice exposed to an acute immunosuppressive dose of 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD)

Flow cytometry detection of lysosomal presence and lysosomal membrane integrity in the three-spined stickleback (*Gasterosteus aculeatus* L.) immune cells: applications in environmental aquatic immunotoxicology

Flucloxacillin-induced immune thrombocytopenia

Fluconazole induces genotoxicity in cultured human peripheral blood mononuclear cells via immunomodulation of TNF-alpha, IL-6, and IL-10: new challenges for safe therapeutic regimens

Fluoride exposure abates pro-inflammatory response and induces invivo apoptosis rendering zebrafish (*Danio rerio*) susceptible to bacterial infections

Fluoride-induced immunotoxicity in adult male albino rat: a correlative approach to oxidative stress

Fluoroquinolone-induced acute interstitial nephritis in immunocompromised patients: two case reports

Fluoxetine reduces murine graft-versus-host disease by induction of T cell immunosuppression

Fluoxetine suppresses the immune responses of blood clams by reducing haemocyte viability, disturbing signal transduction and imposing physiological stress

Focused Screening Identifies Evoxine as a Small Molecule That Counteracts CO<sub>2</sub>-Induced Immune Suppression

Follow-up and immunologic findings in drug-induced myasthenia

Fondaparinux cross-reactivity in heparin-induced thrombocytopenia successfully treated with high-dose intravenous immunoglobulin and rivaroxaban

Food Protein Induced Proctocolitis: A Benign Condition with an Obscure Immunologic Mechanism

Food safety

Food-grade TiO<sub>2</sub> impairs intestinal and systemic immune homeostasis, initiates preneoplastic lesions and promotes aberrant crypt development in the rat colon

For Better or Worse, Iron Overload by Superparamagnetic Iron Oxide Nanoparticles as a MRI Contrast Agent for Chronic Liver Diseases

Fos-like immunoreactivity in auditory and nonauditory brain structures of hamsters previously exposed to intense sound

Fraction From *Lycium barbarum* Polysaccharides Reduces Immunotoxicity and Enhances Antitumor Activity of Doxorubicin in Mice

*Francisella tularensis* Catalase Restricts Immune Function by Impairing TRPM2 Channel Activity

From immunotoxicity to carcinogenicity: the effects of carbamate pesticides on the immune system

From immunotoxicity to nanotherapy: the effects of nanomaterials on the immune system

FTY720 immunosuppression impairs effector T cell peripheral homing without affecting induction, expansion, and memory

Full but impaired activation of innate immunity effectors and virus-specific T cells during CMV and EBV disease following cord blood transplantation

Fully human anti-BAFF inhibitory monoclonal antibody tabalumab does not adversely affect T-dependent antibody responses in cynomolgus monkey (*Macaca fascicularis*): A summary of three pre-clinical immunotoxicology evaluations

Functional activity of cells suppressing the humoral immune response in internal irradiation

Functional disorder of primary immunity responding to respiratory syncytial virus infection in offspring mice exposed to a flame retardant, decabrominated diphenyl ether, perinatally

Functional suppression of macrophages derived from THP-1 cells by environmentally-relevant concentrations of arsenite

Functional testing of the immunotoxicity of the phenoxyalkane acid mecoprop in rats and prevention of immunotoxic effects by adrenalectomy

Functionalization impacts the effects of carbon nanotubes on the immune system of rainbow trout, *Oncorhynchus mykiss*

Further evaluation of the incorporation of an immunotoxicological functional assay for assessing humoral immunity for hazard identification purposes in rats in a standard toxicology study

Fusaric Acid immunotoxicity and MAPK activation in normal peripheral blood mononuclear cells and Thp-1 cells

Fusidic acid induced acute immunologic thrombocytopenia

Ganoderma atrum polysaccharide improves age-related oxidative stress and immune impairment in mice

GC-TOF/MS-based metabolomics approach to study the cellular immunotoxicity of deoxynivalenol on murine macrophage ANA-1 cells

Gender differences in developmental immunotoxicity to lead in the chicken: analysis following a single early low-level exposure in ovo

Gender differences in the disposition and toxicity of metals

Gender-based profiles of developmental immunotoxicity to lead in the rat: assessment in juveniles and adults

Gender-related early immune changes in mice exposed to airborne suspended matter

Gene expression alterations in immune system pathways following exposure to immunosuppressive chemicals

Gene expression alterations in immune system pathways in the thymus after exposure to immunosuppressive chemicals

Gene expression changes in the skin of rats induced by prolonged 35 GHz millimeter-wave exposure.

Gene expression of heat shock protein 70, interleukin-1beta and tumor necrosis factor alpha as tools to identify immunotoxic effects on *Xenopus laevis*: a dose-response study with benzo[a]pyrene and its degradation products

Gene expression of mesothelioma in vinylidene chloride-exposed F344/N rats reveal immune dysfunction, tissue damage, and inflammation pathways

Gene expression profiling in spleens of deoxynivalenol-exposed mice: immediate early genes as primary targets

Gene expression profiling of Bis(tri-n-butyltin)oxide (TBTO)-induced immunotoxicity in mice and rats

Gene profiling of narrowband UVB-induced skin injury defines cellular and molecular innate immune responses

Generalized cryptococcosis in secondary immunodeficiency induced by long-term steroid therapy (on the 100th anniversary of the day of first discovery in 1894)

Generation of free radicals in haemocytes of mussels after exposure to low molecular weight PAH components: immune activation, oxidative and genotoxic effects

Genetic and immune-toxicologic studies on abnormal thyroid functions in hospital employees exposed to cytostatic drugs

Genetic differences in lethality of newborn mice treated in utero with coplanar versus non-coplanar hexabromobiphenyl

Genetic study of gold-salt-induced immune disorders in the rat

Genetic susceptibility and immune-mediated destruction in beryllium-induced disease

Genipin attenuates sepsis-induced immunosuppression through inhibition of T lymphocyte apoptosis

Geno- and immunotoxic effects on populations living near a mine: a case study of Panasqueira mine in Portugal

Genotoxic and apoptotic effects of Goeckerman therapy for psoriasis

Genotoxic and Cytotoxic Effects on the Immune Cells of the Freshwater Bivalve *Dreissena polymorpha* Exposed to the Environmental Neurotoxin BMAA

Genotoxic and immunotoxic effects of cellulose nanocrystals in vitro

Genotoxic and immunotoxic effects of the organophosphate metabolite diethyldithiophosphate (DEDTP) in Vivo

Genotoxic and immunotoxic potential effects of selected psychotropic drugs and antibiotics on blue mussel (*Mytilus edulis*) hemocytes

Genotoxicity and immunotoxic effects of 1,2-dichloroethane in Wistar rats

Genotoxicity induced by monomethylarsonous acid (MMA+3) in mouse thymic developing T cells

Genotoxicity of stack gas condensates of Bavarian waste incineration plants. I. Stack gas condensate-- sample collection techniques

Gestational cadmium exposure-induced ovotoxicity delays puberty through oxidative stress and impaired steroid hormone levels

Gestational exposure to perfluorooctane sulfonate suppresses immune function in B6C3F1 mice

Gestational zinc deficiency impairs humoral and cellular immune responses to hepatitis B vaccination in offspring mice

Glutathione Suppresses Macrophage Immune Function by Subverting Phosphatidylinositol 3,4,5-Trisphosphate Homeostasis

Glivec induced autoimmune hepatitis

Global effects of subchronic treatment of microcystin-LR on rat splenic protein levels

Glue solvent inhalation impairs host resistance to Mycobacterium bovis-induced infection in hamsters

Glutamic acid decarboxylase and GABA immunoreactivities in the cerebellar cortex of adult rat after prenatal exposure to a low concentration of carbon monoxide

Glutathione antagonized cyclophosphamide- and acrolein-induced cytotoxicity of PC3 cells and immunosuppressive actions in mice

Glutathione S-transferase polymorphisms and ochratoxin A toxicity in primary human urothelial cells

Glutathione synthesis is not involved in protection by N-acetylcysteine against UVB-induced systemic immunosuppression in mice

Glycation of extracellular matrix proteins impairs migration of immune cells

Glycogen synthase kinase-3 facilitates con a-induced IFN-gamma-- mediated immune hepatic injury

Glycopeptide-induced cutaneous adverse reaction: results of an immunoallergic investigation in eight patients

Glycoprotein IIb/IIIa complex is the target in mirtazapine-induced immune thrombocytopenia

Glycyrrhizic acid modulates the atrazine-induced apoptosis in rabbit spleen

Gonadotropin-releasing hormone agonist prevents l-arginine induced immune dysfunction independent of gonadal steroids: Relates with a decline in elevated thymus and brain nitric oxide levels

Granulocyte colony stimulating factor-induced exacerbation of fungus-related immune restoration inflammatory syndrome: a case of chronic disseminated candidiasis exacerbation

Granulocyte-macrophage colony stimulating factor-induced immune priming of cyclophosphamide, doxorubicin, vincristine, and prednisone with rituximab chemoimmunotherapy in previously untreated patients with diffuse large B-cell lymphoma and mantle cell lymphoma

Granulocyte-macrophage colony-stimulating factor and interferon-gamma prevent dexamethasone-induced immunosuppression of antifungal monocyte activity against *Aspergillus fumigatus* hyphae

Grape Seed Proanthocyanidin Extract Alleviates AflatoxinB-Induced Immunotoxicity and Oxidative Stress via Modulation of NF-kappaB and Nrf2 Signaling Pathways in Broilers

Graphene Oxide Modulates B Cell Surface Phenotype and Impairs Immunoglobulin Secretion in Plasma Cell

Growth, blood health, antioxidant status and immune response in juvenile yellow catfish *Pelteobagrus fulvidraco* exposed to alpha-ethinylestradiol (EE2)

Growth, blood health, antioxidant status, immune response and resistance to *Aeromonas hydrophila* of juvenile yellow catfish exposed to di-2-ethylhexyl phthalate (DEHP)

Guidance on setting of acute reference dose (ARfD) for pesticides

Gut dysbiosis and impairment of immune system homeostasis in perinatally-exposed mice to Bisphenol A precede obese phenotype development

Hair growth modulation by topical immunophilin ligands: induction of anagen, inhibition of massive catagen development, and relative protection from chemotherapy-induced alopecia

Halogenated aromatic hydrocarbons (HAH) as immunotoxicants

Halogenated aromatic hydrocarbons and toxicity equivalency factors (TEFs) from the public health assessment perspective

Halogenated aryl hydrocarbon-induced suppression of the in vitro plaque-forming cell response to sheep red blood cells is not dependent on the Ah receptor

Halogenated flame retardants: do the fire safety benefits justify the risks

Haloperidol suppresses murine dendritic cell maturation and priming of the T helper 1-type immune response

Halothane-induced suppression of cell-mediated immunity in normal and tumor-bearing C3Hf/He mice

Hapten-induced model of murine inflammatory bowel disease: mucosa immune responses and protection by tolerance

HCMV pUL135 remodels the actin cytoskeleton to impair immune recognition of infected cells

Health assessment of gasoline and fuel oxygenate vapors: immunotoxicity evaluation

Health effects following subacute exposure to geogenic dusts from arsenic-rich sediment at the Nellis Dunes Recreation Area, Las Vegas, NV

Heavy metal mediated innate immune responses of the Indian green frog, *Euphlyctis hexadactylus* (Anura: Ranidae): Cellular profiles and associated Th1 skewed cytokine response

Heavy metal-induced immunotoxicity and its mechanisms

Heavy metal-induced toxicity in the Indian green frog: Biochemical and histopathological alterations

Heavy metal-specific inhibition of phagocytosis and different in vitro sensitivity of heterogeneous coelomocytes from *Lumbricus terrestris* (Oligochaeta)

Helminth derived Immunomodulatory Glycan LNFP3 Impairs Pathogenesis of Peripheral Neuropathic Pain and Spinal Glial Activation

Helper T cell subpopulations from women are more susceptible to the toxic effect of sodium arsenite in vitro

Hemato-immunologic impact of subchronic exposure to melamine and/or formaldehyde in mice

Hematological and immunological impairment following in-utero and postnatal exposure to aluminum sulfate in female offspring of albino rats

Hematotoxic and immunotoxic effects of nitrogen dioxide

Heparin induced autoimmune thrombocytopenia: assessment of a rapid functional assay

Heparin-induced immune thrombocytopenia - a clinical or laboratory diagnosis

Heparin-induced thrombocytopenia: correction of severe bleeding complication with intravenous immune globulin

Heparin-induced thrombocytopenia: further evidence for a unique immune response

Heparin-induced thrombocytopenia--response to intravenous immunoglobulin in vivo and in vitro

Hepatic dysfunction in patients with carcinoma who are severely thrombocytopenic and immunosuppressed

Hepatic Transcriptome Responses of Domesticated and Wild Turkey Embryos to Aflatoxin B

Hepatic veno-occlusive disease following sirolimus-based immune suppression

Hepatitis B immunoglobulin-induced hypercoagulability complicating liver transplantation necessitating ECMO, rescue hepatectomy, and retransplantation

Hepatitis B virus infection is associated with impaired immunological recovery during antiretroviral therapy in the Swiss HIV cohort study

Hepatitis B virus infection status and infertility causes in couples seeking fertility treatment-Indicator of impaired immune response

Hepatitis C virus infection prevalence and liver dysfunction in a cohort of B-cell non-Hodgkin's lymphoma patients treated with immunochemotherapy

Hepatitis caused by clometacin (Duperan). Retrospective study of 30 cases. A model of autoimmune drug-induced hepatitis

Hepatotoxic and immunotoxic effects produced by 1,3-dibromopropane and its conjugation with glutathione in female BALB/c mice



Hepatotoxicants induce cytokine imbalance in response to innate immune system

Hepatotoxicity and immunotoxicity of MC-LR on silver carp

Hepatotoxicity induced by new immunosuppressants

Heterogeneous V beta gene usage in mercury-induced immune disorders in rats

Hexachlorobenzene-induced immunomodulation and skin and lung lesions: a comparison between brown Norway, Lewis, and Wistar rats

Hexavalent chromium and its effect on health: possible protective role of garlic (*Allium sativum* Linn

High altitude impairs in vivo immunity in humans

High dispersity of carbon nanotubes diminishes immunotoxicity in spleen

High frequency of immune dysfunctions in asbestos workers and in patients with malignant mesothelioma

High immunologic risk living donor kidney transplant using bortezomib in a novel induction regimen without acute antibody mediated rejection

High-Content Imaging and Gene Expression Approaches To Unravel the Effect of Surface Functionality on Cellular Interactions of Silver Nanoparticles

High-dose dietary exposure of mice to perfluorooctanoate or perfluorooctane sulfonate exerts toxic effects on myeloid and B-lymphoid cells in the bone marrow and these effects are partially dependent on reduced food consumption

High-dose granulocyte-macrophage colony-stimulating factor-producing vaccines impair the immune response through the recruitment of myeloid suppressor cells

High-dose intravenous immunoglobulin monotherapy for drug-induced hypersensitivity syndrome

High-fat diet-derived free fatty acids impair the intestinal immune system and increase sensitivity to intestinal epithelial damage

Highly sensitive and selective immuno-capture/electrochemical assay of acetylcholinesterase activity in red blood cells: a biomarker of exposure to organophosphorus pesticides and nerve agents

Hippocampal neuronal loss, decreased GFAP immunoreactivity and cognitive impairment following experimental intoxication of rats with aluminum citrate

Histological and immunohistochemical effects of *Curcuma longa* on activation of rat hepatic stellate cells after cadmium induced hepatotoxicity

Histological and immunohistochemical evaluation of the chemopreventive role of lycopene in tongue carcinogenesis induced by 4-nitroquinoline-1-oxide

Histological and immunohistochemical study of the expression of p53 and ki-67 proteins in the mucosa of the tongue, pharynx and larynx of rats exposed to cigarette smoke

Histological, immunohistochemical and morphometric changes in lung tissue in juvenile mice experimentally exposed to *Stachybotrys chartarum* spores

Histological, ultrastructural and immunohistochemical studies on the protective effect of ginger extract against cisplatin-induced nephrotoxicity in male rats

Histone deacetylase inhibitors impair innate immune responses to Toll-like receptor agonists and to infection

Histone deacetylase inhibitors impair the host immune response against *Mycobacterium tuberculosis* infection

Histopathologic approaches to detect changes indicative of immunotoxicity

Histopathological and immunohistochemical analysis of Tilapia (*Oreochromis niloticus*) exposed to cylindrospermopsin and the effectiveness of N-Acetylcysteine to prevent its toxic effects

Histopathological and Immunohistochemical Characterization of Methyl Eugenol-induced Nonneoplastic and Neoplastic Neuroendocrine Cell Lesions in Glandular Stomach of Rats

Histopathological Indices and Inflammatory Response in the Digestive Gland of the Mussel *Mytilus galloprovincialis* as Biomarker of Immunotoxicity to Silver Nanoparticles

Histopathology and analyses of inflammation intensity in the gills of mussels exposed to silver nanoparticles: role of nanoparticle size, exposure time, and uptake pathways

HIV impairment of immune responses in dendritic cells

HLA-G impairs host immune response and predicts poor prognosis in pancreatic cancer

Hochu-ekki-to Treatment Improves Reproductive and Immune Modulation in the Stress-Induced Rat Model of Polycystic Ovarian Syndrome

Hormonal contraception and HIV-1 infection: medroxyprogesterone acetate suppresses innate and adaptive immune mechanisms

Host resistance assays as predictive models in styrene immunomodulation

How can a chemical element elicit complex immunopathology? Lessons from mercury-induced autoimmunity

How does optimism suppress immunity? Evaluation of three affective pathways

How microwave treatment of gluten affects its toxicity for celiac patients? A study on the effect of microwaves on the structure, conformation, functionality and immunogenicity of gluten

How Regulatory T-Cell Induction by Statins May Impair Influenza Vaccine Immunogenicity and Effectiveness

HPV-16 E7 but not E6 oncogenic protein triggers both cellular immunosuppression and angiogenic processes

HSP27 modulates survival signaling in endosulfan-exposed human peripheral blood mononuclear cells treated with curcumin

Huangqin-tang ameliorates dextran sodium sulphate-induced colitis by regulating intestinal epithelial cell homeostasis, inflammation and immune response

Human anti-mitochondria autoantibodies appearing in iproniazid-induced immunoallergic hepatitis recognize human liver monoamine oxidase B

Human cytomegalovirus induces systemic immune activation characterized by a type 1 cytokine signature

Human herpesvirus 8 (HHV-8) inhibits monocyte differentiation into dendritic cells and impairs their immunostimulatory activity

Human immunodeficiency virus treatment-induced adipose tissue pathology and lipodystrophy: prevalence and metabolic consequences

Human immunodeficiency virus type 1 gp120 and ethanol coexposure in rat organotypic brain slice cultures: Curtailment of gp120-induced neurotoxicity and neurotoxic mediators by moderate but not high ethanol concentrations

Human Immunodeficiency Virus-1 Impairs IFN-Alpha Production Induced by TLR-7 Agonist in Plasmacytoid Dendritic Cells

Human immunodeficiency virus-1 protein Tat induces excitotoxic loss of presynaptic terminals in hippocampal cultures

Human immunodeficiency virus-1 Tat protein and methamphetamine interact synergistically to impair striatal dopaminergic function

Human immunosuppression induced by sea nettle (*Chrysaora quinquecirrha*) venom

Human intravenous immunoglobulin-induced aseptic meningitis: a case report

Human leukocyte antigen (HLA)-B\*57:01-restricted activation of drug-specific T cells provides the immunological basis for flucloxacillin-induced liver injury

Human macrophages constitute targets for immunotoxic inorganic arsenic

Human Papillomavirus (HPV) induced cancers and prevention by immunization

Human Rhinovirus Impairs the Innate Immune Response to Bacteria in Alveolar Macrophages in Chronic Obstructive Pulmonary Disease

Human T cells are highly sensitive to suppression of mitogenesis by polycyclic aromatic hydrocarbons and this effect is differentially reversed by alpha-naphthoflavone

Humoral immune dysfunction as a result of prenatal exposure to diphenylhydantoin: correlation with the occurrence of physical defects

Humoral immunological profile of workers exposed to asbestos in asbestos mines

Humoral immunotoxicity of polychlorinated diphenyl ethers, phenoxyphenols, dioxins and furans present as contaminants of technical grade pentachlorophenol

Hydantoin-induced cutaneous pseudolymphoma with clinical, pathologic, and immunologic aspects of Sezary syndrome

Hydrolyzed fumonisin B1 induces less inflammatory responses than fumonisin B1 in the co-culture model of porcine intestinal epithelial and immune cells

Hydrophobic sodium fluoride-based nanocrystals doped with lanthanide ions: assessment of in vitro toxicity to human blood lymphocytes and phagocytes

Hydroxyethyl radicals in ethanol hepatotoxicity

Hydroxylated fullerenes inhibit neutrophil function in fathead minnow (*Pimephales promelas* Rafinesque, 1820

Hygienic evaluation of immune and endocrine systems and modifications of their relationship in reproductive-age women working under exposure to chemical factors in activated carbon emissions

Hypercholesterolemia impairs immunity to tuberculosis

Hypergravity-induced immunomodulation in a rodent model: hematological and lymphocyte function analyses

Hyperinsulinemia induced by highly active antiretroviral therapy in an adolescent with polycystic ovary syndrome who was infected with human immunodeficiency virus

Hypertension induced by immunosuppressive drugs: a comparative analysis between sirolimus and cyclosporine

Hyperthyroidism induced by amiodarone and hyperthyroidism induced by iodine. Histologic, immunohistochemical and ultrastructural aspects

Hypothermal stress induced differential expression profiles of the immune response gene, warm-temperature-acclimation associated 65-kDa protein (Wap65), in the liver of fresh water and seawater milkfish, *Chanos chanos*

Hypothesis: is Alzheimer's disease a metal-induced immune disorder

Hypoxia inducible factor-1 $\alpha$  contributes to UV radiation-induced inflammation, epidermal hyperplasia and immunosuppression in mice

Latrogenic skin cancer: induction by psoralen/ultraviolet A and immunosuppression of organ transplant recipients

Latrogenically induced spondylodiskitis due to *Mycobacterium xenopi* in an immunocompetent patient

Idelalisib impairs T-cell-mediated immunity in chronic lymphocytic leukemia

Identification of AhR-regulated genes involved in PAH-induced immunotoxicity using a highly-sensitive DNA chip, 3D-Gene Human Immunity and Metabolic Syndrome 9k

Identification of crassin acetate as a new immunosuppressant triggering heme oxygenase-1 expression in dendritic cells

Identification of informative features for predicting proinflammatory potentials of engine exhausts

Identification of novel long non-coding RNAs involved in bisphenol A induced immunotoxicity in fish primary macrophages

Identification of novel signature genes attesting arsenic-induced immune alterations in adult zebrafish (*Danio rerio*)

Identification of signal pathways for immunotoxicity in the spleen of common carp exposed to chlorpyrifos

Identification of Signaling Pathways Targeted by the Food Contaminant FB1: Transcriptome and Kinome Analysis of Samples from Pig Liver and Intestine

IgE antibodies to betalactams: relationship between the triggering hapten and the specificity of the immune response

IgY Reduces AFB1-Induced Cytotoxicity, Cellular Dysfunction, and Genotoxicity in Human L-02 Hepatocytes and Swan 71 Trophoblasts

IL-15 Superagonist-Mediated Immunotoxicity: Role of NK Cells and IFN-gamma

Imbalance of Th1 and Th2 cells in cardiac injury induced by ambient fine particles

Imbalanced immune responses involving inflammatory molecules and immune-related pathways in the lung of acute and subchronic arsenic-exposed mice

Imidacloprid induced alterations in oxidative stress, biochemical, genotoxic, and immunotoxic biomarkers in non-mammalian model organism *Galleria mellonella* L. (Lepidoptera: Pyralidae)

Immediate posttransplantation cotrimoxazole-induced immune thrombocytopenia

Immune activation and autoantibodies in humans with long-term inhalation exposure to formaldehyde

Immune activation during pregnancy in rats leads to a postpubertal emergence of disrupted latent inhibition, dopaminergic hyperfunction, and altered limbic morphology in the offspring: a novel neurodevelopmental model of schizophrenia

Immune alterations in humans exposed to the termiticide technical chlordane

Immune alterations in lead-exposed workers

Immune alterations in mice exposed to the herbicide simazine

Immune and reproductive system impairment in adult sea urchin exposed to nanosized ZnO via food

Immune cells and cardiovascular health in premenopausal women of rural India chronically exposed to biomass smoke during daily household cooking

Immune Checkpoint Inhibitor-Induced Colitis: Diagnosis and Management

Immune Checkpoint Inhibitor-Induced Hypoparathyroidism Associated With Calcium-Sensing Receptor-Activating Autoantibodies

Immune complex-mediated haemolytic anaemia and Evans syndrome induced by diclofenac

Immune Complex-Mediated Proliferative Glomerulonephritis Induced by Paclitaxel Treatment

Immune disorders and chronic exposure to organic solvents. Case-control study in the operators who are in organic synthesis laboratories

Immune disorders induced by exposure to pyrethroid insecticides

Immune disruption occurs through altered gut microbiome and NOD2 in arsenic induced mice: Correlation with colon cancer markers

Immune Dysfunction After Cardiac Surgery with Cardiopulmonary Bypass: Beneficial Effects of Maintaining Mechanical Ventilation

Immune dysfunction and increased oxidative stress state in diet-induced obese mice are reverted by nutritional supplementation with monounsaturated and n-3 polyunsaturated fatty acids

Immune dysfunction and liver damage of mice following exposure to lanthanoids

Immune dysfunction during alcohol consumption and murine AIDS: the protective role of dehydroepiandrosterone sulfate

Immune dysfunction in acute alcoholic hepatitis

Immune dysfunction in children after corrective surgery for congenital heart disease

Immune dysfunction in rabbits associated with chronic administration of enemas and rectal insemination

Immune dysregulation after cardiothoracic surgery and incidental thymectomy: maintenance of regulatory T cells despite impaired thymopoiesis

Immune effects of nickel

Immune Enhancement Effects of Codium fragile Anionic Macromolecules Combined with Red Ginseng Extract in Immune-Suppressed Mice

Immune function biomarkers in children exposed to lead and organochlorine compounds: a cross-sectional study

Immune function in female B(6)C(3)F(1) mice is modulated by DE-71, a commercial polybrominated diphenyl ether mixture

Immune function in offspring of nonhuman primates (*Macaca nemestrina*) exposed weekly to 1.8 g/kg ethanol during pregnancy: preliminary observations

Immune function is impaired with a mini nutritional assessment score indicative of malnutrition in nursing home elders with pressure ulcers

Immune haemolytic anaemia induced by allopurinol after liver transplantation

Immune hemolytic anemia with drug-induced antibodies to carboplatin and vincristine in a pediatric patient with an optic pathway glioma

Immune heparin-induced thrombocytopenia can occur in patients receiving clopidogrel and aspirin

Immune homeostasis impairment in acute carbon tetrachloride intoxicated rats corrected by administration of tocopherol acetate and unithiol

Immune modulation by cadmium and lead in the acute reporter antigen-popliteal lymph node assay

Immune modulation of ovalbumin-induced lung injury in mice using beta-glucosylceramide and a potential role of the liver

Immune modulatory effects of the foodborne contaminant citrinin in mice

Immune parameters are affected differently after cyclosporine A exposure in Fischer 344 rats and B6C3F1 mice: implications for immunotoxicology

Immune pathogenesis of heparin-induced thrombocytopenia

Immune response due to silica exposure in Egyptian phosphate mines

Immune response impairment, genotoxicity and morphological transformation induced by *Taenia solium* metacestode

Immune response in mouse experimental cholangitis associated with colitis induced by dextran sulfate sodium

Immune response of adult rats is altered by administration of diazepam in the first postnatal week

Immune response of the bay scallop, *Argopecten irradians*, after exposure to the algicide palmitoleic acid

Immune response proteins as predictive biomarkers of doxorubicin-induced cardiotoxicity in breast cancer patients

Immune response reduced by intense intellectual stress: changes in lymphocyte proliferation in medical students

Immune response to abamectin-induced oxidative stress in Chinese mitten crab, *Eriocheir sinensis*

Immune response to exhaust gases derived from two-cycle combustion engine following experimental exposure

Immune responses and expression of immune-related genes in swimming crab *Portunus trituberculatus* exposed to elevated ambient ammonia-N stress

Immune responses and resistance to viral-induced myocarditis in mice exposed to cadmium

Immune responses and ultrastructural changes of hemocytes in freshwater crab *Sinopotamon henanense* exposed to elevated cadmium

## Immune Responses in Rhinovirus-Induced Asthma Exacerbations

Immune responses in the aquatic gastropod *Lymnaea stagnalis* under short-term exposure to pharmaceuticals of concern for immune systems: Diclofenac, cyclophosphamide and cyclosporine A

Immune responses induced by diclofenac or carbamazepine in an oral exposure model using TNP-Ficoll as reporter antigen

Immune responses induced by intranasal imiquimod and implications for therapeutics in rhinovirus infections

Immune responses of a wall lizard to whole-body exposure to radiofrequency electromagnetic radiation.

Immune responses to different patterns of exposure to ovalbumin in a mouse model of allergic rhinitis

Immune state of workers exposed to nickel aerosols in galvanic production

Immune suppression and skin cancer development: regulation by NKT cells

Immune suppression by combination therapy with basiliximab and cyclosporin in high risk keratoplasty. A pilot study

Immune suppression in transcerebral exposure to 460-MHz microwaves

Immune suppression leading to hepatitis C virus re-emergence after sustained virological response

Immune suppression of IgG response against dairy proteins in major depression

Immune suppression of the echinoderm *Asterias rubens* (L.) following long-term ocean acidification

Immune suppression therapy in aplastic anemia: influencing factors on response and survival

Immune suppression with cyclosporin A-optimism and caution

Immune surveillance and sunlight-induced skin cancer

Immune system alteration in the rat after indirect exposure to methyl mercury chloride or methyl mercury sulfide

Immune system development in infants born to mothers with autoimmune disease, exposed in utero to immunosuppressive agents

Immune System Modifications Induced in a Mouse Model of Chronic Exposure to (90)Sr

Immune system. Relationship to anxiety disorders

Immune thrombocytopenia induced by fludarabine successfully treated with rituximab

Immune Thrombocytopenia Induced by Nivolumab in a Metastatic Non-Small Cell Lung Cancer Patient

Immune/Inflammatory Response and Hypocontractility of Rabbit Colonic Smooth Muscle After TNBS-Induced Colitis

Immune-associated toxicities induced by in vivo and in vitro exposure to interferon-alpha alone or in combination with nucleoside analogs



Immune-enhancing activity of *C. militaris* fermented with *Pediococcus pentosaceus* (GRC-ON89A) in CY-induced immunosuppressed model

Immune-mediated liver dysfunction after antiviral treatment in liver transplanted patients with hepatitis C: allo or autoimmune de novo hepatitis

Immune-mediated myopathy related to anti 3-hydroxy-3-methylglutaryl-coenzymeA reductase antibodies as an emerging cause of necrotizing myopathy induced by statins

Immune-mediated pancytopenia induced by oxaliplatin: a case report

Immune-mediated processes implicated in chemotherapy-induced peripheral neuropathy

Immune-protective, antioxidant and relative genes expression impacts of beta-glucan against fipronil toxicity in Nile tilapia, *Oreochromis niloticus*

Immune-regulatory transcriptional responses in multiple organs of Atlantic salmon after tributyltin exposure, alone or in combination with forskolin

Immune-related Colitis Induced by the Long-term Use of Nivolumab in a Patient with Non-small Cell Lung Cancer

Immune-related eosinophilia induced by anti-programmed death 1 or death-ligand 1 antibodies

Immune-suppressive activity of punicalagin via inhibition of NFAT activation

Immune-suppressive effect of thalidomide

Immunity induced by immunomodulators during echinococcal immunosuppression

Immunity to Polyomavirus BK Infection: Immune Monitoring to Regulate the Balance between Risk of BKV Nephropathy and Induction of Alloimmunity

Immunization of mice with LRP4 induces myasthenia similar to MuSK-associated myasthenia gravis

Immunization with *Porphyromonas gingivalis* enolase induces autoimmunity to mammalian alpha-enolase and arthritis in DR4-IE-transgenic mice

Immunization with the glutamate receptor-derived peptide GluR3B induces neuronal death and reactive gliosis, but confers partial protection from pentylentetrazole-induced seizures

Immuno- and neurotoxicological investigation of combined subacute exposure with the carbamate pesticide propoxur and cadmium in rats

Immuno-, neuro-, and general toxicologic animal studies on a synthetic pyrethroid: cypermethrin

Immunoallergic drug-induced hepatitis: lessons from halothane

Immunoassay approach for diagnosis of exposure to pyrrolizidine alkaloids

Immunobiological effect of bitemporal exposure of rabbits to microwaves.

Immunobiological efficacy and immunotoxicity of novel synthetically prepared fluoroquinolone ethyl 6-fluoro-8-nitro-4-oxo-1,4-dihydroquinoline-3-carboxylate

Immunochemical detection of UV-induced DNA damage and repair

Immunochemical determination of xenobiotics with endocrine disrupting effects

Immunochemical evidence supporting 2-pentylpyrrole formation on proteins exposed to 4-hydroxy-2-nonenal

Immunochemical methods of studying the mechanism of diclofenac-induced hepatitis

Immunochemical methods of studying the mechanism of diclofenac-induced hepatitis [corrected

Immunochemical study of epidermal growth factor in rats with mercuric chloride-induced acute renal failure

Immunocompetence and alterations in hepatic gene expression in rainbow trout exposed to CdS/CdTe quantum dots

Immunocompetence over the lifespan of mice exposed in utero to carbofuran or diazinon: I. Changes in serum immunoglobulin concentrations

IMMUNOCYTOCHEMICAL ANALYSIS OF THE DISTURBANCES IN THE STRUCTURE OF SYNAPTONEMAL COMPLEXES IN SPERMATOCYTE NUCLEI IN MICE UNDER EXPOSURE TO ROCKET FUEL COMPONENT

Immunocytochemical identification of DNA adducts, O6-methylguanine and 7-methylguanine, in respiratory and other tissues of rat, mouse and Syrian hamster exposed to 4-(methylnitrosamino)-1-(3-pyridyl)-1-butanone

Immunocytotoxicity, cytogenotoxicity and genotoxicity of cadmium-based quantum dots in the marine mussel *Mytilus galloprovincialis*

Immunodeficiency induced by immunosuppressive agents

Immuno-enhancement effects of ethanol extract from *Cyrtomium macrophyllum* (Makino) Tagawa on cyclophosphamide-induced immunosuppression in BALB/c mice

Immunoenhancement Effects of Glycosaminoglycan from *Apostichopus japonicus*: In Vitro and In Cyclophosphamide-Induced Immunosuppressed Mice Studies

Immuno-enhancement effects of *Platycodon grandiflorum* extracts in splenocytes and a cyclophosphamide-induced immunosuppressed rat model

Immuno-enhancement effects of Shenqi Fuzheng Injection on cyclophosphamide-induced immunosuppression in Balb/c mice

Immuno-enhancement effects of Yifei Tongluo Granules on cyclophosphamide-induced immunosuppression in Balb/c mice

Immunofluorescence evaluation of 4-hydroxynonenal and 8-hydroxy-2-deoxyguanosine activation in zebrafish (*Danio rerio*) larvae brain exposed (microinjected) to propyl gallate

Immunogenetically controlled autoimmune reactions induced by mercury, gold and D-penicillamine in laboratory animals: a review from the vantage point of premarketing safety studies

Immunogenetics of sunlight-induced skin cancer

Immunogenicity, Inflammation, and Lipid Accumulation in Cynomolgus Monkeys Infused with a Lipidated Tetranectin-ApoA-I Fusion Protein

Immunoglobulin and lymphocyte decrease concurrent with adverse reactions induced by methotrexate for RA

Immunoglobulin and lymphocyte responses following silica exposure in New Zealand mixed mice

Immunoglobulin E and autoantibodies in mercury-exposed workers

Immunoglobulin G Fc receptor FcγRIIIa 158 V/F polymorphism correlates with rituximab-induced neutropenia after autologous transplantation in patients with non-Hodgkin's lymphoma

Immunoglobulin G from patients with heparin-induced thrombocytopenia binds to a complex of heparin and platelet factor 4

Immunoglobulin levels in workers exposed to hexachlorobenzene

Immunoglobulin production is impaired in protein-deprived mice and can be restored by dietary protein supplementation

Immunoglobulin-induced aseptic meningitis: a case report

Immunoglobulins in infants of rubella-exposed mothers

Immunoglobulins in periodontal tissues. III. Concentrations of immunoglobulins in dilantin-induced and idiopathic gingival hyperplastic tissues

Immunoglobulins in persons with long-term exposure to halothane

Immunoglobulins, serum proteins & lactate dehydrogenase levels in workers exposed to cotton dust

Immunogold study of effects of prenatal exposure to lipopolysaccharide and/or valproic acid on the rat blood-brain barrier vessels

Immunohistochemical evaluation of sarcoglycans and integrins in gingival epithelium of multiple myeloma patients with bisphosphonate-induced osteonecrosis of the jaw

Immunohistochemical analysis of dichloroacetic acid (DCA)-induced hepatocarcinogenesis in male Fischer (F344) rats

Immunohistochemical analysis of Th1/Th2 cytokine profiles and androgen receptor expression in the pathogenesis of nifedipine-induced gingival overgrowth

Immunohistochemical and molecular study on the protective effect of curcumin against hepatic toxicity induced by paracetamol in Wistar rats

Immunohistochemical and ultrastructural study of hepatic sinusoidal linings during dichloropropanol-induced acute hepatic necrosis

Immunohistochemical changes in the mouse striatum induced by the pyrethroid insecticide permethrin

Immunohistochemical changes induced by repeated footshock stress: revelations of gender-based differences

Immunohistochemical characterization of fibroblast subpopulations in normal peritoneal tissue and in peritoneal dialysis-induced fibrosis

Immunohistochemical characterization of spontaneous and acrylonitrile-induced brain tumors in the rat

Immunohistochemical characterization of the basement membrane epitopes in bis(2-chloroethyl) sulfide-induced toxicity in mouse ear skin

Immunohistochemical demonstration of increased prostaglandin F $\alpha$  levels in the rat hippocampus following kainic acid-induced seizures

Immunohistochemical detection of apoptotic proteins, p53/Bax and JNK/FasL cascade, in the lung of rats exposed to cigarette smoke

Immunohistochemical detection of polyunsaturated fatty acid oxidation markers in acetaminophen-induced liver injury in rats

Immunohistochemical distribution of activated nuclear factor kappaB and peroxisome proliferator-activated receptors in carbon tetrachloride-induced chronic liver injury in rats

Immunohistochemical evaluation of cardiac connexin43 in rats exposed to low-frequency noise

Immunohistochemical evaluation of chemically induced rhabdomyosarcomas in rats: diagnostic utility of MyoD1

Immunohistochemical evaluation of oxidative stress in murine lungs after cigarette smoke exposure

Immunohistochemical Expression of Cyclin D1, Cytokeratin 20, and Uroplakin III in Proliferative Urinary Bladder Lesions Induced by o-Nitroanisole in Fischer 344/N Rats

Immunohistochemical expression of retinoblastoma and phospho-retinoblastoma protein in sheep lung exposed to fluoro-edenite fibers

Immunohistochemical features of 3,3',4,4'-tetrachloroazobenzene-induced rat gingival lesions

Immunohistochemical localization of brain-derived neurotrophic factor and glial cell line-derived neurotrophic factor in the superior olivary complex of mice after radiofrequency exposure

Immunohistochemical localization of neuronal and glial calcium-binding proteins in hippocampus of chronically low level lead exposed rhesus monkeys

Immunohistochemical localization of surfactant protein A in N-bis (2-hydroxypropyl) nitrosamine-induced lung tumors in rats

Immunohistochemical localization of toluene-induced c-Fos protein expression in the rat brain

Immunohistochemical localization of transforming growth factor beta, basic fibroblast growth factor and heparan sulphate glycosaminoglycan in gingival hyperplasia induced by nifedipine and phenytoin

Immunohistochemical study of 30 cases of cyclosporin A-induced gingival overgrowth

Immunohistochemical study of cyclosporin-induced gingival overgrowth in renal transplant recipients

Immunohistochemical study of macrophage migration inhibitory factor in rat liver fibrosis induced by thioacetamide

Immunohistochemical study of rat Peyer's patches in peptidoglycan-polysaccharide complex-induced chronic enteritis

Immunohistochemical study of rat renal interstitial fibrosis induced by repeated injection of cisplatin, with special reference to the kinetics of macrophages and myofibroblasts

Immunohistochemical study on galactosamine-induced subacute hepatitis in rats of JCL: Wistar-TGN (ARGHGEN) 1 Nts strain (Mini rats)

Immunohistochemical, morphological and histometrical analyses of follicular development in *Astyanax bimaculatus* (Teleostei: Characidae) exposed to an organochlorine insecticide

Immunohistochemistry of aberrant neuronal development induced by 6-propyl-2-thiouracil in rats

Immunohistological study of eosinophilic infiltration of nasal polyps in aspirin-induced asthma

Immunohistopathological analysis of frizzled-4-positive immature melanocytes from hair follicles of patients with Rhododenol-induced leukoderma

Immunolocalization of metallothionein in hepatocytes of guppy fish (*Poecilia reticulata*) exposed to tannery effluent: A biomarker study

Immunolocalization of MMP 2, 9 and 13 in prednisolone induced osteoporosis in mice

Immunologic abnormalities in humans exposed to chlorpyrifos: preliminary observations

Immunologic and endocrine effects of the flame-retardant pentabromodiphenyl ether (DE-71) in C57BL/6J mice

Immunologic and genetic markers of exposure to aromatic hydrocarbons in workers

Immunologic basis for allopurinol-induced severe cutaneous adverse reactions: HLA-B\*58:01-restricted activation of drug-specific T cells and molecular interaction

Immunologic dysfunction among PBB-exposed Michigan dairy farmers

Immunologic effects and tolerability profile of in-season initiation of a standardized-quality grass allergy immunotherapy tablet: a phase III, multicenter, randomized, double-blind, placebo-controlled trial in adults with grass pollen-induced rhinoconjunctivitis

Immunologic effects of background prenatal and postnatal exposure to dioxins and polychlorinated biphenyls in Dutch infants

Immunologic effects of cocaine in prenatally exposed rats and mice

Immunologic effects of nickel. II. Suppression of natural killer cell activity

Immunologic suppression by methotrexate in dermatology

Immunological abnormalities in workers exposed to pollutants at an Egyptian copper company

Immunological and cytotoxicological responses of the Asian clam, *Corbicula fluminea* (M.), experimentally exposed to cadmium

Immunological and genotoxic effects of occupational exposure to alpha-cypermethrin pesticide

Immunological and hematological effects observed in B6C3F1 mice exposed to JP-8 jet fuel for 14 days

Immunological and hematological toxicities challenging clinical translation of nucleic acid-based therapeutics

Immunological and neurobiochemical alterations induced by repeated oral exposure of phenol in mice

Immunological and physiological responses of the periwinkle *Littorina littorea* during and after exposure to the toxic dinoflagellate *Alexandrium minutum*

Immunological and respiratory reactions in workers exposed to organic dusts

Immunological aspects and therapeutic significance of an autoantibody against histone H1 in a rat model of concanavalin A-induced hepatitis

Immunological aspects of tri-o-tolyl phosphate-induced delayed neurotoxicity in chickens

Immunological biomarkers in salt miners exposed to salt dust, diesel exhaust and nitrogen oxides

Immunological changes in hydroxyurea-induced collagen disease

Immunological changes produced in rats by prenatal exposure to carbon monoxide

Immunological consequences of asbestos exposure

Immunological determinants in a murine model of toluene diisocyanate-induced asthma

Immunological disorders in men exposed to metallic mercury vapour. A review

Immunological dysfunction in chronic arsenic exposure: From subclinical condition to skin cancer

Immunological effects of 2-methoxyethanol administered dermally or orally to Fischer 344 rats

Immunological effects of chlorinated dibenzo-p-dioxins

Immunological effects of iron oxide nanoparticles and iron-based complex drug formulations: Therapeutic benefits, toxicity, mechanistic insights, and translational considerations

Immunological effects of silica/asbestos

Immunological evaluation of four arc welders exposed to fumes from ignited polyurethane (isocyanate) foam: antibodies and immune profiles

Immunological function in mice exposed to JP-8 jet fuel in utero

Immunological impacts of oil sands-affected waters on rainbow trout evaluated using an in situ exposure

Immunological impairment and susceptibility to infection after splenectomy

Immunological investigations in a group of workers exposed to various levels of polycyclic aromatic hydrocarbons

Immunological involvement in pulmonary fibrosis induced by peplomycin

Immunological mechanism in development of allergic dermatitis in guinea pig induced by trichloroethylene in vitro

Immunological profile of arsenic toxicity: a hint towards arsenic-induced carcinogenesis

Immunological response of white shrimp (*Litopenaeus vannamei*) to sublethal concentrations of malathion and endosulfan, and their mixture

Immunological responses of weanling cotton rats (*Sigmodon hispidus*) to acute benzene and cyclophosphamide exposure

Immunological studies in a case of hydrochlorothiazide-induced pulmonary edema

Immunological studies of the mucosa in colitis induced by sodium dextran sulfate in rats

Immunological studies on cataracts under conditions of exposure to low-dose radiation

Immunological surveillance and toxicity in mice exposed to the organophosphate pesticide, leptophos

Immunological tolerance in a mouse model of immune-mediated liver injury induced by 16,16 dimethyl PGE2 and PGE2-containing nanoscale hydrogels

Immunological, chemical and clinical aspects of exposure to mixtures of contact allergens

Immunology-related perturbations induced by copper and chitosan in carp (*Cyprinus carpio* L

Immunomodulating effects after perinatal exposure to methylmercury in mice

Immunomodulating effects induced by the regulators of energy metabolism and the exposure to heat and cold in case of toxic affection of the kidneys

Immunomodulation by gastrointestinal carbon black nanoparticle exposure in ovalbumin T cell receptor transgenic mice

Immunomodulation by metals

Immunomodulation in allogeneic marrow transplantation: use of intravenous immune globulin to suppress acute graft-versus-host disease

Immunomodulation in eastern oysters, *Crassostrea virginica*, exposed to a PAH-contaminated, microphytobenthic diatom

Immunomodulation of cell-mediated cytotoxicity after chronic exposure to vapors

Immunomodulation of phloretin by impairing dendritic cell activation and function

Immunomodulation of Rheum tanguticum polysaccharide (RTP) on the immunosuppressive effects of dexamethasone (DEX) on the treatment of colitis in rats induced by 2,4,6-trinitrobenzene sulfonic acid

Immunomodulation of the inflammatory response induced by Androctonus australis hector neurotoxins: biomarker interactions

Immunomodulative effects of aflatoxins and selenium on human peripheral blood lymphocytes

Immunomodulatory activity of a novel polysaccharide from Lonicera japonica in immunosuppressed mice induced by cyclophosphamide

Immunomodulatory activity of fucoidan against aspirin-induced gastric mucosal damage in rats

Immunomodulatory activity of Melaleuca alternifolia concentrate (MAC): inhibition of LPS-induced NF-kappaB activation and cytokine production in myeloid cell lines

Immunomodulatory activity of methanolic extract of Amorphophallus commutatus var. wayanadensis under normal and cyclophosphamide induced immunosuppressive conditions in mice models

Immunomodulatory activity of TNF-alpha during acute liver injury induced by D-galactosamine and its protection by PGE1 in rats

Immunomodulatory and anti-inflammatory effects of hydro-ethanolic extract of Ocimum basilicum leaves and its effect on lung pathological changes in an ovalbumin-induced rat model of asthma

Immunomodulatory and immunotoxic effects of bilirubin: molecular mechanisms

Immunomodulatory Effect of Agave tequilana Evaluated on an Autoimmunity Like-SLE Model Induced in Balb/c Mice with Pristane

Immunomodulatory effect of candesartan on indomethacin-induced gastric ulcer in rats

Immunomodulatory effect of Moringa oleifera Lam. extract on cyclophosphamide induced toxicity in mice

Immunomodulatory effect of new quinolone derivative against cisplatin/gamma radiation-induced renal and brain toxicity in mice

Immunomodulatory Effect of Tremella Polysaccharides against Cyclophosphamide-Induced Immunosuppression in Mice

Immunomodulatory effects in workers exposed to naturally occurring asbestos fibers

Immunomodulatory effects of azithromycin on serum amyloid A production in lipopolysaccharide-induced endotoxemia in mice

Immunomodulatory effects of estradiol and cadmium in adult female rats

Immunomodulatory effects of exposure to polychlorinated biphenyls and perfluoroalkyl acids in East Greenland ringed seals (Pusa hispida)

Immunomodulatory effects of gyokuheifusan on INF-gamma/IL-4 (Th1/Th2) balance in ovalbumin (OVA)-induced asthma model mice



Immunomodulatory effects of L-carnitine and q10 in mouse spleen exposed to low-frequency high-intensity magnetic field

Immunomodulatory effects of maternal atrazine exposure on male Balb/c mice

Immunomodulatory effects of *Premna tomentosa* (L. Verbenaceae) extract in J 779 macrophage cell cultures under chromate (VI)-induced immunosuppression

Immunomodulatory effects of *Premna tomentosa* extract against Cr (VI) induced toxicity in splenic lymphocytes--an in vitro study

Immunomodulatory effects of synthetic endocrine disrupting chemicals on the development and functions of human immune cells

Immunomodulatory effects of the fungicide Mancozeb in agricultural workers

Immunomodulatory effects upon in vitro exposure of California sea lion and southern sea otter peripheral blood leukocytes to domoic acid

Immunomodulatory mechanism of Bushen Huoxue Recipe alleviates cyclophosphamide-induced diminished ovarian reserve in mouse model

Immunomodulatory role of *Ocimum gratissimum* and ascorbic acid against nicotine-induced murine peritoneal macrophages in vitro

Immunomodulatory treatments for aspirin exacerbated respiratory disease

Immunomorphologic changes in the testes upon exposure to a microwave electromagnetic field.

Immunopathologic effects of silicone breast implants

Immunopathology of fatal soybean dust-induced asthma

Immunopharmacological studies on collagen-induced arthritis in dark Agouti (DA) rats

Immunophenotype, ras oncogenes and p53 onco-suppressor gene in benzo(a)pyrene induced malignant soft tissue tumours in Wistar rats

Immunophenotypic and functional effects of bunker C fuel oil on the immune system of American mink (*Mustela vison*)

Immunophenotypical analysis of myofibroblasts and mesenchymal cells in the bleomycin-induced rat scleroderma, with particular reference to their origin

Immuno-physiological alterations from AFB1 in rats counteracted by treatments with *Lactobacillus paracasei* BEJ01 and montmorillonite clay mixture

Immunoproteasome dysfunction augments alternative polarization of alveolar macrophages

Immunoprotective activity and antioxidant properties of cactus (*Opuntia ficus indica*) extract against chlorpyrifos toxicity in rats

Immunoprotective effect of epigallocatechin-3-gallate on oral anticancer drug-induced alpha-defensin reduction in Caco-2 cells

Immunoregulatory role of CD1d in the hydrocarbon oil-induced model of lupus nephritis

Immunoresponsiveness in endometriosis: implications of estrogenic toxicants

Immunostaining of proinflammatory cytokines in renal cortex and medulla of rats exposed to gold nanoparticles

Immunostimulating effects of protein A in immunosuppressed aflatoxin-intoxicated rats

Immunostimulative effects of repeated inhalation exposure to microvesicle-bound endotoxin of *Pantoea agglomerans*

Immunostimulatory oligodeoxynucleotide containing TTTCGTTT motif from *Lactobacillus rhamnosus* GG DNA potentially suppresses OVA-specific IgE production in mice

Immunosuppressant FK506 induces interleukin-6 production through the activation of transcription factor nuclear factor (NF)-kappa(B). Implications for FK506 nephropathy

Immunosuppressant Medication-Induced Lower Extremity Pain After Combined Liver and Kidney Transplant: A Case Report

Immunosuppressant prograf (tacrolimus) induces histopathological disorders in the peritubular tissue of rat testes

Immunosuppressant therapy and bone loss in ligature-induced periodontitis--a study in rats

Immunosuppressant-induced endothelial damage and pulmonary arterial hypertension

Immunosuppressant-induced nephropathy: pathophysiology, incidence and management

Immunosuppression and induction of tolerance by 3-(2-chloroethyl)-2(2-mesyloxyethylamino)-tetrahydro-2H-1,3,2-oxazaphosphorine-2-oxide (ASTA 5122)

Immunosuppression and oxidative stress induced by acute and chronic exposure to cocaine in rat

Immunosuppression and oxidative stress induced by subchronic exposure to carbosulfan in rat spleen: immunomodulatory and antioxidant role of N-acetylcysteine

Immunosuppression as initial treatment for gold induced aplastic anemia

Immunosuppression by arsenic: a comparison of cathepsin L inhibition and apoptosis

Immunosuppression by chronic exposure to N-nitrosodimethylamine (NDMA) in mice

Immunosuppression following 7,12-dimethylbenz[a]anthracene exposure in B6C3F1 mice--II. Altered cell-mediated immunity and tumor resistance

Immunosuppression following 7,12-dimethylbenz[a]anthracene exposure in B6C3F1 mice. I. Effects on humoral immunity and host resistance

Immunosuppression following exposure to 7,12-dimethylbenz[a]anthracene (DMBA) in Ah-responsive and Ah-nonresponsive mice

Immunosuppression in adult female B6C3F1 mice by chronic exposure to ethanol in a liquid diet

Immunosuppression in mice induced by cold water stress

Immunosuppression in mice induced by dioxin (TCDD) in feed

Immunosuppression in the infaunal bivalve *Scrobicularia plana* environmentally exposed to mercury and association with its accumulation

Immunosuppression in the mouse induced by long-term exposure to cigarette smoke

Immunosuppression in the northern leopard frog (*Rana pipiens*) induced by pesticide exposure

Immunosuppression in weanling and adult Sprague-Dawley rats induced by acute exposure to 3,3',4,4'-tetrachloroazoxybenzene

Immunosuppression induced by acute solar-simulated ultraviolet exposure in humans: prevention by a sunscreen with a sun protection factor of 15 and high UVA protection

Immunosuppression induced by ultraviolet radiation: relevance to public health

Immunosuppression induced by ultraviolet rays. Importance for development of non-melanoma skin cancer

Immunosuppression induction with daclizumab and antithymocyte globulin in cardiac transplantation: clinical experience with 8 cases

Immunosuppression of canine, monkey, and baboon allografts by FK 506: with special reference to synergism with other drugs and to tolerance induction

Immunosuppression of pulmonary natural killer activity by exposure to ozone

Immunosuppression With FTY720 Reverses Cardiac Dysfunction in Hypomorphic ApoE Mice Deficient in SR-BI Expression That Survive Myocardial Infarction Caused by Coronary Atherosclerosis

Immunosuppression without liver induction by subchronic exposure to 2,7-dichlorodibenzo-p-dioxin in adult female B6C3F1 mice

Immunosuppression-induced alterations in fish gut microbiota may increase the susceptibility to pathogens

Immunosuppression-induced leukoencephalopathy from tacrolimus (FK506)

Immunosuppressive activities of polychlorinated dibenzofuran congeners: quantitative structure-activity relationships and interactive effects

Immunosuppressive activity of a new pteridine derivative (4AZA1378) alleviates severity of TNBS-induced colitis in mice

Immunosuppressive activity of deer antler extracts of *Cervus korean* TEMMINCK var. *mantchuricus* Swinhoe, on type II collagen-induced arthritis

Immunosuppressive activity of polychlorinated biphenyl mixtures and congeners: nonadditive (antagonistic) interactions

Immunosuppressive drug-induced diabetes

Immunosuppressive drug-induced leukoencephalopathy in patients with liver transplant

Immunosuppressive effect of FK506 on collagen-induced arthritis in rats

Immunosuppressive effect of polycyclic aromatic hydrocarbons by induction of apoptosis of pre-B lymphocytes of bone marrow

Immunosuppressive effect of prolactin-induced protein

Immunosuppressive effect of subchronic exposure to a mixture of eight heavy metals, found as groundwater contaminants in different areas of India, through drinking water in male rats

Immunosuppressive effect of the decimeter-band electromagnetic field.

Immunosuppressive effects and associated compensatory responses in zebrafish after full life-cycle exposure to environmentally relevant concentrations of cadmium

Immunosuppressive effects of 2,3,7,8-tetrachlorodibenzo-p-dioxin in strains of mice with different susceptibility to induction of aryl hydrocarbon hydroxylase

Immunosuppressive effects of arsenic in broiler chicks exposed to Newcastle disease virus

Immunosuppressive effects of highly chlorinated biphenyls and diphenyl ethers on T-cell dependent and independent antigens in mice

Immunosuppressive potential of several polycyclic aromatic hydrocarbons (PAHs) found at a Superfund site: new model used to evaluate additive interactions between benzo[a]pyrene and TCDD

Immunosuppressive strategies and chronic graft dysfunction in kidney transplantation

Immunosuppressive switch to sirolimus in renal dysfunction after liver transplantation

Immunosuppressive therapy exacerbates autoimmunity in NOD mice and diminishes the protective activity of regulatory T cells

Immunosuppressive therapy for effective suppression of life threatening ventricular tachyarrhythmias in chronic myocarditis

Immunosuppressive therapy of cyclosporin A for severe benzene-induced haematopoietic disorders and a 6-month follow-up

Immunosuppressive TOR kinase inhibitor everolimus (RAD) suppresses growth of cells derived from posttransplant lymphoproliferative disorder at allograft-protecting doses

Immunosurgical studies on cytological and cytogenetic toxicity analysis of rat blastocysts after in vivo exposure to cyclophosphamide

Immunoteratology of chlordane: cell-mediated and humoral immune responses in adult mice exposed in utero

Immunotherapy in benzene-induced aplastic anemia using a biotherapeutic approach--is it durable

Immunotherapy with Chinese medicinal herbs. II. Reversal of cyclophosphamide-induced immune suppression by administration of fractionated *Astragalus membranaceus* in vivo

Immunotoxic activity of ochratoxin A

Immunotoxic and cancerostatic effects of ethyl-4-isothiocyanatobutanoate in female Lewis rats with implanted fibrosarcoma

Immunotoxic and cytotoxic effects of atrazine, permethrin and piperonyl butoxide to rainbow trout following in vitro exposure

Immunotoxic and genotoxic potential of arsenic and its chemical species in goats

Immunotoxic and hematotoxic effects of occupational exposures

Immunotoxic and hepatotoxic effects of perfluoro-n-decanoic acid (PFDA) on female Harlan Sprague-Dawley rats and B6C3F1/N mice when administered by oral gavage for 28 days

Immunotoxic catecholamine lesions attenuate 2DG-induced increase of AGRP mRNA

Immunotoxic changes associated with a 7-day oral exposure to perfluorooctanesulfonate (PFOS) in adult male C57BL/6 mice

Immunotoxic damage in floriculturists exposed to pesticide mixtures

Immunotoxic destruction of distinct catecholamine subgroups produces selective impairment of glucoregulatory responses and neuronal activation

Immunotoxic effect of asbestos

Immunotoxic effect of beta-chlorolactic acid on murine splenocyte and peritoneal macrophage function in vitro

Immunotoxic effect of herbicide simazine exposure in BALB/c mice

Immunotoxic effect of low dose cisplatin in mice

Immunotoxic effect of thiamethoxam in immunized mice with *Brucella abortus* cultural filtrate antigen

Immunotoxic effects during acute ethylene glycol poisoning

Immunotoxic effects of 2-bromopropane in male Sprague-Dawley rats: a 28-day exposure study

Immunotoxic effects of 4-nonylphenol on *Clarias gariepinus*: Cytopathological changes in hepatic melanomacrophages

Immunotoxic effects of a new antineoplastic agent S-1 in mice--comparison with S-1, UFT and 5-FU

Immunotoxic effects of an industrial waste incineration site on groundwater in rainbow trout (*Oncorhynchus mykiss*)

Immunotoxic effects of atrazine and its main metabolites at environmental relevant concentrations on larval zebrafish (*Danio rerio*)

Immunotoxic effects of carbon tetrachloride--the effect on morphology and function of the immune system in mice

Immunotoxic effects of chemicals: A matrix for occupational and environmental epidemiological studies

Immunotoxic effects of cis-urocanic acid exposure in C57BL/6N and C3H/HeN mice

Immunotoxic effects of copper and cadmium in the sea bass *Dicentrarchus labrax*

Immunotoxic effects of cyclophosphamide and cyclosporine in the dog

Immunotoxic effects of dialkyltins used for stabilization of plastics

Immunotoxic effects of diethylstilbestrol (DES) and cadmium chloride (CAD) on host resistance: comparison with cyclophosphamide (CPS)

Immunotoxic effects of diethylstilbestrol on host resistance: comparison with cyclophosphamide

Immunotoxic effects of environmental pollutants in marine mammals

Immunotoxic effects of environmental toxicants in fish - how to assess them

Immunotoxic effects of exposure of rats to xenobiotics via maternal lactation. Part I 2,3,7,8-tetrachlorodibenzo-p-dioxin

Immunotoxic effects of gold and silver nanoparticles: Inhibition of mitogen-induced proliferative responses and viability of human and murine lymphocytes in vitro

Immunotoxic effects of hexachlorobenzene on the pathogenesis of systemic, pneumonic and hepatic virus infections in the mouse

Immunotoxic effects of imidacloprid following 28 days of oral exposure in BALB/c mice

Immunotoxic effects of in vitro exposure of dolphin lymphocytes to Louisiana sweet crude oil and Corexit

Immunotoxic effects of inorganic lead on host resistance of mice with different circling behavior preferences

Immunotoxic effects of lead on birds

Immunotoxic effects of mercuric compounds on human lymphocytes and monocytes. IV. Alterations in cellular glutathione content

Immunotoxic effects of mercuric compounds on human lymphocytes and monocytes. III. Alterations in B-cell function and viability

Immunotoxic effects of mercuric compounds on human lymphocytes and monocytes. I. Suppression of T-cell activation

Immunotoxic effects of MPT-IP containing 60% methylparathion in mice

Immunotoxic effects of nickel in the mud crab *Scylla serrata*

Immunotoxic effects of Ochratoxin A in Wistar rats after oral administration

Immunotoxic effects of oil sands-derived naphthenic acids to rainbow trout

Immunotoxic effects of organotin compounds in *Tapes philippinarum*

Immunotoxic effects of perfluorononanoic acid on BALB/c mice

Immunotoxic effects of perfluorooctane sulfonate and di(2-ethylhexyl) phthalate on the marine fish *Oryzias melastigma*

Immunotoxic effects of prolonged dietary exposure of male rats to 2,3,7,8-tetrachlorodibenzo-p-dioxin

Immunotoxic effects of short-term atrazine exposure in young male C57BL/6 mice

Immunotoxic effects of single and combined pharmaceuticals exposure on a harbor seal (*Phoca vitulina*) B lymphoma cell line

Immunotoxic effects of smokeless tobacco on the accessory cell function of rat oral epithelium

Immunotoxic effects of sodium tungstate dihydrate on female B6C3F1/N mice when administered in drinking water

Immunotoxic effects of T-2 mycotoxin on cell-mediated resistance to *Listeria monocytogenes* infection

Immunotoxic effects of T-2 toxin on cell-mediated immunity to listeriosis in mice: comparison with cyclophosphamide

Immunotoxic effects of TCDD and toxic equivalency factors

Immunotoxic effects of the color additive caramel color III: immune function studies in rats

Immunotoxic effects of thymus in mice following exposure to nanoparticulate TiO<sub>2</sub>

Immunotoxic effects of triclosan in the clam *Ruditapes philippinarum*

Immunotoxic organotins as possible model compounds in studying apoptosis and thymocyte differentiation

Immunotoxic potencies of polychlorinated biphenyl (PCB), dibenzofuran (PCDF) and dibenzo-p-dioxin (PCDD) congeners in C57BL/6 and DBA/2 mice

Immunotoxic potential of aeration lagoon effluents for the treatment of domestic and hospital wastewaters in the freshwater mussel *Elliptio complanata*

Immunotoxic potential of antiviral drugs: effects of ganciclovir and (S)-1-(3-hydroxy-2-phosphonylmethoxy propyl) cytosine on lymphocyte transformation and delayed-type hypersensitivity responses

Immunotoxic Potential of Bisphenol F Mediated through Lipid Signaling Pathways on Macrophages

Immunotoxic potential of N-ethyl, N-nitrosourea (ENU) in CD1 mice

Immunotoxic response of oleic acid anilide and its hydrolysis products in female MRL (+/+) mice

Immunotoxic responses of chronic exposure to cypermethrin in common carp

Immunotoxic responses of cypermethrin, a synthetic pyrethroid insecticide in rats

Immunotoxicities of microplastics and sertraline, alone and in combination, to a bivalve species: size-dependent interaction and potential toxication mechanism

Immunotoxicity activity from the essential oils of coriander (*Coriandrum sativum*) seeds

Immunotoxicity activity of natural furocoumarins from milky sap of *Ficus carica* L. against *Aedes aegypti* L

Immunotoxicity and allergenic potential induced by topical application of perfluorooctanoic acid (PFOA) in a murine model

Immunotoxicity and allergic potential induced by topical application of dimethyl carbonate (DMC) in a murine model

Immunotoxicity and biodistribution analysis of arsenic trioxide in C57Bl/6 mice following a 2-week inhalation exposure

Immunotoxicity and disease resistance in Japanese quail (*Coturnix coturnix japonica*) exposed to malathion

Immunotoxicity and environment: immunodysregulation and systemic inflammation in children

Immunotoxicity and genotoxicity of single-walled carbon nanotubes co-exposed with cadmium in the freshwater mussel, *Elliptio complanata*

Immunotoxicity and genotoxicity testing of PLGA-PEO nanoparticles in human blood cell model

Immunotoxicity and hematotoxicity induced by tetrachloroethylene in egyptian dry cleaning workers

Immunotoxicity and hepatic function evaluation in Nile tilapia (*Oreochromis niloticus*) exposed to diazinon

Immunotoxicity and oxidative stress in the Arctic scallop *Chlamys islandica*: effects of acute oil exposure

Immunotoxicity and pulmonary toxicity induced by paints in Egyptian painters

Immunotoxicity and sensitizing capacity of metal compounds depend on speciation

Immunotoxicity and transcriptome analysis of zebrafish embryos in response to glufosinate-ammonium exposure



Immunotoxicity assessment of CdSe/ZnS quantum dots in macrophages, lymphocytes and BALB/c mice

Immunotoxicity assessment of gentamycin and liquamycin

Immunotoxicity assessment of rice-derived recombinant human serum albumin using human peripheral blood mononuclear cells

Immunotoxicity assessment of sub-chronic oral administration of acetamiprid in Wistar rats

Immunotoxicity Considerations for Next Generation Cancer Nanomedicines

Immunotoxicity derived from manipulating leukocytes with lipid-based nanoparticles

Immunotoxicity effects of carbaryl in vivo and in vitro

Immunotoxicity evaluation by subchronic oral administration of clothianidin in Sprague-Dawley rats

Immunotoxicity evaluation of novel bioactive composites in male mice as promising orthopaedic implants

Immunotoxicity Following Pre- and Post-natal Aluminum Exposure in Rats

Immunotoxicity from checkpoint inhibitor therapy: clinical features and underlying mechanisms

Immunotoxicity in ascidians: antifouling compounds alternative to organotin-IV. The case of zinc pyrithione

Immunotoxicity in ascidians: antifouling compounds alternative to organotin: III--the case of copper(I) and Irgarol 1051

Immunotoxicity in ascidians: antifouling compounds alternative to organotin - II. The case of Diuron and TCMS pyridine

Immunotoxicity in C57BL/6 mice exposed to benzene and Aroclor 1254

Immunotoxicity in channel catfish, *Ictalurus punctatus*, following acute exposure to tributyltin

Immunotoxicity in green mussels under perfluoroalkyl substance (PFAS) exposure: Reversible response and response model development

Immunotoxicity in invertebrates: measurement and ecotoxicological relevance

Immunotoxicity in mice induced by short-term exposure to methoxychlor, parathion, or piperonyl butoxide

Immunotoxicity induced by acute subtoxic doses of paraquat herbicide: implication of shifting cytokine gene expression toward T-helper (T(H))-17 phenotype

Immunotoxicity induced in mice by subacute exposure to berberine

Immunotoxicity monitoring of hospital staff occupationally exposed to cytostatic drugs

Immunotoxicity of 180 day exposure to polydimethylsiloxane (silicone) fluid, gel and elastomer and polyurethane disks in female B6C3F1 mice

Immunotoxicity of 1-methyl-3-octylimidazolium bromide on brocarded carp (*Cyprinus carpio* L

Immunotoxicity of 2',3'-dideoxyinosine in female B6C3F1 mice

Immunotoxicity of 2,4-diaminotoluene in female B6C3F1 mice

Immunotoxicity of 2-methoxyethanol following oral administration in Fischer 344 rats

Immunotoxicity of 3 chemical forms of beryllium following inhalation exposure

Immunotoxicity of a reconstituted polynuclear aromatic hydrocarbon mixture in B6C3F1 mice

Immunotoxicity of a standardized citrus polymethoxylated flavone extract

Immunotoxicity of acrylamide in female BALB/c mice

Immunotoxicity of acute acephate exposure in control or IL-1-challenged rats: correlation between the immune cell composition and corticosteroid concentration in blood

Immunotoxicity of aflatoxin B1 in rats: effects on lymphocytes and the inflammatory response in a chronic intermittent dosing study

Immunotoxicity of aflatoxin B1: impairment of the cell-mediated response to vaccine antigen and modulation of cytokine expression

Immunotoxicity of aflatoxin M1 : as a potent suppressor of innate and acquired immune systems in a subacute study

Immunotoxicity of alcohol in young and old mice. I. In vitro suppressive effects of ethanol on the activities of T and B immune cells of aging mice

Immunotoxicity of alcohol in young and old mice. II. Impaired T cell proliferation and T cell-dependent antibody responses of young and old mice fed ethanol-containing liquid diet

Immunotoxicity of aluminum

Immunotoxicity of aluminum chloride

Immunotoxicity of aminocarb. III. Exposure route-dependent immunomodulation by aminocarb in mice

Immunotoxicity of atrazine in Balb/c mice

Immunotoxicity of AZT: inhibitory effect on thymocyte differentiation and peripheral T cell responsiveness to gp120 of human immunodeficiency virus

Immunotoxicity of benzodiazepines. Influence on contact hypersensitivity to picryl chloride in mice

Immunotoxicity of Benzothiazole on *Mytilus Edulis* Following In Vitro Exposure of Hemocytes

Immunotoxicity of beryllium

Immunotoxicity of beta-Diketone Antibiotic Mixtures to Zebrafish (*Danio rerio*) by Transcriptome Analysis

Immunotoxicity of bis(tri-n-butyltin) oxide in the rat

Immunotoxicity of bis(tri-n-butyltin)oxide in the rat: effects on thymus-dependent immunity and on nonspecific resistance following long-term exposure in young versus aged rats

Immunotoxicity of bisphenol A to *Carassius auratus* lymphocytes and macrophages following in vitro exposure

Immunotoxicity of bisphenol S and F are similar to that of bisphenol A during zebrafish early development

Immunotoxicity of carbaryl in chicken

Immunotoxicity of carbon black nanoparticles to blue mussel hemocytes

Immunotoxicity of clonazepam in adult albino rats

Immunotoxicity of cobalt and nickel--experimental study on cytotoxicity of immuno-sensitive metals

Immunotoxicity of cocaethylene

Immunotoxicity of cocaine and crack

Immunotoxicity of commercial-mixed glyphosate in broad snouted caiman (*Caiman latirostris*)

Immunotoxicity of copper alginate fibers in guinea pigs and mice

Immunotoxicity of copper nanoparticle and copper sulfate in a common Indian earthworm

Immunotoxicity of cupravit and previcur fungicides in mice

Immunotoxicity of environmentally relevant concentrations of butyltins on human natural killer cells in vitro

Immunotoxicity of epicutaneously applied anticoagulant rodenticide warfarin: evaluation by contact hypersensitivity to DNCB in rats

Immunotoxicity of estrogen and nonylphenol on apoptosis and expression of ERs in goldfish macrophage: Opening new avenue for discovering the role of experimental model systems and sexes

Immunotoxicity of ethyl carbamate in female BALB/c mice: role of esterase and cytochrome P450

Immunotoxicity of ethyl-4-isothiocyanatobutanoate in male Wistar rats

Immunotoxicity of explosives-contaminated soil before and after bioremediation

Immunotoxicity of four nanoparticles to a marine bivalve species, *Tegillarca granosa*

Immunotoxicity of gallium arsenide on antigen presentation: comparative study of intratracheal and intraperitoneal exposure routes

Immunotoxicity of heavy metals in relation to Great Lakes

Immunotoxicity of hexachlorobenzene

Immunotoxicity of hydrocortisone on Th1/Th2-related cytokine production is associated with yang-deficient state in traditional Chinese medicine

Immunotoxicity of in vitro vanadium exposures: effects on interleukin-1, tumor necrosis factor-alpha, and prostaglandin E2 production by WEHI-3 macrophages

Immunotoxicity of low level cadmium exposure in fish: an alternative animal model for immunotoxicological studies

Immunotoxicity of mercury: Pathological and toxicological effects

Immunotoxicity of microplastics and two persistent organic pollutants alone or in combination to a bivalve species

Immunotoxicity of monoclonal antibodies

Immunotoxicity of mono-nitrotoluenes in female B6C3F1 mice: I. Para-nitrotoluene

Immunotoxicity of mono-nitrotoluenes in female B6C3F1 mice: II. Meta-nitrotoluene

Immunotoxicity of mycotoxins

Immunotoxicity of N,N-diethylaniline in mice: effect on natural killer activity, cytotoxic T lymphocyte activity, lymphocyte proliferation response and cellular components of the spleen

Immunotoxicity of nanoparticle nTiO2 to a commercial marine bivalve species, Tegillarca granosa

Immunotoxicity of nanoparticles

Immunotoxicity of nanoparticles: a computational study suggests that CNTs and C60 fullerenes might be recognized as pathogens by Toll-like receptors

Immunotoxicity of nitrobenzene in female B6C3F1 mice

Immunotoxicity of nivalenol after subchronic dietary exposure to rats

Immunotoxicity of nucleic acid reduced BioProtein--a bacterial derived single cell protein--in Wistar rats

Immunotoxicity of ochratoxin A and aflatoxin B1 in combination is associated with the nuclear factor kappa B signaling pathway in 3D4/21 cells

Immunotoxicity of ochratoxin A to growing gilts

Immunotoxicity of organophosphate flame retardants TPHP and TDCIPP on murine dendritic cells invitro

Immunotoxicity of organophosphorous pesticides

Immunotoxicity of paraquat after subacute exposure to mice

Immunotoxicity of particulate lead: in vitro exposure alters pulmonary macrophage tumor necrosis factor production and activity

Immunotoxicity of PCBs (Aroclors) in relation to Great Lakes

Immunotoxicity of pentachlorophenol (PCP): increased susceptibility to tumor growth in adult mice fed technical PCP-contaminated diets

Immunotoxicity of pentachlorophenol on macrophage immunity and IgM secretion of the crucian carp (*Carassius auratus*)

Immunotoxicity of perfluorinated alkylates: calculation of benchmark doses based on serum concentrations in children

Immunotoxicity of perfluorinated compounds: recent developments

Immunotoxicity of perfluorooctanoic acid and perfluorooctane sulfonate and the role of peroxisome proliferator-activated receptor alpha

Immunotoxicity of pesticides

Immunotoxicity of pesticides: perspectives and trends

Immunotoxicity of phosphamidon following subchronic exposure in albino rats

Immunotoxicity of poly (lactic-co-glycolic acid) nanoparticles: influence of surface properties on dendritic cell activation

Immunotoxicity of polychlorinated biphenyls (PCB) in free-ranging gray seal pups with special emphasis on dioxin-like congeners

Immunotoxicity of polychlorinated biphenyls: present status and future considerations

Immunotoxicity of polyunsaturated fatty acids in serum-free medium

Immunotoxicity of pyrethroid metabolites in an in vitro model

Immunotoxicity of repeated low level exposure to T-2 toxin, a trichothecene mycotoxin, in CD-1 mice

Immunotoxicity of silicon dioxide nanoparticles with different sizes and electrostatic charge

Immunotoxicity of silicone: implications of oxidant balance towards adjuvant activity

Immunotoxicity of Silver Nanoparticles (AgNPs) on the Leukocytes of Common Bottlenose Dolphins (*Tursiops truncatus*)

Immunotoxicity of silver nanoparticles in an intravenous 28-day repeated-dose toxicity study in rats

Immunotoxicity of skin acid secretion produced by the sea slug *Berthellina citrina* in mice spleen: Histological and Immunohistochemical study

Immunotoxicity of sodium bromate in female B6C3F1 mice: a 28-day drinking water study

Immunotoxicity of soluble and insoluble salts of cadmium instilled intratracheally

Immunotoxicity of soluble beta-glucans induced by indomethacin treatment

Immunotoxicity of subchronic versus chronic exposure to aldicarb in mice

Immunotoxicity of sulfuric acid aerosol: effects on pulmonary macrophage effector and functional activities critical for maintaining host resistance against infectious diseases

Immunotoxicity of surface waters contaminated by municipal effluents to the snail *Lymnaea stagnalis*

Immunotoxicity of tannery effluent to the freshwater fish *Cyprinus carpio*

Immunotoxicity of the anticancer drug CI-994 in rats: effects on lymphoid tissue

Immunotoxicity of the colour additive caramel colour III; a review on complicated issues in the safety evaluation of a food additive

Immunotoxicity of the commercial polybrominated diphenyl ether mixture DE-71 in ranch mink (*Mustela vison*)

Immunotoxicity of the organochlorine pesticide methoxychlor in female ICR, BALB/c, and C3H/He mice

Immunotoxicity of the pyrethroid insecticides deltamethrin and alpha-cypermethrin

Immunotoxicity of the pyrrolizidine alkaloid monocrotaline following subchronic administration to C57Bl/6 mice

Immunotoxicity of the semiconductor gallium arsenide in female B6C3F1 mice

Immunotoxicity of the xenoestrogen 4-nonylphenol to the cockle *Cerastoderma glaucum*

Immunotoxicity of titanium dioxide nanoparticles via simultaneous induction of apoptosis and multiple toll-like receptors signaling through ROS-dependent SAPK/JNK and p38 MAPK activation

Immunotoxicity of trenbolone acetate in Japanese quail

Immunotoxicity of tributyltin oxide in rats exposed as adults or pre-weanlings

Immunotoxicity of trichloroethylene: a study with MRL-lpr/lpr mice

Immunotoxicity of tri-n-butyltin oxide (TBTO) and tri-n-butyltin chloride (TBTC) in the rat

Immunotoxicity of washing soda in a freshwater sponge of India

Immunotoxicity of zearalenone in Balb/c mice in a high subchronic dosing study counteracted by *Raphanus sativus* extract

Immunotoxicity of zinc oxide nanoparticles with different size and electrostatic charge

Immunotoxicity risks associated with land-treatment of petrochemical wastes revealed using an in situ rodent model

Immunotoxicity studies in mink (*Mustela vison*) chronically exposed to dietary bleached kraft pulp mill effluent

Immunotoxicity studies of PCB (Aroclor 1254) in the adult rhesus (*Macaca mulatta*) monkey--preliminary report

Immunotoxicity study of repeated small doses of dimethoate and methylparathion administered to rats over three generations

Immunotoxicity testing of mycotoxins T-2 and patulin on Balb/c mice

Immunotoxicity, genotoxicity and epigenetic toxicity of nanomaterials: New strategies for toxicity testing

Immunotoxicity: the risk is real

Immunotoxicologic effects of 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD) in rabbits

Immunotoxicologic effects of cyclosporine on tumor progression in models of squamous cell carcinoma and B-cell lymphoma in C3H mice

Immunotoxicologic effects of ethylene dibromide in the mouse and their modulation by the estrous cycle

Immunotoxicologic effects of inhaled chromium: role of particle solubility and co-exposure to ozone

Immunotoxicologic effects of polychlorinated biphenyls on the cell-mediated and humoral immune systems

Immunotoxicologic studies with vinyl chloride in rabbits and mice

Immunotoxicological and biochemical effects of aflatoxins in rats prevented by Tunisian montmorillonite with reference to HSCAS

Immunotoxicological and neurotoxicological profile of health effects following subacute exposure to geogenic dust from sand dunes at the Nellis Dunes Recreation Area, Las Vegas, NV

Immunotoxicological assessment of cyclosporin A by conventional pathological techniques and immune function testing in the rat

Immunotoxicological characteristics of sodium methylthiocarbamate

Immunotoxicological consequences of perinatal chemical exposures

Immunotoxicological consequences of perinatal chemical exposures: a plea for inclusion of immune parameters in reproduction studies

Immunotoxicological effects of a sub-chronic exposure to selected current-use pesticides in rainbow trout (*Oncorhynchus mykiss*)

Immunotoxicological effects of Agent Orange exposure to the Vietnam War Korean veterans

Immunotoxicological effects of an activated-sludge-treated effluent on rainbow trout (*Oncorhynchus mykiss*)

Immunotoxicological Effects of Aripiprazole: In vivo and In vitro Studies

Immunotoxicological effects of arsenic bioaccumulation on spatial metallomics and cellular enzyme response in the spleen of male Wistar rats after oral intake

Immunotoxicological effects of cadmium on *Labeo rohita*, with emphasis on the expression of HSP genes

Immunotoxicological effects of dermal application of scum of waste crankcase oil in mice

Immunotoxicological effects of environmental contaminants on marine bivalves

Immunotoxicological effects of inorganic arsenic on gilthead seabream (*Sparus aurata* L

Immunotoxicological effects of JP-8 jet fuel exposure

Immunotoxicological effects of piperine in mice

Immunotoxicological effects of repeated combined exposure by cypermethrin and the heavy metals lead and cadmium in rats

Immunotoxicological effects of streptozotocin and alloxan: in vitro and in vivo studies

Immunotoxicological effects on piglets of feeding sows diets containing aflatoxins

Immunotoxicological evaluation of corn genetically modified with *Bacillus thuringiensis* Cry1Ah gene by a 30-day feeding study in BALB/c mice

Immunotoxicological evaluation of toluene exposure via drinking water in mice

Immunotoxicological examination of repeated dose combined exposure by dimethoate and two heavy metals in rats

Immunotoxicological impact and biodistribution assessment of bismuth selenide (Bi<sub>2</sub>Se<sub>3</sub>) nanoparticles following intratracheal instillation in mice

Immunotoxicological impact of engineered nanomaterial exposure: mechanisms of immune cell modulation

Immunotoxicological impact of occupational and environmental nanoparticles exposure: The influence of physical, chemical, and combined characteristics of the particles

Immunotoxicological investigation in rats dosed repeatedly with combinations of cypermethrin, As(III), and Hg(II)

Immunotoxicological investigation of SCMF, a new pyrethroid pesticide in mice

Immunotoxicological investigation of subacute combined exposure by permethrin and the heavy metals arsenic(III) and mercury(II) in rats

Immunotoxicological investigation of subacute combined exposure with low doses of Pb, Hg and Cd in rats

Immunotoxicological investigation of the effects of a pesticide; cypermethrin

Immunotoxicological investigation using pharmaceutical drugs. In vitro evaluation of immune effects using rodent or human immune cells

Immunotoxicological investigations on rats treated subcutely with dimethoate, As<sup>3+</sup> and Hg<sup>2+</sup> in combination

Immunotoxicological profile of morphine sulfate in B6C3F1 female mice

Immunotoxicological properties of airborne particles at landfill, urban and rural sites and their relation to microbial concentrations



Immunotoxicological response of the earthworm *Lumbricus terrestris* following exposure to cement kiln dusts

Immunotoxicological risk of mycotoxins for domestic animals

Immunotoxicological screening of morphine and methadone in an extended 28 day study in rats

Immunotoxicological study of one of the most common over-the-counter pyrethroid insecticide products in Egypt

Immunotoxicological, biochemical, and histopathological studies on Roundup and Stomp herbicides in Nile catfish (*Clarias gariepinus*)

Immunotoxicology and its application in risk assessment

Immunotoxicology assessment in the pharmaceutical industry

Immunotoxicology in occupational and environmental medicine: perspectives, limitations, and research objectives

Immunotoxicology in wood mice along a heavy metal pollution gradient

Immunotoxicology of arc welding fume: worker and experimental animal studies

Immunotoxicology of beryllium lung disease

Immunotoxicology of brown recluse spider (*Loxosceles reclusa*) venom

Immunotoxicology of cadmium

Immunotoxicology of cadmium and mercury on B-lymphocytes--I. Effects on lymphocyte function

Immunotoxicology of cigarette smoke condensates: suppression of macrophage responsiveness to interferon gamma

Immunotoxicology of heavy metals

Immunotoxicology of non-functionalized engineered nanoparticles in aquatic organisms with special emphasis on fish--review of current knowledge, gap identification, and call for further research

Immunotoxicology of organic acid anhydrides (OAAs)

Immunotoxicology of silica

Immunotoxicology studies of sodium arsenate-effects of exposure on tumor growth and cell-mediated tumor immunity

Immunotoxicology studies on lead: effects of exposure on tumor growth and cell-mediated tumor immunity after syngeneic or allogeneic stimulation

Immunotoxicology studies on octoxynol-9 and nonoxynol-9 in mice

Immunotoxicology: environmental contamination by polybrominated biphenyls and immune dysfunction among residents of the State of Michigan

Immunotoxicology: role of inflammation in chemical-induced hepatotoxicity

Immunotoxins constructed with chimeric, short-lived anti-CD22 monoclonal antibodies induce less vascular leak without loss of cytotoxicity

Impact of aluminum exposure on the immune system: a mini review

Impact of an immunosuppressive human pharmaceutical on the interaction of a bacterial parasite and its invertebrate host

Impact of dexamethasone-induced immunosuppression on the duration and level of shedding of *Escherichia coli* O157:H7 in calves

Impact of dietary *Trichosporon* mycotoxinivorans on ochratoxin A induced immunotoxicity; In vivo study

Impact of exposure time, particle size and uptake pathway on silver nanoparticle effects on circulating immune cells in *Mytilus galloprovincialis*

Impact of in ovo-administered lead and testosterone on developing female thymocytes

Impact of in vitro gallium arsenide exposure on macrophages

Impact of maternal HIV exposure, feeding status, and microbiome on infant cellular immunity

Impact of nitrite exposure on plasma biochemical parameters and immune-related responses in *Takifugu rubripes*

Impact of PBDE-209 exposure during pregnancy and lactation on immune function of offspring rats

Impact of perfluorooctanesulfonate and perfluorooctanoic acid on human peripheral leukocytes

Impact of radiofrequency radiation on DNA damage and antioxidants in peripheral blood lymphocytes of humans residing in the vicinity of mobile phone base stations.

Impacts of simultaneous exposure to arsenic (III) and copper (II) on inflammatory response, immune homeostasis, and heat shock response in chicken thymus

Impaired Antibody-Independent Immune Response of B Cells in Patients With Acute Dengue Infection

Impaired B cell immunity in acute myeloid leukemia patients after chemotherapy

Impaired CD8(+) T cell immunity after allogeneic bone marrow transplantation leads to persistent and severe respiratory viral infection

Impaired cell-mediated immunity in the first week after burn injury: investigation of spontaneous blastogenic transformation, PHA, IL-2 response and plasma suppressive activity

Impaired cellular immune response to diphtheria and tetanus vaccines in children after thoracic transplantation

Impaired cellular immune response to injected bacteria after knockdown of ferritin genes in the hard tick *Haemaphysalis longicornis*

Impaired Cytomegalovirus Immunity in Idiopathic Pulmonary Fibrosis Lung Transplant Recipients with Short Telomeres

Impaired humoral and cellular immune responses to influenza vaccination in chronic obstructive pulmonary disease patients

Impaired immune defense in hemodialysis patients: role of alpha-defensins

Impaired immune function and structural integrity in the gills of common carp (*Cyprinus carpio* L.) caused by chlorpyrifos exposure: Through oxidative stress and apoptosis

Impaired Immune Function in Patients Undergoing Surgery for Bone Cancer

Impaired immune regulation after radioiodine therapy for Graves' disease and the protective effect of Methimazole

Impaired Immune Response in Elderly Burn Patients: New Insights Into the Immune-senescence Phenotype

Impaired Immune Response in Old Mice Suffering from Obesity and Premature Immunosenescence in Adulthood

Impaired immune response in severe human lower tract respiratory infection by respiratory syncytial virus

Impaired immune response to vaccination against infection with human respiratory syncytial virus at advanced age

Impaired immune responses following spinal cord injury lead to reduced ability to control viral infection

Impaired immune responses in streptozotocin-induced type I diabetes in mice. Involvement of high glucose

Impaired immunity in obesity: suppressed but reversible lymphocyte responsiveness

Impaired immunity: risk groups and consequences for general practice

Impaired immunocompetence associated with iron deficiency

Impaired immunoglobulin M production by incubation of hybridoma cells with ethanol

Impaired immunosuppressive response to ultraviolet radiation in interleukin-10-deficient mice

Impaired innate immune alveolar macrophage response and the predilection for COPD exacerbations

Impaired innate immunity of ocular surface is the key bridge between extended contact lens wearing and occurrence of *Acanthamoeba* keratitis

Impaired innate mucosal immunity in aged mice permits prolonged *Streptococcus pneumoniae* colonization

Impaired intestinal immune barrier and physical barrier function by phosphorus deficiency: Regulation of TOR, NF-kappaB, MLCK, JNK and Nrf2 signalling in grass carp (*Ctenopharyngodon idella*) after infection with *Aeromonas hydrophila*

Impaired intestinal immunity and barrier function: a cause for enhanced bacterial translocation in alcohol intoxication and burn injury

Impaired intestinal mucosal immunity in specific-pathogen-free chickens after infection with very virulent infectious bursal disease virus

Impaired long-term immune protection following pneumococcal 13-valent/23-valent polysaccharide vaccine in systemic lupus erythematosus (SLE)

Impaired Microcirculation in Children After Kidney Transplantation: Everolimus Versus Mycophenolate Based Immunosuppression Regimen

Impaired mononuclear cell immune function in extreme obesity is corrected by weight loss

Impaired non-viral specific immune function of dendritic cell does not interfere with clearance and cytotoxic T lymphocyte response to HBV or HCV

Impaired outcome of continuous ambulatory peritoneal dialysis in immunosuppressed patients

Impaired phosphate handling of renal allografts is aggravated under rapamycin-based immunosuppression

Impaired pulmonary immunity post-bone marrow transplant

Impaired responses to gliadin and gut microbes of immune cells from mice with altered stress-related behavior and premature immune senescence

Impaired serotype-specific immune function following pneumococcal vaccination in infants with prior carriage

Impairment of cell-mediated immune responses by *Pseudomonas aeruginosa*

Impairment of Immunonutritional Status During Treatment is a Factor Associated With Oncologic Outcomes in Patients With Rectal Cancer Treated With Preoperative Chemoradiotherapy

Impairment of Immunoproteasome Function by Cigarette Smoke and in Chronic Obstructive Pulmonary Disease

Impairment of innate immune function by hydroxyethyl starch

Impairment of innate immune killing mechanisms by bacteriostatic antibiotics

Impairment of innate immune responses of airway epithelium by infection with bovine viral diarrhea virus

Impairment of non-specific immunity in patients under persistent vegetative state resulting from trauma

Impairment of renal function after islet transplant alone or islet-after-kidney transplantation using a sirolimus/tacrolimus-based immunosuppressive regimen

Impairment of several immune functions in anxious women

Impairment of T cell immunity by the respiratory syncytial virus: targeting virulence mechanisms for therapy and prophylaxis

Impairment of the immune response after transcuticular introduction of the insect gonadoinhibitory and hemocytotoxic peptide Neb-colloostatin: A nanotech approach for pest control

Impairment of the immune system in GH-overexpressing transgenic zebrafish (*Danio rerio*)

Implementation of toxicokinetics in toxicity studies--Toxicokinetics of 4-methylanisole and its metabolites in juvenile and adult rats

Implications of the new FDA/CDER immunotoxicology guidance for drugs

Implications of UV-induced inflammation and immunomodulation

Importance of innate immunity and collagen binding integrin  $\alpha 1\beta 1$  in TNBS-induced colitis

In altering the release of glucocorticoids, ketorolac exacerbates the effects of systemic immune stimuli on expression of proinflammatory genes in the brain

In contrast to morphine, buprenorphine enhances macrophage-induced humoral immunity and, as oxycodone, slightly suppresses the effector phase of cell-mediated immune response in mice

In situ (mesocosm) assessment of immunotoxicity risks to small mammals inhabiting petrochemical waste sites

In utero arsenic exposure is associated with impaired thymic function in newborns possibly via oxidative stress and apoptosis

In utero exposure to arsenic alters lung development and genes related to immune and mucociliary function in mice

In Utero exposure to genistein enhanced intranasal house dust mite allergen-induced respiratory sensitization in young adult B6C3F1 mice

In utero exposure to immunosuppressive drugs

In utero exposure to immunosuppressive drugs: experimental and clinical studies

In utero ultrafine particulate matter exposure causes offspring pulmonary immunosuppression

In vitro analysis of verapamil-induced immunosuppression: potent inhibition of T cell motility and lymphocytic transmigration through allogeneic endothelial cells

In vitro and in vivo comparison of the immunotoxicity of single- and multi-layered graphene oxides with or without pluronic F-127

In vitro and in vivo immunotoxic and immunomodulatory effects of nonsupplemented and selenium-supplemented cigarette smoke condensate

In vitro assays supporting the safety assessment of immunomodulatory monoclonal antibodies

In vitro characterization of cutaneous immunotoxicity of immortalized human keratinocytes (HaCaT) exposed to reactive and disperse textile dyes

In vitro characterization of the immunotoxic potential of several perfluorinated compounds (PFCs)

In vitro cytokine release from human peripheral blood mononuclear cells in the assessment of the immunotoxic potential of chemicals

In vitro cytopathic effects of mycotoxin T-2 on human peripheral blood T lymphocytes

In vitro effect of 4-pentylphenol and 3-methyl-4-nitrophenol on murine splenic lymphocyte populations and cytokine/granzyme production

In vitro enhancement of mouse T helper 2 cell sensitization to ovalbumin allergen by carbon black nanoparticles

In vitro evaluation of the effects of perfluorooctanesulfonic acid (PFOS) and perfluorooctanoic acid (PFOA) on IL-2 production in human T-cells

In vitro evaluation of the immunotoxic potential of perfluorinated compounds (PFCs)

In Vitro Exposure of Harbor Seal Immune Cells to Aroclor 1260 Alters Phocine Distemper Virus Replication

In vitro hematological and in vivo immunotoxicity assessment of dextran stabilized iron oxide nanoparticles

In vitro immune toxicity of polybrominated diphenyl ethers on murine peritoneal macrophages: apoptosis and immune cell dysfunction

In vitro immunotoxicity and cytotoxicity of trichosanthin against human normal immunocytes and leukemia-lymphoma cells

In vitro immunotoxicity of bis(tri-n-butyltin)oxide (TBTO) studied by toxicogenomics

In vitro immunotoxicity of untreated and treated urban wastewaters using various treatment processes to rainbow trout leucocytes

In vitro immunotoxicological effects of heavy metals on European sea bass (*Dicentrarchus labrax* L.) head-kidney leucocytes

In vitro immunotoxicology of quantum dots and comparison with dissolved cadmium and tellurium

In vitro lymphotoxicity and selective T cell immunotoxicity of high doses of acyclovir and its derivatives in mice

In vitro Modulating Activity of aqueous extracts from American Plants on Chlorpyrifos-induced toxicity on Murine Splenocytes

In vitro nanoparticle toxicity to rat alveolar cells and coelomocytes from the earthworm *Lumbricus rubellus*

In vitro nephrotoxicity induced by chloronitrobenzenes in renal cortical slices from Fischer 344 rats

In vitro nephrotoxicity induced by propanil

In vitro potency of inhibition by antiviral drugs of hematopoietic progenitor colony formation correlates with exposure at hemotoxic levels in human immunodeficiency virus-positive humans

In vitro stimulatory effect of N-acetyl tryptophan-glucopyranoside against gamma radiation induced immunosuppression

In vitro studies on immunotoxic potential of Orange II in splenocytes

In vitro study on the immune effects of the exposure to palladium nanoparticles

In vitro tests to evaluate immunotoxicity: a preliminary study

In vitro toxicity and interactions of environmental contaminants (Arochlor 1254 and mercury) and immunomodulatory agents (lipopolysaccharide and cortisol) on thymocytes from lake trout (*Salvelinus namaycush*)

In vitro VLA-4 blockade results in an impaired NK cell-mediated immune surveillance against melanoma

In vitro" comparative immune effects of different titanium compounds

In vitro" effects of different arsenic compounds on PBMC (preliminary study)

In vivo activation of a T helper 2-driven innate immune response in lung fibrosis induced by multi-walled carbon nanotubes

In vivo and in vitro immunosuppressive effects of benzo[k]fluoranthene in female Balb/c mice

In vivo effects of immunostimulating lipopeptides on mouse liver microsomal cytochromes P-450 and on paracetamol-induced toxicity

In vivo immunotoxicity of  $\text{SiO}_2@(\text{Y}_{0.5}\text{Gd}_{0.45}\text{Eu}_{0.05})_2\text{O}_3$  as dual-modality nanoprobes

In vivo induction of apoptosis and immune responses in mice by administration of lipopolysaccharide from *Porphyromonas gingivalis*

In vivo short-term exposure to residual oil fly ash impairs pulmonary innate immune response against environmental mycobacterium infection

In vivo studies on the immunotoxic effects of microcystins on rabbit

In Vivo toxicological assessment of biologically synthesized silver nanoparticles in adult Zebrafish (*Danio rerio*)

Inactivation of cadmium induced immunotoxicological alterations in rats by Tunisian montmorillonite clay

Incidence and etiology of hepatic dysfunction in heart transplant recipients receiving a cyclosporine-based triple immunosuppressive therapy

Incidence and natural history of intravenous immunoglobulin-induced aseptic meningitis: a retrospective review at a single tertiary care center

Incidence of cytomegalovirus in cardiac transplant recipients receiving induction immunosuppression with antithymocyte globulin

Incident stressful and traumatic life events and human immunodeficiency virus sexual transmission risk behaviors in a longitudinal, multisite cohort study

Inconsistently impairment of immune function and structural integrity of head kidney and spleen by vitamin A deficiency in grass carp (*Ctenopharyngodon idella*)

Increase in level of tumor necrosis factor- $\alpha$  in 6-hydroxydopamine-lesioned striatum in rats is suppressed by immunosuppressant FK506

Increased affective ultrasonic communication during fear learning in adult male rats exposed to maternal immune activation

Increased expression of immune modulator proteins and decreased expression of apolipoprotein A-1 and haptoglobin in blood plasma of sarin exposed rats

Increased expression of platelet IgG Fc receptors in immune heparin-induced thrombocytopenia

Increased immune and inflammatory responses to dust mite antigen in rats exposed to 5 ppm NO<sub>2</sub>

Increased immunoreactive rat lung ICAM-1 in oleic acid-induced lung injury

Increased immunoreactivity of POMC-derived neuropeptides and immediate-early gene-derived proteins (c-Fos and Egr-1 proteins) as an early step of acute cocaine-induced stressor effects: comparison with the effects of immobilization stress

Increased levels of glutathione S transferases and appearance of novel alpha class isoenzymes in kidneys of mice exposed to mercuric chloride. I. Biochemical and immunohistochemical studies

Increased levels of numerical chromosome aberrations after in vitro exposure of human peripheral blood lymphocytes to radiofrequency electromagnetic fields for 72 hours.

Increased total and mite-specific immunoglobulin E in patients with aspirin-induced urticaria and angioedema

Indicators of immunotoxicity in populations of cotton rats (*Sigmodon hispidus*) inhabiting an abandoned oil refinery

Individual and combined cytotoxicity of major trichothecenes type B, deoxynivalenol, nivalenol, and fusarenon-X on Jurkat human T cells

Individual responsiveness to induction of micronuclei in human lymphocytes after exposure in vitro to 1800-MHz microwave radiation.

Individual susceptibility to hexavalent chromium of workers of shoe, hide, and leather industries. Immunological pattern of HLA-B8, DR3-positive subjects

Indoxacarb interaction alters immunotoxic and genotoxic potential of endotoxin



Indoxyl sulfate (IS)-mediated immune dysfunction provokes endothelial damage in patients with end-stage renal disease (ESRD)

Induced resistance to *Candida albicans* in BALB/c mice during short-term immunosuppression with cyclophosphamide

Induced sensitization to nickel in guinea pigs immunized with mycobacteria by injection of purified protein derivative with nickel

Inducible headkidney cytochrome P450 contributes to endosulfan immunotoxicity in walking catfish *Clarias gariepinus*

Induction and activation of adaptive immune populations during acute and chronic phases of a murine model of experimental colitis

Induction and Amelioration of Methotrexate-Induced Gastrointestinal Toxicity are Related to Immune Response and Gut Microbiota

Induction and modulation of pulmonary inflammation by organic dusts: cytokines, immune complexes and 'all of those things'

Induction Immunosuppression in High-risk Kidney Transplant Recipients

Induction immunosuppression in kidney transplant recipients older than 60 years of age : safety and efficacy of ATGAM, OKT3 and Simulect

Induction immunosuppression with interleukin-2 receptor antibodies (basiliximab and daclizumab) in renal transplant recipients

Induction immunosuppression with OKT3 monoclonal antibody in cardiac transplant recipients

Induction immunosuppressive therapy in kidney transplantation

Induction immunosuppressive therapy in renal transplantation: does basiliximab make the difference

Induction immunosuppressive therapy in the elderly kidney transplant recipient in the United States

Induction immunosuppressive therapy stratified according to risk categories in renal transplantation

Induction of a type 1 immune response to a recombinant antigen from *Mycobacterium tuberculosis* expressed in *Mycobacterium vaccae*

Induction of alloreactive immunosuppression by 1,4-bis [( 2-aminoethyl)amino]-5,8-dihydroxy-9,10-anthracenedione dihydrochloride (CL 232,468)

Induction of an autologous immune-complex glomerulonephritis in the rat by intravenous injection of heterologous anti-rat kidney tubular antibody IV: Effect of injection of HgCl<sub>2</sub> prior to the antibody

Induction of antiviral cytotoxic T cells by plasmacytoid dendritic cells for adoptive immunotherapy of posttransplant diseases

Induction of apoptotic lesions in liver and lymphoid tissues and modulation of cytokine mRNA expression by acute exposure to deoxynivalenol in piglets

Induction of autoimmunity through bystander effects. Lessons from immunological disorders induced by heavy metals

Induction of cancer, actinic keratosis, and specific p53 mutations by UVB light in human skin maintained in severe combined immunodeficient mice

Induction of colitis in mice with food allergen-specific immune response

Induction of eruptive benign melanocytic naevi by immune suppressive agents, including biologicals

Induction of experimental systemic lupus erythematosus in mice by immunization with a monoclonal anti-La autoantibody

Induction of IDO-1 by immunostimulatory DNA limits severity of experimental colitis

Induction of Immune Reaction in Benign Melanocytic Nevi Without Halo During Nivolumab Therapy in a Patient With Melanoma

Induction of immunomediated diseases by recombinant human granulocyte-macrophage colony-stimulating factor during cancer treatment

Induction of immunosuppression with rabbit antithymocyte globulin: five-year experience in cardiac transplantation

Induction of immunotoxicity by polycyclic hydrocarbons: role of the Ah locus

Induction of immunotoxicity in mice by polyhalogenated biphenyls

Induction of innate immune gene expression following methyl methanesulfonate-induced DNA damage in sea urchins

Induction of Maternal Immune Activation in Mice at Mid-gestation Stage with Viral Mimic Poly(I:C

Induction of multiple melanocytic nevus cell nevi in 2 children with malignant hematologic systemic diseases and chemotherapy-induced immunosuppression

Induction of neutralizing antibodies against Tityus serrulatus toxins by immunization with a recombinant nontoxic protein

Induction of neutralizing antibodies in mice immunized with scorpion toxins detoxified by liposomal entrapment

Induction of oxidative stress by Taxol vehicle Cremophor-EL triggers production of interleukin-8 by peripheral blood mononuclear cells through the mechanism not requiring de novo synthesis of mRNA

Induction of sarcomas by a single subcutaneous injection of 7,12-dimethylbenz[a]anthracene into neonatal male Sprague-Dawley rats: histopathological and immunohistochemical analyses

Induction of serum-borne immunomodulatory factors in B6C3F1 mice by carbon tetrachloride. I. Carbon tetrachloride-induced suppression of helper T-lymphocyte function is mediated by a serum borne factor

Induction of TGF-beta 1, not regulatory T cells, impairs antiviral immunity in the lung following bone marrow transplant

Induction Therapy With ATG Compared With Anti-IL2 Basiliximab in Low-Immunologic Risk Kidney Transplant Recipients

Induction versus noninduction in renal transplant recipients with tacrolimus-based immunosuppression

Industry experience in the identification of the immunotoxic potential of agrochemicals

Ineffectiveness of folic acid supplementation against phenytoin-induced decrease in salivary immunoglobulin A concentration of epileptic patients

Infection with the Lyme disease pathogen suppresses innate immunity in mice with diet-induced obesity

Infection with Vpr-positive human immunodeficiency virus type 1 impairs NK cell function indirectly through cytokine dysregulation of infected target cells

Inflammation caused by peripheral immune cells across into injured mouse blood brain barrier can worsen postoperative cognitive dysfunction induced by isoflurane

Inflammation induced by influenza virus impairs human innate immune control of pneumococcus

Inflammatory and immunological responses to subchronic exposure to endotoxin-contaminated metalworking fluid aerosols in F344 rats

Inflammatory and oxidative stress-related effects associated with neurotoxicity are maintained after exclusively prenatal trichloroethylene exposure

Inflammatory potential in relation to the microbial content of settled dust samples collected from moisture-damaged and reference schools: results of HITEA study

Infliximab in the treatment of anti-CTLA4 antibody (ipilimumab) induced immune-related colitis

Influence of 1.8 GHz microwave on DNA damage induced by 4 chemical mutagens.

Influence of arecoline on immune system: II. Suppression of thymus-dependent immune responses and parameter of non-specific resistance after short-term exposure

Influence of cadmium on murine thymocytes: potentiation of apoptosis and oxidative stress

Influence of chlorpyrifos on the profile of subpopulations of immunoactive cells and their phagocytic activity in an experimental in vivo model

Influence of environmental and industrial immunotoxic hazards on clinical course of HIV-infection

Influence of Human Leukocyte Antigen (HLA) Alleles and Killer Cell Immunoglobulin-Like Receptors (KIR) Types on Heparin-Induced Thrombocytopenia (HIT)

Influence of immunization on the pulmonary inflammatory response of rabbits induced by Pasteurella haemolytica A1 lipopolysaccharide

Influence of immunization with non-genotoxic PAH-KLH conjugates on the resistance of organisms exposed to benzo(a)pyrene

Influence of induction therapy, immunosuppressive regimen and anti-viral prophylaxis on development of lymphomas after heart transplantation: data from the Spanish Post-Heart Transplant Tumour Registry

Influence of life stress on immunological reactivity to mild psychological stress

Influence of low-level exposure to *Fusarium* mycotoxins on selected immunological and hematological parameters in young swine

Influence of melatonin on immunotoxicity of cadmium

Influence of melatonin on immunotoxicity of lead

Influence of perinatal ochratoxin A exposure on the immune system in mice

Influence of stress on DDT-induced humoral immune responsiveness in mice

Influence of subchronic exposure to lindane on humoral immunity in mice

Influence of the Ah locus on the humoral immunotoxicity of 2,3,7,8-tetrachlorodibenzo-p-dioxin: evidence for Ah-receptor-dependent and Ah-receptor-independent mechanisms of immunosuppression

Influenza A virus protein PB1-F2 impairs innate immunity by inducing mitophagy

Influenza A virus protein PB1-F2 translocates into mitochondria via Tom40 channels and impairs innate immunity

Ingestion of low doses of deoxynivalenol does not affect hematological, biochemical, or immune responses of piglets

Inhalation toxicity studies of methyl isocyanate (MIC) in rats: Part IV--Immunologic response of rats one week after exposure: effect on body and organ weights, phagocytic and DTH response

Inhaled isobutyl nitrite compromises T-dependent, but not T-independent, antibody induction

Inhaled particles and respiratory disease

Inhibition of 3,3',4,4',5-pentachlorobiphenyl-induced fetal cleft palate and immunotoxicity in C57BL/6 mice by 2,2',4,4',5,5'-hexachlorobiphenyl

Inhibition of cytochrome P-450 with 2-diethylamino-ethyl-2,2-diphenylpropylacetate (SKF-525A) reduces immunotoxicity of chlorinated carbohydrates

Inhibition of HIV-1-specific T-cells and increase of viral load during immunosuppressive treatment in an HIV-1 infected patient with *Chlamydia trachomatis* induced arthritis

Inhibition of immune responses and related proteins in *Rhamdia quelen* exposed to diclofenac

Inhibition of immune-mediated concanavalin A-induced liver damage by free-radical scavengers

Inhibition of in vitro cytokine production by human peripheral blood mononuclear cells treated with xenobiotics: implications for the prediction of general toxicity and immunotoxicity

Inhibition of LPS-induced splenocyte proliferation by ortho-substituted polychlorinated biphenyl congeners

Inhibition of mite-induced immunoglobulin E synthesis, airway inflammation, and hyperreactivity by herbal medicine STA-1

Inhibition of TC-1 cytokine production, effector cytotoxic T lymphocyte development and alloantibody production by 2,3,7,8-tetrachlorodibenzo-p-dioxin

Inhibition of tumor necrosis factor activity fails to restore 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD)-induced suppression of the antibody response to sheep red blood cells

Inhibition of type II collagen induced arthritis in mice by an immunosuppressive extract of *Tripterygium wilfordii* Hook f

Inhibitory effect of *Platycodon grandiflorum* on T(H)1 and T(H)2 immune responses in a murine model of 2,4-dinitrofluorobenzene-induced atopic dermatitis-like skin lesions

Inhibitory effect of the immunosuppressant FK506 on apoptotic cell death induced by HIV-1 gp120

Initial experience of combined immunosuppressive induction therapy with polyclonal antithymocyte antibody, FK 506 (tacrolimus), and prednisolone in clinical liver transplantation

Initial immunosuppression in liver transplant recipients with impaired renal function

Innate immune parameters and haemolymph protein expression profile to evaluate the immunotoxicity of tributyltin on abalone (*Haliotis diversicolor supertexta*)

Innate immune processes are sufficient for driving cigarette smoke-induced inflammation in mice

Innate immune response of silver catfish (*Rhamdia quelen*) exposed to atrazine

Innate immunity conferred by Toll-like receptors 2 and 4 and myeloid differentiation factor 88 expression is pivotal to monosodium urate monohydrate crystal-induced inflammation

Innate immunity drives xenobiotic-induced murine autoimmune cholangitis

Innate Lymphoid Cells Mediate Pulmonary Eosinophilic Inflammation, Airway Mucous Cell Metaplasia, and Type 2 Immunity in Mice Exposed to Ozone

Inorganic and methylated arsenic compounds induce cell death in murine macrophages via different mechanisms

Inorganic arsenic causes apoptosis cell death and immunotoxicity on European sea bass (*Dicentrarchus labrax*)

Insights into the heavy metal-induced immunotoxic and genotoxic alterations as health indicators of *Clarias gariepinus* inhabiting a rivulet

Integration of the human lymphocyte into immunotoxicological investigations

Integrative assessment of enantioselectivity in endocrine disruption and immunotoxicity of synthetic pyrethroids

Intensive postgrafting immune suppression combined with nonmyeloablative conditioning for transplantation of HLA-identical hematopoietic cell grafts: results of a pilot study for treatment of primary immunodeficiency disorders

Interaction of aflatoxin B1 and fumonisin B1 in mice causes immunotoxicity and oxidative stress: Possible protective role using lactic acid bacteria

Interaction of *Lactobacillus plantarum* MON03 with Tunisian montmorillonite clay and ability of the composite to immobilize Zearalenone in vitro and counteract immunotoxicity in vivo

Interactions of malnutrition and immune impairment, with specific reference to immunity against parasites

Interactions of nanomaterials with the immune system

Interferon-gamma and immunoglobulin enhance mineral dust-induced production of reactive oxygen metabolites by human macrophages

Interleukin-1 receptor signaling is required to overcome the effects of pertussis toxin and for efficient infection- or vaccination-induced immunity against *Bordetella pertussis*

Interleukin-10 promoter microsatellite polymorphisms influence the immune response to heparin and the risk of heparin-induced thrombocytopenia

Interleukin-2 receptor antibody (basiliximab) for immunosuppressive induction therapy after liver transplantation: a protocol with early elimination of steroids and reduction of tacrolimus dosage

Interrelationship between splenocyte immunomodulating factors in animals undergoing combined exposure to ethanol and tetrachloromethane

Interruption of immune responses in primary macrophages exposed to nonylphenol provides insights into the role of ER and NF- $\kappa$ B in immunotoxicity of Persian sturgeon

Intervention of Grape Seed Proanthocyanidin Extract on the Subchronic Immune Injury in Mice Induced by Aflatoxin B1

Intestinal T lymphocytes of different rat strains in immunotoxicity

Intimate male partner violence impairs immune control over herpes simplex virus type 1 in physically and psychologically abused women

Intracellular human immunodeficiency virus Tat expression in astrocytes promotes astrocyte survival but induces potent neurotoxicity at distant sites via axonal transport

Intracellular immunoglobulins in plasma cells of nasal biopsies taken from vanadium-exposed workers. A retrospective case control study by the peroxidase-antiperoxidase (PAP) method

Intracellular zinc stores protect the intestinal epithelium from Ochratoxin A toxicity

Intraoperative anticoagulation and limb amputations in patients with immune heparin-induced thrombocytopenia who require vascular surgery

Intraoperative anti-thymocyte globulin-Fresenius (ATG-F) administration as induction immunosuppressive therapy in kidney transplantation

Intrauterine cocaine exposure of rabbits: persistent elevation of GABA-immunoreactive neurons in anterior cingulate cortex but not visual cortex

Intravenous cyclosporine to induce immunosuppression in cardiac allograft recipients

Intravenous Immune Globulin for Statin-Triggered Autoimmune Myopathy

Intravenous immunoglobulin as an adjunct therapy in persisting heparin-induced thrombocytopenia

Intravenous immunoglobulin as prophylaxis of chemotherapy-induced oral mucositis

Intravenous immunoglobulin followed by platelet transfusion in the acute treatment of trimethoprim/sulfamethoxazole-induced immune thrombocytopenia

Intravenous immunoglobulin G use in patients with chronic inflammatory demyelinating polyneuropathy: An uncommon cause of drug-induced discoid lupus erythematosus

Intravenous immunoglobulin in the treatment of severe methotrexate-induced acral erythema

Intravenous immunoglobulin induced meningoencephalitis

Intravenous immunoglobulin induced-nephropathy: a complication of IVIG therapy

Intravenous immunoglobulin preparation attenuates LPS-induced production of pro-inflammatory cytokines in human monocytic cells by modulating TLR4-mediated signaling pathways

Intravenous immunoglobulin-induced acute thrombocytopenia

Intravenous immunoglobulin-induced haemolysis: a case report and review of the literature

Intravenous immunoglobulin-induced hemolytic anemia after thoracoscopic thymectomy for myasthenia gravis

Intravenous immunoglobulin-induced hemolytic anemia in a patient with juvenile dermatomyositis

Intravenous immunoglobulin-induced lichenoid dermatitis: a unique adverse reaction

Intravenous immunoglobulin-induced neutropenia

Intravenous immunoglobulin-induced neutrophil apoptosis in the lung during murine endotoxemia

Intravenous immunoglobulin-induced osmotic nephrosis

Intravenous Immunoglobulin-Induced Profound Bradycardia in a Patient With Idiopathic Thrombocytopenic Purpura

Intravenous Immunoglobulin-Induced Pulmonary Embolism: It Is Time to Act

Intravenous immunoglobulin-induced reversible neutropenia in a renal transplant patient

Intravenous immunoglobulin-induced, non-eczematous, vesiculobullous eruptions in Stevens-Johnson syndrome

Intravenous immunoglobulins-induced eczematous eruption: a long-term follow-up study

Investigating the Effects of Particulate Matter on House Dust Mite and Ovalbumin Allergic Airway Inflammation in Mice

Investigation of a screening battery for immunotoxicity of pharmaceuticals within a 28-day oral toxicity study using azathioprine and cyclosporin A as model compounds

Investigation of immunotoxicity of supercypermethrin forte in the Wistar rat

Investigation of neurotoxic and immunotoxic effects of some plant growth regulators at subacute and subchronic applications on rats

Investigation of sensitive biomarkers to determine cadmium inducing hepato- and nephro-toxicity in cattle by immunofluorescence method

Investigation of the adjuvant and immuno-suppressive effects of benzyl butyl phthalate, phthalic acid and benzyl alcohol in a murine injection model

Investigation of the preventive effect of Sijunzi decoction on mitomycin C-induced immunotoxicity in rats by <sup>1</sup>H NMR and MS-based untargeted metabolomic analysis

Investigations into the Immunotoxicity and Allergic Potential Induced by Topical Application of N-Butylbenzenesulfonamide (NBBS) in a Murine Model

Investigations of immunotoxicity and allergic potential induced by topical application of triclosan in mice

In vitro immunotoxicity of environmentally representative antibiotics to the freshwater mussel *Elliptio complanata*

In vivo immunotoxicity of perfluorooctane sulfonate in BALB/c mice: Identification of T-cell receptor and calcium-mediated signaling pathway disruption through gene expression profiling of the spleen

Involvement of altered B7 expression in dioxin immunotoxicity: B7 transfection restores the CTL but not the autoantibody response to the P815 mastocytoma

Involvement of cellular immunity and humoral immunity in mixed allergy induced by trichloroethylene

Involvement of immune-related factors in diclofenac-induced acute liver injury in mice

Involvement of oxidative stress in tri-ortho-cresyl phosphate-induced autophagy of mouse Leydig TM3 cells in vitro

Iodine induced lymphocytic thyroiditis in the BB/W rat: early and late immune phenomena

Ipilimumab-induced Guillain-Barre Syndrome Presenting as Dysautonomia: An Unusual Presentation of a Rare Complication of Immunotherapy

Ipilimumab-induced immune-related renal failure--a case report

Ipomoea obscura ameliorates cyclophosphamide-induced toxicity by modulating the immune system and levels of proinflammatory cytokine and GSH



Iron Deficiency Impairs Intra-Hepatic Lymphocyte Mediated Immune Response

Iron oxide particles modulate the ovalbumin-induced Th2 immune response in mice

Is Chronic Exposure to Low-Dose Organochlorine Pesticides a New Risk Factor of T-cell Immunosenescence

Is hydroxylamine-induced cytotoxicity a valid marker for hypersensitivity reactions to sulfamethoxazole in human immunodeficiency virus-infected individuals

Is impaired immunity a consequence of surgery in patients infected by the human immunodeficiency virus

Is murine gammaherpesvirus-68 (MHV-68) a suitable immunotoxicological model for examining immunomodulatory drug-associated viral recrudescence

Is opioid-induced immunosuppression a clinically relevant problem? Innocent until proven guilty

Is there a human health risk associated with indirect exposure to perfluoroalkyl carboxylates (PFCA)s

Is there a potential immune dysfunction with anabolic androgenic steroid use?: A review

Is there evidence that environmental noise is immunotoxic

Is type 1 diabetes a disease of the gut immune system triggered by cow's milk insulin

Isocyanate exposure and hypersensitivity pneumonitis--report of a probable case and prevalence of specific immunoglobulin G antibodies among exposed individuals

Isoflavonoid compounds from red clover (*Trifolium pratense*) protect from inflammation and immune suppression induced by UV radiation

Isoflurane anesthesia impairs the expression of immune neuromodulators in the hippocampus of aged mice

Isolated herpes simplex in the adult larynx as a rare complication of methotrexate-induced immunosuppression

Isolation and characterization of microglia from adult mouse brain: selected applications for ex vivo evaluation of immunotoxicological alterations following in vivo xenobiotic exposure

Issues raised by the reference doses for perfluorooctane sulfonate and perfluorooctanoic acid

iTRAQ: a method to elucidate cellular responses to mycotoxin zearalenone

Jeju ground water containing vanadium induced immune activation on splenocytes of low dose gamma-rays-irradiated mice

Jet fuel-induced immunotoxicity

JP-8 induces immune suppression via a reactive oxygen species NF-kappaB-dependent mechanism

JP-8 jet fuel exposure results in immediate immunotoxicity, which is cumulative over time

JP-8 jet fuel exposure suppresses the immune response to viral infections

Kaposi's sarcoma-associated herpesvirus-encoded replication and transcription activator impairs innate immunity via ubiquitin-mediated degradation of myeloid differentiation factor 88

Kawasaki syndrome. Association with exposure to carpet shampoo and successful therapy with immunoglobulins in the second week of the illness

Keratinocyte-derived cytokines and UVB-induced immunosuppression

Ketoprofen impairs immunosuppression induced by severe sepsis and reveals an important role for prostaglandin E2

Keyhole limpet haemocyanin - a model antigen for human immunotoxicological studies

Keynote lecture in the 13th Japanese Society of Immunotoxicology (JSIT 2006) : -Pathophysiological Development and Immunotoxicology: what we have found from research related to silica and silicate such as asbestos

Kidney disorders induced by non-steroidal anti-inflammatory agents and immunomodulators

Kinetic determination of vitellogenin induction in the epidermis of cyprinid and perciform fishes: Evaluation of sensitive enzyme-linked immunosorbent assays

Kinetic of the CMV-specific T-cell immune response and CMV infection in CMV-seropositive kidney transplant recipients receiving rabbit anti-thymocyte globulin induction therapy: A pilot study

Kinin B1 receptor deficiency attenuates cisplatin-induced acute kidney injury by modulating immune cell migration

Klotho protects human monocytes from LPS-induced immune impairment associated with immunosenescent-like phenotype

Laboratory diagnosis of heparin-associated thrombocytopenia and comparison of platelet aggregation test, heparin-induced platelet activation test, and platelet factor 4/heparin enzyme-linked immunosorbent assay

Laboratory diagnosis of heparin-induced thrombocytopenia in Asian Indians as investigated with functional and immunologic methods

Laboratory diagnosis of immune heparin-induced thrombocytopenia

Lack of autoantibody induction by mercury exposure in artisanal gold mining settings in Colombia: Findings and a review of the epidemiology literature

Lack of effect of Vitamin A on corticosteroid-induced immunosuppression

Lack of immunotoxicity of saquinavir (Ro 31-8959) used alone or in double or triple combination with AZT and ddC

Lactic acidemia in human immunodeficiency virus-uninfected infants exposed to perinatal antiretroviral therapy

Lactobacillus casei HY7213 ameliorates cyclophosphamide-induced immunosuppression in mice by activating NK, cytotoxic T cells and macrophages

Lactobacillus delbrueckii ssp. bulgaricus OLL1073R-1 feeding enhances humoral immune responses, which are suppressed by the antiviral neuraminidase inhibitor oseltamivir in influenza A virus-infected mice

Lactobacillus johnsonii provides a dose-dependent protection against UVR-induced immunosuppression

Lactobacillus paracasei BEJ01 prevents immunotoxic effects during chronic zearalenone exposure in Balb/c mice

Lactobacillus plantarum KLD51.0318 Ameliorates Impaired Intestinal Immunity and Metabolic Disorders in Cyclophosphamide-Treated Mice

Lactobacillus plantarum NCU116 Attenuates Cyclophosphamide-Induced Immunosuppression and Regulates Th17/Treg Cell Immune Responses in Mice

Lactobacillus reuteri CRL 1098 and Lactobacillus acidophilus CRL 1014 differently reduce in vitro immunotoxic effect induced by Ochratoxin A

Lamotrigine-induced common variable immune deficiency

Lamotrigine-induced toxic epidermal necrolysis treated with intravenous cyclosporin: a discussion of pathogenesis and immunosuppressive management

L-arginine and L-glutamine as immunonutrients and modulating agents for oxidative stress and toxicity induced by sodium nitrite in rats

L-Arginine reverses radiation-induced immune dysfunction: the need for optimum treatment window

Lasting effects of an impairment of Th1-like immune response in gamma-irradiated mice: A resemblance between irradiated mice and aged mice

Late conversion from tacrolimus to a belatacept-based immuno-suppression regime in kidney transplant recipients improves renal function, acid-base derangement and mineral-bone metabolism

Late onset neutropenia and immunoglobulin suppression of the patients with malignant lymphoma following autologous stem cell transplantation with rituximab

Late-onset pericardial tamponade, bilateral pleural effusions and recurrent immune monoarthritis induced by ipilimumab use for metastatic melanoma

Lead alters the immunogenicity of two neural proteins: a potential mechanism for the progression of lead-induced neurotoxicity

Lead and immune function

Lead exposure reduces carotenoid-based coloration and constitutive immunity in wild mallards

Lead induced modulation of splenic macrophage responses on humoral and cell mediated immunity

Lead toxicity on non-specific immune mechanisms of freshwater fish Channa punctatus

Lead toxicity: from overt to subclinical to subtle health effects

Leakage of mitochondrial and cytosolic GOT in isolated rat hepatocytes exposed to halothane: an immunohistochemical study

Lethal and sub-lethal evaluation of Indigo Carmine dye and byproducts after TiO<sub>2</sub> photocatalysis in the immune system of *Eisenia andrei* earthworms

Letter: Tinea faciei exacerbated by topical immunomodulation: Two case reports

Leukopenia and altered hematopoietic activity in mice exposed to the abused inhalant, isobutyl nitrite

Levels of serum oxidoreductases (LD, MD, ICD), blood proteins, immunoglobulins, and ceruloplasmin in a group of workers exposed to carbon disulfide

Levodopa administration in multiple sclerosis patients with immunosuppressive therapy-induced fatigue

Lichen planus and lichenoid drug-induced eruption: a histological and immunohistochemical study

Life events, frontal electroencephalogram laterality, and functional immune status after acute psychological stressors in adolescents

Life stress and cervical squamous intraepithelial lesions in women with human papillomavirus and human immunodeficiency virus

Lifelong persistent viral infection alters the naive T cell pool, impairing CD8 T cell immunity in late life

Lifespan-extending caloric restriction or mTOR inhibition impair adaptive immunity of old mice by distinct mechanisms

Life-threatening piperacillin-induced immune haemolysis in a patient with cystic fibrosis

Life-threatening rituximab-induced pyoderma gangrenosum successfully treated with intravenous immunoglobulin

Life-threatening ceftriaxone-induced immune hemolytic anemia in a child with Crohn's disease

Limited immunotoxic potential of technical formulation of the herbicide atrazine (AAtrex) in mice

Lindane-induced immunological alterations in human poisoning cases

Linear immunoglobulin A bullous dermatosis induced by gemcitabine

Linear immunoglobulin A bullous dermatosis possibly induced by mefenamic acid

Links between prolonged exposure to xenobiotics, increased incidence of hepatopathies, immunological disturbances and exacerbation of latent Epstein-Barr virus infections

Lipoic acid attenuates high fat diet-induced chronic oxidative stress and immunosuppression in mice jejunum: a microarray analysis

Lipopolysaccharide (LPS) induced activation of the immune system in control rats and rats chronically exposed to a low level of the organothiophosphate insecticide, acephate

Lipopolysaccharide impairs mucin secretion and stimulated mucosal immune stress response in respiratory tract of neonatal chicks

Lipopolysaccharide induces paired immunoglobulin-like receptor B (PirB) expression, synaptic alteration, and learning-memory deficit in rats

Lipopolysaccharide levels adherent to PM2.5 play an important role in particulate matter induced-immunosuppressive effects in mouse splenocytes

Lipopolysaccharide-induced immune activation impairs attention but has little effect on short-term working memory

Liposome-mediated combinatorial cytokine gene therapy induces localized synergistic immunosuppression and promotes long-term survival of cardiac allografts

Liver antioxidant and plasma immune responses in juvenile golden grey mullet (*Liza aurata*) exposed to dispersed crude oil

Liver enzyme function and humoral immunity in workers exposed to vinyl chloride

Local immunity and methods for its assessment upon exposure to ambient air-polluting biologicals

Local versus systemic immunotoxicity of isobutyl nitrite following subchronic inhalation exposure of female B6C3F1 mice

Localization and composition of renal immunodeposits in mice developing HgCl<sub>2</sub>-induced autoimmune process

Long term effects of carbaryl exposure on antiviral immune responses in *Xenopus laevis*

Long-term effect of *Spirulina platensis* extract on DMBA-induced hamster buccal pouch carcinogenesis (immunohistochemical study)

Long-term effects of immunotoxic cholinergic lesions in the septum on acquisition of the cone-field task and noncognitive measures in rats

Long-term effects on humoral immunity among workers exposed to 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD)

Long-term effects on the immune system following local irradiation for breast cancer. Pokeweed mitogen induced immunoglobulin secretion by blood lymphocytes and serum immunoglobulin levels

Long-term exposure to arsenic affects head kidney and impairs humoral immune responses of *Clarias batrachus*

Long-term exposure to high levels of decabrominated diphenyl ether inhibits CD4 T-cell functions in C57Bl/6 mice

Long-term exposure to PCB in broiler chickens--metabolic and immunotoxic effects

Long-term immune dysfunction after radiotherapy to the head and neck area

Long-term immune dysfunction in rainbow trout (*Oncorhynchus mykiss*) exposed as embryos to aflatoxin B1

Long-term Immunotoxic Effects of Oral Prenatal and Neonatal Atrazine Exposure

Long-term moderate calorie restriction inhibits inflammation without impairing cell-mediated immunity: a randomized controlled trial in non-obese humans

Long-term sun exposure alters the collagen of the papillary dermis. Comparison of sun-protected and photoaged skin by northern analysis, immunohistochemical staining, and confocal laser scanning microscopy

Long-term toxicity of immune suppression in juvenile rheumatic diseases

Long-Time Instead of Short-Time Exposure in Vitro and Administration in Vivo of Ochratoxin A Is Consistent in Immunosuppression

Long-wave UVA offers partial protection against UVB-induced immune suppression in human skin

Loperamide-induced expression of immune and inflammatory genes in Collies associated with ivermectin sensitivity

Low dose exposure to cadmium and its health effects. (3) Toxicity in laboratory animals and cultured cells

Low dose TBT exposure decreases amphipod immunocompetence and reproductive fitness

Low doses of monocrotaline in rats cause diminished bone marrow cellularity and compromised nitric oxide production by peritoneal macrophages

Low level exposure to chemicals and immune system

Low levels of residual oil fly ash (ROFA) impair innate immune response against environmental mycobacteria infection in vitro

Low or excess levels of dietary cholesterol impaired immunity and aggravated inflammation response in young grass carp (*Ctenopharyngodon idella*)

Low-dose azathioprine effectively suppresses clinical and immunological manifestations of generalized myasthenia

Low-dose exposure to inorganic mercury accelerates disease and mortality in acquired murine lupus

Low-dose inorganic mercury increases severity and frequency of chronic coxsackievirus-induced autoimmune myocarditis in mice

Low-dose interleukin-2 impairs host anti-tumor immunity and inhibits therapeutic responses in a mouse model of melanoma

Low-dose synergistic immunosuppression of T-dependent antibody responses by polycyclic aromatic hydrocarbons and arsenic in C57BL/6J murine spleen cells

Lower endotoxin immunity predicts increased cognitive dysfunction in elderly patients after cardiac surgery

Low-level dietary deoxynivalenol and acute exercise stress result in immunotoxicity in BALB/c mice

Low-level methylmercury exposure causes human T-cells to undergo apoptosis: evidence of mitochondrial dysfunction

LPS regulation of the immune response: Bacteroides endotoxin induces mitogenic, polyclonal, and antibody responses in classical LPS responsive but not C3H/HeJ mice

LPS-induced systemic inflammation reveals an immunomodulatory role for the prion protein at the blood-brain interface

Lycopene alleviates AFB1-induced immunosuppression by inhibiting oxidative stress and apoptosis in the spleen of mice

Lymphocyte dysfunction after DNA damage by toxic oxygen species. A model of immunodeficiency

Lymphocyte loss and immunosuppression following acetaminophen-induced hepatotoxicity in mice as a potential mechanism of tolerance

Lymphocyte proliferative response and tissue distribution of methylmercury sulfide and chloride in exposed rats

Lymphocyte subsets and Langerhans cells in sun-protected and sun-exposed skin of immunosuppressed renal allograft recipients

Macrophages are involved in hexachlorobenzene-induced adverse immune effects

Mafofamide induces DNA fragmentation and apoptosis in human T-lymphocytes. A possible mechanism of its immunosuppressive action

Maintenance rituximab following induction chemo-immunotherapy for mantle cell lymphoma: long-term follow-up of a pilot study from the Wisconsin Oncology Network

MAIT cells launch a rapid, robust and distinct hyperinflammatory response to bacterial superantigens and quickly acquire an anergic phenotype that impedes their cognate antimicrobial function: Defining a novel mechanism of superantigen-induced immunopathology and immunosuppression

Maize prolamins could induce a gluten-like cellular immune response in some celiac disease patients

Malathion immunotoxicity in the American lobster (*Homarus americanus*) upon experimental exposure

Malathion, lindane, and piperonyl butoxide, individually or in combined mixtures, induce immunotoxicity via apoptosis in murine splenocytes in vitro

Malnutrition and impaired immune response to infection

Malnutrition, urocanic acid, and sun may interact to suppress immunity in sojourners to high altitude

Mammalian immunoassays for predicting the toxicity of malathion in a laboratory fish model

Management of adverse events induced by next-generation immunomodulatory drug and proteasome inhibitors in multiple myeloma

Management of immune dysfunction after adult cardiac surgery

Managing immune checkpoint-inhibitor-induced severe autoimmune-like hepatitis by liver-directed topical steroids

Manganese tissue accumulation and tyrosine hydroxylase immunostaining response in the Neotropical freshwater crab, *Dilocarcinus pagei*, exposed to manganese

Manifestations of cellular immunity in the rat after prolonged asbestos inhalation. II. Alveolar macrophage-induced splenic lymphocyte proliferation

Markers for immunotoxic effects in rodents and man

Mast cells in the intestine and gills of the sea bream, *Sparus aurata*, exposed to a polychlorinated biphenyl, PCB 126

Mast cells mediate the immune suppression induced by dermal exposure to JP-8 jet fuel

Maternal adrenalectomy alters the immune and endocrine functions of fetal alcohol-exposed male offspring

Maternal and early postnatal polychlorinated biphenyl exposure in relation to total serum immunoglobulin concentrations in 6-month-old infants

Maternal exposure to airborne particulate matter causes postnatal immunological dysfunction in mice offspring

Maternal exposure to fish oil primes offspring to harbor intestinal pathobionts associated with altered immune cell balance

Maternal exposure to particulate matter alters early post-natal lung function and immune cell development

Maternal exposure to silver nanoparticles are associated with behavioral abnormalities in adulthood: Role of mitochondria and innate immunity in developmental toxicity

Maternal Immune Activation Causes Behavioral Impairments and Altered Cerebellar Cytokine and Synaptic Protein Expression

Maternal Lipopolysaccharide Exposure Promotes Immunological Functional Changes in Adult Offspring CD4<sup>+</sup> T Cells

Maternal lipopolysaccharide-induced inflammation during pregnancy programs impaired offspring innate immune responses

Maternal obesity programs offspring nonalcoholic fatty liver disease by innate immune dysfunction in mice

Maternal separation in early life impairs tumor immunity in adulthood in the F344 rat



Maternal smoking is associated with impaired neonatal toll-like-receptor-mediated immune responses

Maternal transfer and sublethal immune system effects of brevetoxin exposure in nesting loggerhead sea turtles (*Caretta caretta*) from western Florida

Maternal undernutrition induces neuroendocrine immune dysfunction in male pups at weaning

Maternal-Fetal Cancer Risk Assessment of Ochratoxin A during Pregnancy

MDCO-216 Does Not Induce Adverse Immunostimulation, in Contrast to Its Predecessor ETC-216

Measurement of airborne flour exposure with a monoclonal antibody-based immunoassay

Measures of resting immune function and related physiology in juvenile rainbow trout exposed to a pulp mill effluent

Mechanical ventilation of healthy rats suppresses peripheral immune function

Mechanism of 7,12-dimethylbenz[a]anthracene-induced immunotoxicity: role of metabolic activation at the target organ

Mechanism of deoxynivalenol effects on the reproductive system and fetus malformation: Current status and future challenges

Mechanism of immunosuppression in zebrafish (*Danio rerio*) spleen induced by environmentally relevant concentrations of perfluorooctanoic acid

Mechanism of immunotoxicological effects of tributyltin chloride on murine thymocytes

Mechanism of perfluorooctanesulfonate (PFOS)-induced apoptosis in the immunocyte

Mechanism of systemic immune suppression by UV irradiation in vivo. II. The UV effects on number and morphology of epidermal Langerhans cells and the UV-induced suppression of contact hypersensitivity have different wavelength dependencies

Mechanism of TCDD-induced suppression of antibody production: effect on T cell-derived cytokine production in the primary immune reaction of mice

Mechanism of UVB-induced suppression of the immune response to *Mycobacterium bovis* bacillus Calmette-Guerin: role of cytokines on macrophage function

Mechanisms for how inhaled multiwalled carbon nanotubes suppress systemic immune function in mice

Mechanisms involved in the immunotoxicity induced by dermal application of JP-8 jet fuel

Mechanisms of 1,2,3,4,6,7,8-heptachlorodibenzo-p-dioxin (HpCDD)-induced humoral immune suppression: evidence of primary defect in T-cell regulation

Mechanisms of dimethylbenzanthracene-induced immunotoxicity

Mechanisms of ethanol-induced immunosuppression

Mechanisms of immune dysfunction in stem cell transplantation

Mechanisms of immunotoxic effects of acrylonitrile

Mechanisms of organophosphorus pesticide toxicity in the context of airway hyperreactivity and asthma

Mechanisms underlying the suppression of established immune responses by ultraviolet radiation

Mechanisms underlying UV-induced immune suppression: implications for sunscreen design

Medical management and sublingual immunotherapy practices in patients with house dust mite-induced respiratory allergy: a retrospective, observational study

Melatonin alleviates aluminium chloride-induced immunotoxicity by inhibiting oxidative stress and apoptosis associated with the activation of Nrf2 signaling pathway

Melatonin ameliorates ochratoxin A-induced oxidative stress and apoptosis in porcine oocytes

Mercuric chloride induces a strong immune activation, but does not accelerate the development of dermal fibrosis in tight skin 1 mice

Mercury Accumulation, Structural Damages, and Antioxidant and Immune Status Changes in the Gilthead Seabream (*Sparus aurata* L.) Exposed to Methylmercury

Mercury and autoimmunity: implications for occupational and environmental health

Mercury compounds and the immune system: a review

Mercury exposure and children's health

Mercury Exposure and Heart Diseases

Mercury exposure in protein A immunoadsorption

Mercury exposure, serum antinuclear/antinucleolar antibodies, and serum cytokine levels in mining populations in Amazonian Brazil: a cross-sectional study

Mercury impairment of mouse thymocyte survival in vitro: involvement of cellular thiols

Mercury-induced autoimmune glomerulonephritis in inbred rats. II. Immunohistopathology, histopathology and effects of prostaglandin administration

Meropenem-induced immune thrombocytopenia and the diagnostic process of laboratory testing

Mesenteric lymph node T $\gamma$ ammadelta cells induced by gastrectomy in mice suppress cell-mediated immune response in vitro via released TGF- $\beta$

Metabolic and immune impairments induced by the endocrine disruptors benzo[a]pyrene and triclosan in *Xenopus tropicalis*

Metabolic and immunological disorders induced by long-term treatment with phenytoin

Metabolic pathways of ochratoxin A

Metabolic profiling study on potential toxicity and immunotoxicity-biomarker discovery in rats treated with cyclophosphamide using HPLC-ESI-IT-TOF-MS

Metabolic requirements for induction of contact hypersensitivity to immunotoxic polyaromatic hydrocarbons

Metabolic Syndrome after Hematopoietic Cell Transplantation: At the Intersection of Treatment Toxicity and Immune Dysfunction

Metabolism and toxicity of aflatoxins M1 and B1 in human-derived in vitro systems

Metabolite-specific (IgG) and drug-specific antibodies (IgG, IgM) in two cases of trimethoprim-sulfamethoxazole-induced immune thrombocytopenia

Metal allergens of growing significance: epidemiology, immunotoxicology, strategies for testing and prevention

Metal-induced impairment of the cellular immunity of newborn harbor seals (*Phoca vitulina*)

Metallothionein-null mice are highly susceptible to the hematotoxic and immunotoxic effects of chronic CdCl<sub>2</sub> exposure

Methamphetamine activates nuclear factor kappa-light-chain-enhancer of activated B cells (NF-kappaB) and induces human immunodeficiency virus (HIV) transcription in human microglial cells

Methamphetamine induces trace amine-associated receptor 1 (TAAR1) expression in human T lymphocytes: role in immunomodulation

Method for evaluation of immunotoxicity of dioxin compounds using human T-lymphoblastic cell line, L-MAT

Methods for Analysis of Nanoparticle Immunosuppressive Properties In Vitro and In Vivo

Methods for Assessing Mast Cell Responses to Engineered Nanomaterial Exposure

Methotrexate-induced iatrogenic immunodeficiency-associated lymphoproliferative disorder causing hypercalcaemia

Methylenedioxymethamphetamine ('Ecstasy')-induced immunosuppression: a cause for concern

Methylmercury/copper effects on hemosiderin: possible mechanism of immune suppression in fish

Methylmercury-induced movement and postural disorders in developing rat: loss of somatostatin-immunoreactive interneurons in the striatum

Methylprednisolone-induced immune thrombocytopenia

Methylprednisolone-Induced Lymphocytosis in Patients with Immune-Mediated Inflammatory Disorders

Mice repeatedly exposed to Group-A beta-Haemolytic Streptococcus show perseverative behaviors, impaired sensorimotor gating, and immune activation in rostral diencephalon

Mice with genetically determined high susceptibility to ultraviolet (UV)-induced immunosuppression show enhanced UV carcinogenesis

Microbial Degradation of Cellular Kinases Impairs Innate Immune Signaling and Paracrine TNFalpha Responses

Microcystin-LR exposure to adult zebrafish (*Danio rerio*) leads to growth inhibition and immune dysfunction in F1 offspring, a parental transmission effect of toxicity

Microcystin-LR Induced Immunotoxicity in Mammals

Microcystin-LR induces cytotoxicity and affects carp immune cells by impairment of their phagocytosis and the organization of the cytoskeleton

Microcystin-LR modulates selected immune parameters and induces necrosis/apoptosis of carp leucocytes

Microcystin-LR toxicodynamics, induced pathology, and immunohistochemical localization in livers of blue-green algae exposed rainbow trout (*Oncorhynchus mykiss*)

Microgravity inhibits resting T cell immunity in an exposure time-dependent manner

Micronized, Heat-Treated *Lactobacillus plantarum* LM1004 Alleviates Cyclophosphamide-Induced Immune Suppression

Micronutrient deficiencies are associated with impaired immune response and higher burden of respiratory infections in elderly Ecuadorians

MicroRNA-155 and microRNA-181a, via HO-1, participate in regulating the immunotoxicity of cadmium in the kidneys of exposed *Cyprinus carpio*

Microsomal epoxide hydrolase is required for 7,12-dimethylbenz[a]anthracene (DMBA)-induced immunotoxicity in mice

Midazolam impairs immune functions: it's time to take care of dendritic cells

Miliary *Mycobacterium bovis* induced by intravesical bacille Calmette-Guerin immunotherapy

Minimal immunological changes in structurally malformed rats after prenatal exposure to cyclophosphamide

Minimal immunological effects on workers with prolonged low exposure to inorganic mercury

Minimum doses of ultraviolet radiation required to induce murine skin edema and immunosuppression are different and depend on the ultraviolet emission spectrum of the source

Minocycline-induced immune thrombocytopenia presenting as Schamberg's disease

Minor effects of low exposure to inorganic mercury on the human immune system

MiR-122 partly mediates the ochratoxin A-induced GC-2 cell apoptosis

MiR-582-5p/miR-590-5p targeted CREB1/CREB5-NF-kappaB signaling and caused opioid-induced immunosuppression in human monocytes

Misoprostol impairs female reproductive tract innate immunity against *Clostridium sordellii*

Mitigation of Hydrophobicity-Induced Immunotoxicity by Sugar Poly(orthoesters)

Mitochondrial functions of THP-1 monocytes following the exposure to selected natural compounds

Mitochondrial peroxiredoxin 3 (Prx3) from rock bream (*Oplegnathus fasciatus*): immune responses and role of recombinant Prx3 in protecting cells from hydrogen peroxide induced oxidative stress

Mitotic activity of keratinocytes in nifedipine- and immunosuppressive medication-induced gingival overgrowth

Mixed mold mycotoxicosis: immunological changes in humans following exposure in water-damaged buildings

MNU-induced rat mammary carcinomas: immunohistology and estrogen receptor expression

Modeling and predicting immunological effects of chemical stressors: characterization of a quantitative biomarker for immunological changes caused by atrazine and ethanol

Modeling the Function of TATA Box Binding Protein in Transcriptional Changes Induced by HIV-1 Tat in Innate Immune Cells and the Effect of Methamphetamine Exposure

Modeling toxicodynamic effects of trichloroethylene on liver in mouse model of autoimmune hepatitis

Modelling the immunopathophysiology of *Brucella melitensis* and its lipopolysaccharide in mice infected via oral route of exposure

Moderate prenatal alcohol exposure suppresses the TLR4-mediated innate immune response in the hippocampus of young rats

Modification of immunoreactive EGF and EGF receptor after acute tubular necrosis induced by tobramycin or cisplatin

Modification of primary amines to higher order amines reduces in vivo hematological and immunotoxicity of cationic nanocarriers through TLR4 and complement pathways

Modifications of immunological and neuro-endocrine parameters induced by antiorthostatic bed-rest in human healthy volunteers

Modifications of neuronal phosphorylated tau immunoreactivity induced by NMDA toxicity

Modulating effect of MgO-SiO<sub>2</sub> nanoparticles on immunological and histopathological alterations induced by aflatoxicosis in rats

Modulation of benzo[a]pyrene induced immunotoxicity in mice actively immunized with a B[a]P-diphtheria toxoid conjugate

Modulation of benzo[a]pyrene induced neurotoxicity in female mice actively immunized with a B[a]P-diphtheria toxoid conjugate

Modulation of biochemical parameters by *Hemidesmus indicus* in cumene hydroperoxide-induced murine skin: possible role in protection against free radicals-induced cutaneous oxidative stress and tumor promotion

Modulation of immune and antioxidant responses by azinphos-methyl in the freshwater mussel *Diplodon chilensis* challenged with *Escherichia coli*

Modulation of immune response by organophosphorus pesticides: fishes as a potential model in immunotoxicology

Modulation of immune responses with nanoparticles and reduction of their immunotoxicity

Modulation of immune-associated parameters and antioxidant responses in the crab (*Scylla serrata*) exposed to mercury

Modulation of macrophage activation by ammonium metavanadate

Modulation of murine peritoneal macrophage function by chronic exposure to arsenate in drinking water

Modulation of N-methyl-N-nitrosourea-induced crypt restricted metallothionein immunopositivity in mouse colon by a non-genotoxic diet-related chemical

Modulation of signal transduction pathways in lymphocytes due to sub-lethal toxicity of chlorinated phenol

Modulation of T-helper cell populations: potential mechanisms of respiratory hypersensitivity and immune suppression

Modulatory efficacy of green tea polyphenols on glycoconjugates and immunological markers in 4-Nitroquinoline 1-oxide-induced oral carcinogenesis-A therapeutic approach

Modulatory role of dietary *Chlorella vulgaris* powder against arsenic-induced immunotoxicity and oxidative stress in Nile tilapia (*Oreochromis niloticus*)

Modulatory role of dietary *Thymus vulgaris* essential oil and *Bacillus subtilis* against thiamethoxam-induced hepatorenal damage, oxidative stress, and immunotoxicity in African catfish (*Clarias garipenus*)

Modulatory role of L-carnitine against microcystin-LR-induced immunotoxicity and oxidative stress in common carp

Molecular and immune toxicity of CoCr nanoparticles in MoM hip arthroplasty

Molecular and immunological toxic effects of nanoparticles

Molecular cloning, immunohistochemical localization, characterization and expression analysis of caspase-8 from the blunt snout bream (*Megalobrama amblycephala*) exposed to ammonia

Molecular determinants of UV-induced immunosuppression

Molecular insight of arsenic-induced carcinogenesis and its prevention

Molecular mechanisms of amitraz mammalian toxicity: a comprehensive review of existing data

Molecular mechanisms underlying mancozeb-induced inhibition of TNF- $\alpha$  production

Monitoring antioxidant defenses and free radical production in space-flight, aviation and railway engine operators, for the prevention and treatment of oxidative stress, immunological impairment, and premature cell aging

Monitoring of azathioprine-induced immunosuppression with thiopurine methyltransferase activity in kidney transplant recipients

Monoclonal gammopathy after intense induction immunosuppression in renal transplant patients

Morphine impairs host innate immune response and increases susceptibility to *Streptococcus pneumoniae* lung infection

Morphine induced, behavioural, biochemical and immunological correlations

Morphine or U-50,488 suppresses Fos protein-like immunoreactivity in the spinal cord and nucleus tractus solitarius evoked by a noxious visceral stimulus in the rat

Morphological alteration, lysosomal membrane fragility and apoptosis of the cells of Indian freshwater sponge exposed to washing soda (sodium carbonate)

Morphological analysis of glomerular lesions in renal transplants immunosuppressed with cyclosporine A (CYA): has CYA induced a new transplant glomerular lesion

Morphological and immunobiochemical analysis of the liver in L-arginine induced experimental chronic pancreatitis

Morphological and immunohistochemical changes on rat skin exposed to nitrogen mustard

Morphological and immunohistochemical characteristics of the thymus during chemical carcinogenesis induced by 1,2-dimethylhydrazine administration

Morphological and immunohistochemical studies on porcine serum-induced rat liver fibrosis

Morphological changes in the thyroid and adrenals under the bitemporal action of a UHF electrical field and decimeter waves (experimental research).

Morphological features of the organs of the immune system in conditions of exposure to chromium and benzene

Mouse allergen exposure and immunologic responses: IgE-mediated mouse sensitization and mouse specific IgG and IgG4 levels

Mouse lung immune response after acute exposure to flour dust

Moxifloxacin as the likely cause of drug-induced linear immunoglobulin A bullous dermatosis

Mucosal-associated invariant T cells from patients with tuberculosis exhibit impaired immune response

Multilaboratory evaluation of 15 bioassays for (eco)toxicity screening and hazard ranking of engineered nanomaterials: FP7 project NANOVALID

Multiorgan failure induced by intravenous immunoglobulin

Multiparametric immunotoxicity screening in mice during early drug development

Multiple biomarkers of the cytotoxicity induced by BDE-47 in human embryonic kidney cells

Multiple enlarged metabolically active lymph nodes in 18F-FDG PET/CT after anti-CTLA-4 antibody therapy in metastatic melanoma - disease progression or immunologically induced side effect

Multiple experimental approaches of immunotoxic effects of mercury chloride in the blue mussel, *Mytilus edulis*, through in vivo, in tubo and in vitro exposures

Multiple myeloma: clusters, clues, and dioxins

Murine membranous nephropathy: immunization with alpha3(IV) collagen fragment induces subepithelial immune complexes and Fc gamma R-independent nephrotic syndrome

Murine model of ethanol-induced immunosuppression

Mutagenic and morphologic impacts of 1.8GHz radiofrequency radiation on human peripheral blood lymphocytes (hPBLs) and possible protective role of pre-treatment with Ginkgo biloba (EGb 761).

Mutual augmentation of the induction of the histamine-forming enzyme, histidine decarboxylase, between alendronate and immuno-stimulants (IL-1, TNF, and LPS), and its prevention by clodronate

Myasthenia triggered by immune checkpoint inhibitors: New case and literature review

Mycobacterium infection from a cardiopulmonary bypass heater-cooler unit in a patient with steroid-induced immunosuppression

Mycobiota and Ochratoxin A in laboratory mice feed: preliminary study

Mycolactone-producing *Mycobacterium marinum* infection in captive Hong Kong warty newts and pathological evidence of impaired host immune function

Mycophenolate mofetil and reduced doses of cyclosporine in pediatric liver transplantation with chronic renal dysfunction: changes in the immune responses

Mycophenolate mofetil-based calcineurin inhibitor reduced immunosuppressive protocol for the improvement of renal dysfunction after liver transplantation

Mycophenolate mofetil-based, cyclosporine-free induction and maintenance immunosuppression: first-3-months analysis of efficacy and safety in two cohorts of renal allograft recipients

Mycotoxins in children

Mycotoxins, general view, chemistry and structure

Myeloid-derived suppressor cells accumulate in the liver site after sepsis to induce immunosuppression

N,N'-diethyl-m-toluamide (DEET) suppresses humoral immunological function in B6C3F1 mice

Nanomaterial Exposure Induced Neutrophil Extracellular Traps: A New Target in Inflammation and Innate Immunity

Nanomaterial Induced Immune Responses and Cytotoxicity



Nanoparticles of zinc oxide defeat chlorpyrifos-induced immunotoxic effects and histopathological alterations

Nanosafety studies of nanomaterials about biodistribution and immunotoxicity

Natural killer activity in Fischer-344 rat lungs as a method to assess pulmonary immunocompetence: immunosuppression by phosgene inhalation

Naturally occurring level of mixed aflatoxins B and G stimulate toll-like receptor-4 in bovine mononuclear cells

Necrotizing Autoimmune myopathy: A case report on statin induced rhabdomyolysis requiring immunosuppressive therapy

Negative effect of chronic cadmium exposure on growth, histology, ultrastructure, antioxidant and innate immune responses in the liver of zebrafish: Preventive role of blue light emitting diodes

Neonatal exposure to mild hyperoxia causes persistent increases in oxidative stress and immune cells in the lungs of mice without altering lung structure

Neonatal exposure to UVR alters skin immune system development, and suppresses immunity in adulthood

Neonatal phytoestrogen exposure alters oviduct mucosal immune response to pregnancy and affects preimplantation embryo development in the mouse

Neonicotinoid-induced pathogen susceptibility is mitigated by *Lactobacillus plantarum* immune stimulation in a *Drosophila melanogaster* model

Nerve growth factor produced by activated human monocytes/macrophages is severely affected by ethanol

Neurodegeneration Triggers Peripheral Immune Cell Recruitment into the Forebrain

Neurodevelopmental disorders: Impaired immune system function linked to social behaviour deficits in mice

Neuroendocrine and Eosinophilic Granule Cells in the Gills of Tilapia, *Oreochromis niloticus*: Effects of Waterborne Copper Exposure

Neuroendocrine and immunotoxicity of polyaromatic hydrocarbon, chrysene in crustacean post larvae

Neuro-immune interactions in psychopathology with the example of interferon-alpha-induced depression

Neurologic and immunologic effects of exposure to corticosterone, chlorpyrifos, and multiple doses of tri-ortho-tolyl phosphate over a 28-day period in rats

Neuro-Modulation of Immuno-Endocrine Response Induced by Kalitoxin of *Androctonus* Scorpion Venom

Neuron specific enolase (NSE) immunostaining detection of endocrine cell hyperplasia in adult rats exposed to asbestos

Neuron-immune interactions in the sensitized thalamus induced by mustard oil application to rat molar pulp

Neuropathophysiological effect and immuno-inflammatory response induced by kaliotoxin of androctonus scorpion venom

Neurophysiological, histological and immunohistochemical characterization of bortezomib-induced neuropathy in mice

Neurotoxic Anatoxin-a Can Also Exert Immunotoxicity by the Induction of Apoptosis on *Carassius auratus* Lymphocytes in vitro When Exposed to Environmentally Relevant Concentrations

Neurotoxic and immunotoxic effects of fenthion and omethoate on frogs at acute exposure

Neurotoxic and immunotoxic effects of Indole-3-butyric acid on rats at subacute and subchronic exposure

Neurotoxic effects of ochratoxin A on the subventricular zone of adult mouse brain

Neurotoxicity and immunotoxicity assessment in CBA/J mice with chronic *Toxoplasma gondii* infection and multiple oral exposures to methylmercury

Neurotoxicity and immunotoxicity assessment in CBA/J mice with chronic *Toxoplasma gondii* infection and single-dose exposure to methylmercury

Neurotoxicity of toluene

Neutralizing antibodies obtained in a persistent immune response are effective against deleterious effects induced by the *Thalassophryne nattereri* fish venom

Neutrophil depletion during *Toxoplasma gondii* infection leads to impaired immunity and lethal systemic pathology

Neutrophil function after exposure to polychlorinated biphenyls in vitro

Neutrophil protease inhibition reduces neuromyelitis optica-immunoglobulin G-induced damage in mouse brain

New insights into the mechanisms of polymorphic light eruption: resistance to ultraviolet radiation-induced immune suppression as an aetiological factor

New mechanism of organophosphorus pesticide-induced immunotoxicity

NF-kappaB activation via MyD88-dependent Toll-like receptor signaling is inhibited by trichothecene mycotoxin deoxynivalenol

Nickel - role in human organism and toxic effects

Nicotinamide reduces photodynamic therapy-induced immunosuppression in humans

Nitric oxide appears to be a mediator of solar-simulated ultraviolet radiation-induced immunosuppression in humans

Nitric oxide synthase immunolocalization and expression in the rat hippocampus after sub-acute lead acetate exposure in rats

Nitric oxide synthase inhibitor, aminoguanidine reduces intracerebroventricular colchicine induced neurodegeneration, memory impairments and changes of systemic immune responses in rats

Nitrofurantoin-induced immune-mediated lung and liver disease

Nivolumab-induced acute-onset type 1 diabetes mellitus as an immune-related adverse event: A case report

Nivolumab-Induced Subcutaneous Fat Necrosis: Another FDG-Avid Immune-Related Adverse Event

Nivolumab-induced type 1 diabetes mellitus as an immune-related adverse event

Non-analgesic effects of opioids: mechanisms and potential clinical relevance of opioid-induced immunodepression

Noncardiogenic pulmonary edema triggered by intravenous immunoglobulin in cancer-associated thrombotic thrombocytopenic purpura-hemolytic uremic syndrome

Nonclinical regulatory immunotoxicity testing of nanomedicinal products: Proposed strategy and possible pitfalls

Non-coplanar PCB-mediated modulation of human leukocyte phagocytosis: a new mechanism for immunotoxicity

Non-coplanar polychlorinated biphenyl (PCB)-induced immunotoxicity is coincident with alterations in the serotonergic system

Non-cytotoxic hydroxyl-functionalized exfoliated boron nitride nanoflakes impair the immunological function of insect haemocytes in vivo

Non-immunologic factor: immunosuppressive drug-induced nephrotoxicity

Normal development of tyrosine hydroxylase and serotonin immunoreactive fibers innervating anterior cingulate cortex and visual cortex in rabbits exposed prenatally to cocaine

Normalizing effect of plant-originated glycoprotein (116 kDa) on G0/G1 arrest in cadmium chloride-induced primary cultured mouse myelocytes

Novel Acaricidal Drug Fluazuron Causes Immunotoxicity via Selective Depletion of Lymphocytes T CD8

Novel Approach in Monocyte Intracellular TNF Measurement: Application to Sepsis-Induced Immune Alterations

Novel biomarkers of mercury-induced autoimmune dysfunction: a cross-sectional study in Amazonian Brazil

Novel CD3-specific antibody induces immunosuppression via impaired phosphorylation of LAT and PLCgamma1 following T-cell stimulation

Novel compound 2-methyl-2H-pyrazole-3-carboxylic acid (2-methyl-4-o-tolylazo-phenyl)-amide (CH-223191) prevents 2,3,7,8-TCDD-induced toxicity by antagonizing the aryl hydrocarbon receptor

Novel evidence of microglial immune response in impairment of Dengue infection of CNS

Novel gene markers of immunosuppressive chemicals in mouse lymph node assay

Novel mechanisms of gliadin immunotoxicity

Novel Use of All-Trans-Retinoic Acid in A Model of Lipopolysaccharide-Immunosuppression to Decrease the Generation of Myeloid-Derived Suppressor Cells by Reducing the Proliferation of CD34+ Precursor Cells

Nrf2 Regulates the Risk of a Diesel Exhaust Inhalation-Induced Immune Response during Bleomycin Lung Injury and Fibrosis in Mice

Nrf2-dependent repression of interleukin-12 expression in human dendritic cells exposed to inorganic arsenic

Nutritional aspects of immunosuppression in athletes

O(6)-benzylguanine enhances the in vitro immunotoxic activity of temozolomide on natural or antigen-dependent immunity

Obesity impairs cell-mediated immunity during the second trimester of pregnancy

Obesity is associated with impaired immune response to influenza vaccination in humans

Obesity: influenza immunity impaired in obesity

Obesity-induced tissue free radical generation: an in vivo immuno-spin trapping study

Occupational exposure to formaldehyde and early biomarkers of cancer risk, immunotoxicity and susceptibility

Occurrences and Outcomes of Immune Checkpoint Inhibitors-Induced Vitiligo in Cancer Patients: A Retrospective Cohort Study

Ochratoxin A exerts neurotoxicity in human astrocytes through mitochondria-dependent apoptosis and intracellular calcium overload

Ochratoxin A exposure biomarkers in the Czech Republic and comparison with foreign countries

Ochratoxin A from a toxicological perspective

Ochratoxin A induces nephrotoxicity and immunotoxicity through different MAPK signaling pathways in PK15 cells and porcine primary splenocytes

Ochratoxin A induces oxidative DNA damage and G1 phase arrest in human peripheral blood mononuclear cells in vitro

Ochratoxin A inhibits mouse embryonic development by activating a mitochondrion-dependent apoptotic signaling pathway

Ochratoxin A inhibits the production of tissue factor and plasminogen activator inhibitor-2 by human blood mononuclear cells: another potential mechanism of immune-suppression

Ochratoxin A: developmental and reproductive toxicity-an overview

Ochratoxin A: its cancer risk and potential for exposure

Ochratoxin A: previous risk assessments and issues arising

Ochratoxin A: the continuing enigma

Ochratoxin A: Toxicity, oxidative stress and metabolism

Ofloxacin eyedrops-induced toxic epidermal necrolysis treated with intravenous immunoglobulin

Oil mist exposure in industrial health--a review

OKT3 induction and steroid-free maintenance immunosuppression for treatment of high-risk heart transplant recipients

Omalizumab pretreatment decreases acute reactions after rush immunotherapy for ragweed-induced seasonal allergic rhinitis

Omics analysis of low dose acetaminophen intake demonstrates novel response pathways in humans

On the long-term effects of methyl isocyanate on cell-mediated immunity in Bhopal gas-exposed long-term survivors and their offspring

O-phenylphenol and its sodium and potassium salts: a toxicological assessment

Opioid-Induced Hyperalgesia Is Associated with Dysregulation of Circadian Rhythm and Adaptive Immune Pathways in the Mouse Trigeminal Ganglia and Nucleus Accumbens

Opioid-induced immunosuppression

Opioid-induced immunosuppression. A clinically relevant problem

Opioid-induced immunosuppression: is it centrally mediated or peripherally mediated

Opposite effects of the maternal immune system activated by interleukin-1 $\beta$  vs. PSK and OK432 on 5-azacytidine-induced birth defects

Optimal immunosuppressor induces stable gut microbiota after liver transplantation

Options for induction immunosuppression in liver transplant recipients

Oral (gavage), in utero and postnatal exposure of Sprague-Dawley rats to low doses of tributyltin chloride. Part 1: Toxicology, histopathology and clinical chemistry

Oral (gavage), in utero and post-natal exposure of Sprague-Dawley rats to low doses of tributyltin chloride. Part II: effects on the immune system

Oral administration of lipoteichoic acid from *Lactobacillus rhamnosus* GG overcomes UVB-induced immunosuppression and impairs skin tumor growth in mice

Oral administration of the immunomodulator JBT-3002 induces endogenous interleukin 15 in intestinal macrophages for protection against irinotecan-mediated destruction of intestinal epithelium

Oral benzo[a]pyrene: understanding pharmacokinetics, detoxication, and consequences--Cyp1 knockout mouse lines as a paradigm

Oral Exposure to Atrazine Induces Oxidative Stress and Calcium Homeostasis Disruption in Spleen of Mice

Oral exposure to benzo[a]pyrene in the mouse: detoxication by inducible cytochrome P450 is more important than metabolic activation

Oral exposure to immunostimulating drugs results in early changes in innate immune parameters in the spleen

Oral exposure to inorganic mercury alters T lymphocyte phenotypes and cytokine expression in BALB/c mice

Oral exposure to mineral oils: Is there an association with immune perturbation and autoimmunity

Oral exposure to the herbicide simazine induces mouse spleen immunotoxicity and immune cell apoptosis

Oral mucosal changes induced by anticancer targeted therapies and immune checkpoint inhibitors

Organization of research team for nano-associated safety assessment in effort to study nanotoxicology of zinc oxide and silica nanoparticles

Organochlorine compounds: Risk of non-Hodgkin's lymphoma and breast cancer

Organochlorine contaminant exposure and associations with hematological and humoral immune functional assays with dam age as a factor in free-ranging northern fur seal pups (*Callorhinus ursinus*)

Organochlorine pesticide-induced oxidative stress and immune suppression in rats

Organochlorine pollutants [corrected] in California sea lions revisited

Organophosphorus pesticide immunotoxicity: effects of O,O,S-trimethyl phosphorothioate on cellular and humoral immune response systems

Organotins disrupt the 11 $\beta$ -hydroxysteroid dehydrogenase type 2-dependent local inactivation of glucocorticoids

Oxaliplatin Immune-Induced Syndrome Occurs With Cumulative Administration and Rechallenge: Single Institution Series and Systematic Review Study

Oxaliplatin induces a delayed immune-mediated hemolytic anemia: a case report and review of the literature

Oxaliplatin-induced hypersensitivity reaction displaying marked elevation of immunoglobulin E

Oxaliplatin-induced immune pancytopenia

Oxaliplatin-induced immune thrombocytopenia

Oxaliplatin-induced immune thrombocytopenia in a patient with colon cancer

Oxazolone-Induced Colitis as a Model of Th2 Immune Responses in the Intestinal Mucosa

Oxcarbazepine-induced immunoglobulin deficiency

Oxidative damage and immunotoxicity effect of people who exposed to electronic waste

Oxidative stress and apoptotic changes in murine splenocytes exposed to cadmium

Oxidative stress and immune disturbance after long-term exposure to bisphenol A in juvenile common carp (*Cyprinus carpio*)

Oxidative stress and immunotoxic effects of bisphenol A on the larvae of rare minnow *Gobiocypris rarus*

Oxidative stress and immunotoxic effects of lead and their amelioration with myrrh (*Commiphora molmol*) emulsion

Oxidative stress and immunotoxicity induced by graphene oxide in zebrafish

Oxidative stress, immunological response, and heat shock proteins induction in the Chinese Mitten Crab, *Eriocheir sinensis* following avermectin exposure

Oxidative stress, ion concentration change and immune response in gills of common carp (*Cyprinus carpio*) under long-term exposure to bisphenol A

Oxidative stress, neurotoxicity, and non-specific immune responses in juvenile red sea bream, *Pagrus major*, exposed to different waterborne selenium concentrations

Oxidative stress-induced skeletal muscle injury involves in NF-kappaB/p53-activated immunosuppression and apoptosis response in copper (II) or/and arsenite-exposed chicken

Oxidized lipid depresses canine growth, immune function, and bone formation

Oxidized phospholipid, 1-palmitoyl-2-(9'-oxo-nonanoyl)-glycerophosphocholine (PON-GPC), produced in the lung due to cigarette smoking, impairs immune function in macrophages

Ozone exposure impairs antigen-specific immunity but activates IL-7-induced proliferation of CD4-CD8-thymocytes in BALB/c mice

p38 mitogen-activated protein kinase impairment of innate immune cells is a characteristic feature of HBeAg-negative chronic hepatitis B

p53 and ATM/ATR regulate 7,12-dimethylbenz[a]anthracene-induced immunosuppression

Pact with the devil: alemtuzumab therapy, immune suppression and infectious complications in chronic lymphocytic leukemia

Palivizumab is highly effective in suppressing respiratory syncytial virus in an immunosuppressed animal model

Panitumumab-Induced Immune Complex Glomerulonephritis

Paper and pulp mill effluent-induced immunotoxicity in freshwater fish *Channa punctatus* (Bloch

Parameters of immunological competence in subjects with high consumption of fish contaminated with persistent organochlorine compounds

Paraquat Preferentially Induces Apoptosis of Late Stage Effector Lymphocyte and Impairs Memory Immune Response in Mice

Parasitism and venom of ectoparasitoid *Scleroderma guani* impairs host cellular immunity

Parathion-induced suppression of humoral immunity in inbred mice

Pargyline pretreatment prevents immunological changes induced by MPTP in mice

Partial regression of Merkel cell carcinoma in response to withdrawal of azathioprine in an immunosuppression-induced case of metastatic Merkel cell carcinoma

Past, present and emerging toxicity issues for jet fuel

Pathogenesis of trehalose dimycolate-induced interstitial pneumonitis. IV. Evidence against roles for immunoglobulins and the complement system

Pathogenetic analyses of carbamazepine-induced liver injury in F344 rats focused on immune- and inflammation-related factors

Pathological damage and immunomodulatory effects of zebrafish exposed to microcystin-LR

Pathological findings in rabbits and sheep following the subacute administration of triphenyltin acetate

Pathological, immunological and biochemical markers of subchronic arsenic toxicity in rats

Pathophysiologic and immunologic mechanisms of contrast medium-induced anaphylactoid immediate hypersensitivity--an overview

Patients with quinine-induced immune thrombocytopenia have both "drug-dependent" and "drug-specific" antibodies

Patterns of immunotoxicity associated with chronic as compared with acute exposure to chemical or physical stressors and their relevance with regard to the role of stress and with regard to immunotoxicity testing

Patulin immunotoxicology: effect on phagocyte activation and the cellular and humoral immune system of mice and rabbits

Pb exposure attenuates hypersensitivity in vivo by increasing regulatory T cells

PCB 28 metabolites elimination kinetics in human plasma on a real case scenario: Study of hydroxylated polychlorinated biphenyl (OH-PCB) metabolites of PCB 28 in a highly exposed German Cohort

PCB-induced hepatic Cyp1A induction is associated with innate immune dysfunction in a feral teleost fish



PCB-related alteration of thyroid hormones and thyroid hormone receptor gene expression in free-ranging harbor seals (*Phoca vitulina*)

Peculiarities of immunologic reactivity in workers exposed to various neurotoxic chemicals

Pediatric renal transplantation with mycophenolate mofetil-based immunosuppression without induction: results after three years

Pefloxacin and ciprofloxacin increase UVA-induced edema and immune suppression

Pegylated interferon-induced immune-mediated hepatitis post-liver transplantation

PEGylation is effective in reducing immunogenicity, immunotoxicity, and hepatotoxicity of alpha-momorcharin in vivo

Pembrolizumab-Induced Encephalopathy: A Review of Neurological Toxicities with Immune Checkpoint Inhibitors

Pembrolizumab-Induced Extensive Panniculitis and Nevus Regression: Two Novel Cutaneous Manifestations of the Post-immunotherapy Granulomatous Reactions Spectrum

Pembrolizumab-triggered Uveitis: An Additional Surrogate Marker for Responders in Melanoma Immunotherapy

Penicillinase-resistant antibiotics induce non-immune-mediated cholestasis through HSP27 activation associated with PKC/P38 and PI3K/AKT signaling pathways

Penicillin-induced immune-mediated hemolytic anemia in a horse

Pentamidine-induced hyperglycemia and ketosis in acquired immunodeficiency syndrome

Pentosan-induced thrombocytopenia: support for an immune complex mechanism

Percutaneous interventions in patients with immune-mediated heparin-induced thrombocytopenia

Perfluorinated compounds and immunotoxicity in children

Perfluorinated compounds: emerging POPs with potential immunotoxicity

Perfluoroalkyl acids: a review of monitoring and toxicological findings

Perfluoroalkyl and polyfluoroalkyl substances and indicators of immune function in children aged 12-19 y: National Health and Nutrition Examination Survey

Perfluorooctanoic acid induces mast cell-mediated allergic inflammation by the release of histamine and inflammatory mediators

Performance characteristics of an automated latex immunoturbidimetric assay [HemosIL HIT-Ab(PF4-H)] for the diagnosis of immune heparin-induced thrombocytopenia

Performance characteristics of two commercially available IgG-specific immunoassays in the assessment of heparin-induced thrombocytopenia (HIT)

Pericardial effusion due to pembrolizumab-induced immunotoxicity: A case report and literature review

Perinatal 192 IgG-Saporin as Neuroteratogen

Perinatal exposure of mice to TCDD decreases allergic sensitisation through inhibition of IL-4 production rather than T regulatory cell-mediated suppression

Perinatal exposure to a low dose of bisphenol A impaired systemic cellular immune response and predisposes young rats to intestinal parasitic infection

Perinatal exposure to insecticide methamidophos suppressed production of proinflammatory cytokines responding to virus infection in lung tissues in mice

Perinatal exposure to low doses of PCB 153 and PCB 126 affects maternal and neonatal immunity in goat kids

Perinatal TCDD exposure and the adult onset of autoimmune disease

Periodontitis induced by *Porphyromonas gingivalis* drives periodontal microbiota dysbiosis and insulin resistance via an impaired adaptive immune response

Perioperative agranulocytosis induced by immunosuppressants in a renal graft recipient : a case report

Perioperative recombinant human granulocyte colony-stimulating factor (Filgrastim) treatment prevents immunoinflammatory dysfunction associated with major surgery

Perioperative single high dose ATG-Fresenius S administration as induction immunosuppressive therapy in cadaveric renal transplantation--preliminary results

Perioperative supplementation of EPA reduces immunosuppression induced by postoperative chemoradiation therapy in patients with esophageal cancer

Permethrin-induced oxidative stress and toxicity and metabolism. A review

Persistent perfluorinated compounds cause immunotoxic effects

Persistent effect of in utero meso-2,3-dimercaptosuccinic acid (DMSA) on immune function and lead-induced immunotoxicity

Persistent hematologic and immunologic disturbances in 8-year-old Dutch children associated with perinatal dioxin exposure

Persistent hypogammaglobulinemia due to immunoglobulin class switch impairment by peri-transplant rituximab therapy

Persistent immunotoxic effects of tributyltin in human natural killer cells can be reversed by interleukin 2

Persistent sub-lethal chlorine exposure augments temperature induced immunosuppression in *Cyprinus carpio* advanced fingerlings

Persistent suppression of contact hypersensitivity, and altered T-cell parameters in F344 rats exposed perinatally to 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD)

Persistent suppression of humoral immunity produced by 7,12-dimethylbenz(A)anthracene (DMBA) in B6C3F1 mice: correlation with changes in spleen cell surface markers detected by flow cytometry

Pesticide immunotoxicity

Pesticide induced immunotoxicity in humans: a comprehensive review of the existing evidence

Pesticide mixtures potentiate the toxicity in murine thymocytes

Pesticide-induced immunotoxicity: are Great Lakes residents at risk

PFOS mediates immunomodulation in an avian cell line that can be mitigated via a virus infection

Pharmaceuticals, personal care products and endocrine-disrupting chemicals in U.S. surface and finished drinking waters: a proposed ranking system

Pharmacokinetic and nephroprotective benefits of using Schisandra chinensis extracts in a cyclosporine A-based immune-suppressive regime

Pharmacological correction of immuno-metabolic disorders with the peptide Gly-His-Lys in hepatic damage induced by tetrachloromethane

Phase I study of lenalidomide and alemtuzumab in refractory chronic lymphocytic leukaemia: maintaining immune functions during therapy-induced immunosuppression

Phenprocoumon-induced hepatitis as an immunologically mediated drug allergic complication of antithrombotic therapy

Phenytoin- and cranial radiotherapy-induced toxic epidermal necrolysis treated with combination therapy: systemic steroid and intravenous immunoglobulin

Phosphoproteomic analysis of mouse thymoma cells treated with tributyltin oxide: TBTO affects proliferation and energy sensing pathways

Physical activity and stress resistance: sympathetic nervous system adaptations prevent stress-induced immunosuppression

Physiologic basis of steroid-induced immunosuppression. Effects on lymphocyte migration to lymph nodes

Physiological levels of 5 alpha-dihydrotestosterone depress wound immune function and impair wound healing following trauma-hemorrhage

Physiological stress and innate immune response in gilthead sea bream (*Sparus aurata*) and sea bass (*Dicentrarchus labrax*) exposed to combination of trimethoprim and sulfamethoxazole (TMP-SMX)

PI3K-AKT signaling pathway is involved in hypoxia/thermal-induced immunosuppression of small abalone *Haliotis diversicolor*

Pigs with Severe Combined Immunodeficiency Are Impaired in Controlling Influenza A Virus Infection

Pine resin and biopin ointment: immunotoxic and allergenic activity

Pioglitazone blocks ethanol induction of microglial activation and immune responses in the hippocampus, cerebellum, and cerebral cortex in a mouse model of fetal alcohol spectrum disorders

Piperacillin/tazobactam-induced immune-mediated thrombocytopenia in the intensive care unit

Piperacillin-induced immune hemolysis: new cases and a concise review of the literature

Piracetam-induced immune thrombocytopenia

PK/TD modeling for prediction of the effects of 8C2, an anti-topotecan mAb, on topotecan-induced toxicity in mice

Plasma organochlorine levels and risk of non-Hodgkin lymphoma in a cohort of men

Platelet activating factor receptor binding plays a critical role in jet fuel-induced immune suppression

Platelet-activating factor does not mediate UVB-induced local immune suppression

Platelet-activating factor is crucial in psoralen and ultraviolet A-induced immune suppression, inflammation, and apoptosis

Plutonium-induced proliferative lesions and pulmonary epithelial neoplasms in the rat: immunohistochemical and ultrastructural evidence for their origin from type II pneumocytes

PM 2.5 collected in a residential area induced Th1-type inflammatory responses with oxidative stress in mice

Pneumatosis intestinalis associated with immune-suppressive agents in a case of minimal change disease

Pneumocystis jirovecii pneumonia in HIV-negative patients: a prospective study with focus on immunosuppressive drugs and markers of immune impairment

Poly(D,L-Lactic Acid) Nanoparticle Size Reduction Increases Its Immunotoxicity

Polybrominated biphenyls in model and environmentally contaminated human blood: protein binding and immunotoxicological studies

Polybrominated diphenyl ethers: human tissue levels and toxicology

Polycarbonate and polystyrene nanoplastic particles act as stressors to the innate immune system of fathead minnow (Pimephales promelas)

Polychlorinated biphenyl immunotoxicity: dependence on isomer planarity and the Ah gene complex

Polychlorinated biphenyls (PCBs) modulate both phagocytosis and NK cell activity in vitro in juvenile loggerhead sea turtles (Caretta caretta)

Polychlorinated diphenylethers: origin, analysis, distribution, and toxicity in the marine environment

Polyclonal antibodies against formaldehyde-modified apolipoprotein A-I. An approach to circumventing fixation-induced loss of antigenicity in immunocytochemistry

Polycyclic aromatic hydrocarbon-induced oxidative stress and lipid peroxidation in relation to immunological alteration

Polymeric Nanoparticles Induce NLRP3 Inflammasome Activation and Promote Breast Cancer Metastasis

Polymorphisms in genes involved in innate immunity and susceptibility to benzene-induced hematotoxicity

Polyomavirus disease under new immunosuppressive drugs: a cause of renal graft dysfunction and graft loss

Polypodium leucotomos inhibits ultraviolet B radiation-induced immunosuppression

Pompholyx induced by intravenous immunoglobulin therapy

Popliteal lymph node responses to acetone and ethanol differ from those induced by streptozotocin

Poppers: large cancer increase and immune suppression in animal tests

Porphyromonas gingivalis-impaired innate immune response via NLRP3 proteolysis in endothelial cells

Posited mechanisms of metal immunotoxicity

Possible association between the observed cyclosporine-induced upregulation of in vitro immune functions and posttransplantation lymphoproliferative disorders

Possible effects of repeated exposure to ibuprofen and acetaminophen on the intestinal immune response in young infants

Possible enhancing effect of the immunosuppressive agent, 6-mercaptopurine(6-MP) on focal lesion development in cirrhotic liver induced by carbon tetrachloride but not furfural in F344 rats

Possible functional immunotoxicity of acrylonitrile (VCN)

Possible induction of oral allergy syndrome during specific immunotherapy in patients sensitive to tree pollen

Possible involvement of stress hormones and hyperglycaemia in chronic mild stress-induced impairment of immune functions in diabetic mice

Possible mechanisms of immunotoxicity following in vivo exposure to the inhalant, isobutyl nitrite

Possible protective effect of royal jelly against cyclophosphamide induced prostatic damage in male albino rats; a biochemical, histological and immuno-histo-chemical study

Posterior reversible encephalopathy induced by intravenous immunoglobulin

Post-immune UV irradiation induces Tr1-like regulatory T cells that suppress humoral immune responses

Postoperative hepatitis B virus reactivation and surgery-induced immunosuppression in patients with hepatitis B-related hepatocellular carcinoma

Postoperative immune suppression in visceral surgery: characterisation of an intestinal mouse model

Postoperative period in cardiac surgery: don't forget heparin induced thrombocytopenia

Posttransplant lymphoproliferative disorders in liver recipients treated with OKT3 or ALG induction immunosuppression

Posttraumatic stress symptoms, intrusive thoughts, loss, and immune function after Hurricane Andrew

Postweaning exposure to aflatoxin results in impaired child growth: a longitudinal study in Benin, West Africa

Potassium sorbate suppresses intestinal microbial activity and triggers immune regulation in zebrafish (*Danio rerio*)

Potent immunosuppressive activity of a phosphodiesterase-4 inhibitor N-acylhydrazone in models of lipopolysaccharide-induced shock and delayed-type hypersensitivity reaction

Potent suppression of the adaptive immune response in mice upon dietary exposure to the potent peroxisome proliferator, perfluorooctanoic acid

Potential ability for metallothionein and vitamin E protection against cadmium immunotoxicity in head kidney and spleen of grass carp (*Ctenopharyngodon idellus*)

Potential for amelioration of aflatoxin B1-induced immunotoxic effects in progeny of White Leghorn breeder hens co-exposed to vitamin E

Potential for immunotoxicity due to environmental exposure to ultraviolet radiation

Potential for methotrexate exposure through contamination during parenteral use as an immunosuppressant

Potential immunotoxic effects of trichloroethylene-induced IV allergic reaction in renal impairment

Potential impact of quercetin and idebenone against immuno-inflammatory and oxidative renal damage induced in rats by titanium dioxide nanoparticles toxicity

Potential preventive role of lactic acid bacteria against aflatoxin M immunotoxicity and genotoxicity in mice

Potentially increased sensitivity of pregnant and lactating female rats to immunotoxic agents

Potentiation of the immunotoxicity of ethanol by acetaminophen in mice

Potentiation of the maternal immune system may modify the apoptotic process in embryos exposed to developmental toxicants

Pre-acclimation to low copper mitigated immunotoxic effects in spleen and head-kidney of large yellow croaker (*Pseudosciaena crocea*) when exposed subsequently to high copper

Pre-clinical immunotoxicity studies of nanotechnology-formulated drugs: Challenges, considerations and strategy

Prediction of drug-induced immune-mediated hepatotoxicity using hepatocyte-like cells derived from human embryonic stem cells

Pre-existing yellow fever immunity impairs and modulates the antibody response to tick-borne encephalitis vaccination

Pre-hatching fluoxetine-induced neurochemical, neurodevelopmental, and immunological changes in newly hatched cuttlefish

Preliminary study on immunosuppressive effects evaluation of biomaterial

Prenatal alcohol exposure and fetal programming: effects on neuroendocrine and immune function

Prenatal alcohol exposure selectively suppresses cell-mediated but not humoral immune responsiveness

Prenatal ambient air exposure to polycyclic aromatic hydrocarbons and the occurrence of respiratory symptoms over the first year of life

Prenatal and infant exposure to thimerosal from vaccines and immunoglobulins and risk of autism

Prenatal and postnatal cigarette and cannabis exposure: Effects on Secretory Immunoglobulin A in early childhood

Prenatal cadmium exposure alters postnatal immune cell development and function

Prenatal cadmium exposure produces persistent changes to thymus and spleen cell phenotypic repertoire as well as the acquired immune response

Prenatal exposure of Balb/c mice to ochratoxin A: effects on the immune system in the offspring

Prenatal exposure to bacterial endotoxin reduces the number of GAD67- and reelin-immunoreactive neurons in the hippocampus of rat offspring

Prenatal exposure to perfluorinated chemicals and relationship with allergies and infectious diseases in infants

Pre-natal exposure to perfluoroalkyl substances may be associated with altered vaccine antibody levels and immune-related health outcomes in early childhood

Prenatal exposure to polychlorinated biphenyls and dioxins from the maternal diet may be associated with immunosuppressive effects that persist into early childhood

Prenatal Exposure to Soy Isoflavones Altered the Immunological Parameters in Female Rats

Prenatal HgCl<sub>2</sub> exposure in BALB/c mice: gender-specific effects on the ontogeny of the immune system

Prenatal immunotoxicant exposure and postnatal autoimmune disease

Prenatal paracetamol exposure and risk of asthma and elevated immunoglobulin E in childhood

Prenatal PCB exposure and thymus size at birth in neonates in Eastern Slovakia

Prenatal pesticide exposure and otitis media during early childhood in the PELAGIE mother-child cohort

Prenatal TCDD causes persistent modulation of the postnatal immune response, and exacerbates inflammatory disease, in 36-week-old lupus-like autoimmune SNF1 mice

Pre-natal/juvenile chlorpyrifos exposure associated with immunotoxicity in adulthood in Swiss albino mice

Preoperative detection and management of immune heparin-induced thrombocytopenia in patients undergoing heart surgery with iloprost

Preservation of carrageenan-induced immune suppression with alleviation of toxicity in aprotinin-treated mice

Pretilachlor has the potential to induce endocrine disruption, oxidative stress, apoptosis and immunotoxicity during zebrafish embryo development

Pretreatment with 1,8-cineole potentiates thioacetamide-induced hepatotoxicity and immunosuppression

Prevalence of kidney dysfunction in humans - relationship to cadmium dose, metallothionein, immunological and metabolic factors

Prevention by alpha-difluoromethylornithine of skin carcinogenesis and immunosuppression induced by ultraviolet irradiation

Prevention by chitosan of myelotoxicity, gastrointestinal toxicity and immunocompetent organic toxicity induced by 5-fluorouracil without loss of antitumor activity in mice

Prevention of anti-T-cell receptor alpha beta monoclonal antibody-induced side-effects by treatment with cyclosporin A without interference of monoclonal antibody-induced immunosuppression in mice

Prevention of excessive collagen accumulation by human intravenous immunoglobulin treatment in a murine model of bleomycin-induced scleroderma

Prevention of glucocorticoid-induced osteoporosis in immunobullous diseases with alendronate: a randomized, double-blind, placebo-controlled study

Prevention of graft-versus-host disease and the induction of transplant tolerance by low-dose UV-B irradiation of BM cells combined with cyclosporine immunosuppression

Prevention of ochratoxin A-induced oxidative stress-mediated apoptotic processes and impairment of embryonic development in mouse blastocysts by liquiritigenin

Prevention of photocarcinogenesis and UV-induced immunosuppression in mice by topical tannic acid

Prevention of solar-induced immunosuppression by a new highly protective broadspectrum sunscreen

Prevention of ultraviolet radiation-induced immunosuppression by (-)-epigallocatechin-3-gallate in mice is mediated through interleukin 12-dependent DNA repair

Prevention of ultraviolet radiation-induced immunosuppression by sunscreen in Candida albicans-induced delayed-type hypersensitivity

Preventive and therapeutic effects of sugar cane extract on cyclophosphamide-induced immunosuppression in chickens



Preventive effect of a traditional herbal medicine, Hochu-ekki-to, on immunosuppression induced by surgical stress

Preventive effect of immunosuppressive agents against indomethacin-induced small intestinal ulcers in rats

Preventive effects of diphenyl dimethyl dicarboxylate on the immunotoxicity of carbon tetrachloride in ICR mice

Primary antibody response to keyhole limpet hemocyanin in rat as a model for immunotoxicity evaluation

Primary concept of nickel toxicity - an overview

Prior immunity to homologous and heterologous Salmonella serotypes suppresses local and systemic anti-fragment C antibody responses and protection from tetanus toxin in mice immunized with Salmonella strains expressing fragment C

Prior or coinstantaneous oral exposure to environmental immunosuppressive agents aggravates mite allergen-induced atopic dermatitis-like immunoreaction in NC/Nga mice

Prior oral exposure to environmental immunosuppressive chemicals methoxychlor, parathion, or piperonyl butoxide aggravates allergic airway inflammation in NC/Nga mice

Pro- and anti-inflammatory cytokine expression in carp blood and head kidney leukocytes exposed to cyanotoxin stress--an in vitro study

Proapoptotic and immunotoxic effects of sulfur-fumigated polysaccharides from Smilax glabra Roxb. in RAW264.7 cells

Probable corticosteroid-induced reactivation of latent hepatitis B virus infection in an HIV-positive patient involving immune escape

Problems and challenges in the development and validation of human cell-based assays to determine nanoparticle-induced immunomodulatory effects

Proceedings: Does radiation induced immunosuppression influence distant metastases in mammary carcinoma

Processed Aloe vera gel ameliorates cyclophosphamide-induced immunotoxicity

Profiling the immunotoxicity of chemicals based on in vitro evaluation by a combination of the Multi-ImmunoTox assay and the IL-8 Luc assay

Profound impairment of adaptive immune responses by alkylating chemotherapy

Progression of renal dysfunction in cardiac transplantation after the introduction of everolimus in the immunosuppressive regime

Prolonged ceftriaxone-induced immune thrombocytopenia due to impaired drug clearance: a case report

Prolonged exposure to progesterone prevents induction of protective mucosal responses following intravaginal immunization with attenuated herpes simplex virus type 2

Prolonged Impairment of Immunological Memory After Anti-CD20 Treatment in Pediatric Idiopathic Nephrotic Syndrome

Promoting effects of nanoparticles/materials on sensitive lung inflammatory diseases

Propanil exposure induces delayed but sustained abrogation of cell-mediated immunity through direct interference with cytotoxic T-lymphocyte effectors

Properties of Gluten Intolerance: Gluten Structure, Evolution, Pathogenicity and Detoxification Capabilities

Prophylactic ganciclovir is more effective in HLA-identical family member marrow transplant recipients than in more heavily immune-suppressed HLA-identical unrelated donor marrow transplant recipients. Australasian Bone Marrow Transplant Study Group

Prophylactic role of alpha-lipoic acid and vitamin E against zinc oxide nanoparticles induced metabolic and immune disorders in rat's liver

Prophylactic role of coenzyme Q10 and Cynara scolymus L on doxorubicin-induced toxicity in rats: Biochemical and immunohistochemical study

Prophylaxis with respiratory syncytial virus F-specific humanized monoclonal antibody delays and moderately suppresses the native antibody response but does not impair immunity to late rechallenge

Propranolol Impairs Primary Immune Responses in Rat Experimental Autoimmune Encephalomyelitis

Propylthiouracil-induced diffuse proliferative lupus nephritis: review of immunological complications

Prospects for treatment of paraquat-induced lung fibrosis with immunosuppressive drugs and the need for better prediction of outcome: a systematic review

Prostaglandin E1 attenuates impairment of cellular immunity after cardiopulmonary bypass

Prostaglandin E2 Promotes UV radiation-induced immune suppression through DNA hypermethylation

Prostaglandin E2-Mediated Impairment of Innate Immune Response to A(H1N1)pdm09 Infection in Diet-Induced Obese Mice Could Be Restored by Paracetamol

Prostaglandin E2-prostaglandin E receptor subtype 4 (EP4) signaling mediates UV irradiation-induced systemic immunosuppression

Protamine-induced fatal anaphylaxis. Prevalence of anti-protamine immunoglobulin E antibody

Protecting effect of Chaenomeles speciosa broth on immunosuppressive mice induced by cyclophosphamide

Protection Against Lipopolysaccharide-Induced Immunosuppression by IgG and IgM

Protection against UV-induced systemic immunosuppression in mice by a single topical application of the antioxidant vitamins C and E

Protection from inflammation, immunosuppression and carcinogenesis induced by UV radiation in mice by topical Pycnogenol

Protection from JP-8 jet fuel induced immunotoxicity by administration of aerosolized substance P

Protection from O,O,S-trimethyl phosphorothioate-induced immune suppression

Protection to immune system of mice by N-acetyl tryptophan glucoside (NATG) against gamma radiation induced immune suppression

Protective antibody levels and dose requirements for IV 5% Nabi Hepatitis B immune globulin combined with lamivudine in liver transplantation for hepatitis B-induced end stage liver disease

Protective effect of a marine oligopeptide preparation from chum salmon (*Oncorhynchus keta*) on radiation-induced immune suppression in mice

Protective Effect of Alpha-Tocopherol in Deltamethrin Induced Immunotoxicity

Protective effect of an immune-modulating diet comprising whey peptides and fermented milk products on indomethacin-induced small-bowel disorders in rats

Protective effect of bioactive fraction of *Sphaeranthus indicus* Linn. against cyclophosphamide induced suppression of humoral immunity in mice

Protective effect of *Cassia occidentalis* L. on cyclophosphamide-induced suppression of humoral immunity in mice

Protective effect of centella triterpene saponins against cyclophosphamide-induced immune and hepatic system dysfunction in rats: its possible mechanisms of action

Protective effect of chitooligosaccharides against cyclophosphamide-induced immunosuppression in mice

Protective Effect of Chitosan Oligosaccharides Against Cyclophosphamide-Induced Immunosuppression and Irradiation Injury in Mice

Protective effect of curcumin against arsenic-induced apoptosis in murine splenocytes in vitro

Protective effect of *Devosia* sp. ANSB714 on growth performance, serum chemistry, immunity function and residues in kidneys of mice exposed to deoxynivalenol

Protective effect of *Echinacea purpurea* (Immulant) against cisplatin-induced immunotoxicity in rats

Protective effect of N-acetylcysteine against DNA damage and S-phase arrest induced by ochratoxin A in human embryonic kidney cells (HEK-293)

Protective effect of resveratrol and vitamin E against ethanol-induced oxidative damage in mice: biochemical and immunological basis

Protective effect of thymoquinone against diazinon-induced hematotoxicity, genotoxicity and immunotoxicity in rats

Protective effect of triterpenoid fractions from the rhizomes of *Astilbe chinensis* on cyclophosphamide-induced toxicity in tumor-bearing mice

Protective effect of Vitamin D3 against lead induced hepatotoxicity, oxidative stress, immunosuppressive and calcium homeostasis disorders in rat

Protective effects of [6]-paradol on histological lesions and immunohistochemical gene expression in DMBA induced hamster buccal pouch carcinogenesis

Protective effects of antioxidants on deoxynivalenol-induced damage in murine lymphoma cells

Protective effects of *Butea frondosa* leaves against stress induced immune impairment in sprague dawley rats

Protective effects of diallyl sulfide on N-nitrosodimethylamine-induced immunosuppression in mice

Protective effects of different antioxidants against endosulfan-induced oxidative stress and immunotoxicity in albino rats

Protective effects of gamma-irradiated *Astragalus* polysaccharides on intestinal development and mucosal immune function of immunosuppressed broilers

Protective effects of meat from lambs on selenium nanoparticle supplemented diet in a mouse model of polycyclic aromatic hydrocarbon-induced immunotoxicity

Protective effects of melatonin in reduction of oxidative damage and immunosuppression induced by aflatoxin B1-contaminated diets in young chicks

Protective effects of polysaccharides from *Cordyceps gunnii* mycelia against cyclophosphamide-induced immunosuppression to TLR4/TRAF6/NF-kappaB signalling in BALB/c mice

Protective effects of sea cucumber (*Holothuria atra*) extract on testicular dysfunction induced by immune suppressant drugs in Wistar rats

Protective effects of selenium against cadmium induced hematological disturbances, immunosuppressive, oxidative stress and hepatorenal damage in rats

Protective effects of selenium on mercury induced immunotoxic effects in mice by way of concurrent drinking water exposure

Protective Effects of Sporoderm-Broken Spores of *Ganderma lucidum* on Growth Performance, Antioxidant Capacity and Immune Function of Broiler Chickens Exposed to Low Level of Aflatoxin B

Protective effects of zymosan on heat stress-induced immunosuppression and apoptosis in dairy cows and peripheral blood mononuclear cells

Protective role of ascorbic acid isolated from *Cissus quadrangularis* on NSAID induced toxicity through immunomodulating response and growth factors expression

Protective role of bentonite against aflatoxin B1- and ochratoxin A-induced immunotoxicity in broilers

Protective role of quercetin against hematotoxic and immunotoxic effects of furan in rats

Protein A induced abrogation of cyclophosphamide toxicity is associated with concomitant potentiation of immune function of host

Protein malnutrition impairs the immune control of *Trichinella spiralis* infection

Protein malnutrition impairs the immune response and influences the severity of infection in a hamster model of chronic visceral leishmaniasis

Protein phosphatase 2A regulates innate immune and proteolytic responses to cigarette smoke exposure in the lung

Protein phosphorylation profiling identifies potential mechanisms for direct immunotoxicity

Proteins of 30 and 36 kilodaltons, membrane constituents of the *Staphylococcus aureus* L form, induce production of tumor necrosis factor alpha and activate the human immunodeficiency virus type 1 long terminal repeat

Protracted severe immune dysregulation induced by cardiopulmonary bypass: a predisposing etiologic factor in blood transfusion-related AIDS

*Pseudomonas aeruginosa* colonization causes PD-L1 overexpression on monocytes, impairing the adaptive immune response in patients with cystic fibrosis

*Pseudomonas aeruginosa* induces localized immunosuppression during pneumonia

*Pseudomonas aeruginosa* LasB protease impairs innate immunity in mice and humans by targeting a lung epithelial cystic fibrosis transmembrane regulator-IL-6-antimicrobial-repair pathway

Psoriatic skin lesions induced by tumor necrosis factor antagonist therapy: clinical features and possible immunopathogenesis

Psychological factors in immunomodulation induced by cancer surgery: a review

Psychosocial factors and secretory immunoglobulin A

*Pteromalus puparum* venom impairs host cellular immune responses by decreasing expression of its scavenger receptor gene

Pulmonary abnormalities and serum immunoglobulins in facsimile machine repair technicians exposed to butyl methacrylate fume

Pulmonary function and concentrations of alpha 1-antitrypsin and immunoglobulin E in workers exposed to pancreatic enzymes

Pulmonary glass particles may persist in the lung suppressing function of immune cells

Pulmonary immunotoxic potentials of metals are governed by select physicochemical properties: vanadium agents

Pulmonary immunotoxic potentials of metals are governed by select physicochemical properties: chromium agents

Pulmonary immunotoxicity of inhaled ammonium metavanadate in Fisher 344 rats

Pulmonary immunotoxicology of select metals: aluminum, arsenic, cadmium, chromium, copper, manganese, nickel, vanadium, and zinc

Pulmonary metastases, a potential biologic consequence of anesthetic-induced immunosuppression by thiopental

Pulmonary persistence of graphene nanoplatelets may disturb physiological and immunological homeostasis

Pulmonary toxicity following inhalation exposure to VX in anesthetized rats: Possible roles for compromised immunity and oxidative stress-induced lung injury

Pulse modulated 900 MHz radiation induces hypothyroidism and apoptosis in thyroid cells: a light, electron microscopy and immunohistochemical study

Pulsed' immunosuppressive therapy in the treatment of immunologically induced corneal and scleral disease

Pure red cell aplasia induced by lamivudine without the influence of zidovudine in a patient infected with human immunodeficiency virus

Purinergic system as a potential target for inflammation and toxicity induced by thymol in immune cells and tissues

Pyrethroids used indoors--immune status of humans exposed to pyrethroids following a pest control operation--a one year follow-up study

QSPR/QSAR models for prediction of the physico-chemical properties and biological activity of polychlorinated diphenyl ethers (PCDEs)

Quantitative and immunohistochemical evaluation of PCNA, androgen receptors, apoptosis, and Glutathione-S-Transferase P1 on preneoplastic changes induced by cadmium and zinc chloride in the rat ventral prostate

Quantitative assessment of mesangial immunoglobulin A (IgA) accumulation, elevated circulating IgA immune complexes, and hematuria during vomitoxin-induced IgA nephropathy

Quantitative modeling of suppression of IgG1, IgG2a, IL-2, and IL-4 responses to antigen in mice treated with exogenous corticosterone or restraint stress

Quantitative Proteomic Analysis Reveals That Arctigenin Alleviates Concanavalin A-Induced Hepatitis Through Suppressing Immune System and Regulating Autophagy

Quantitative relationships between the suppression of selected immunological parameters and the area under the corticosterone concentration vs. time curve in B6C3F1 mice subjected to exogenous corticosterone or to restraint stress

Quercetin mitigates Adriamycin-induced anxiety- and depression-like behaviors, immune dysfunction, and brain oxidative stress in rats

Quinine-dependent, platelet-reactive monoclonals mimic antibodies found in patients with quinine-induced immune thrombocytopenia

Quinine-induced immune thrombocytopenia with hemolytic uremic syndrome: clinical and serological findings in nine patients and review of literature

Quinocetone induces mitochondrial apoptosis in HepG2 cells through ROS-dependent promotion of VDAC1 oligomerization and suppression of Wnt1/beta-catenin signaling pathway

Radiation-induced atypia of endocervical epithelium: a histological, immunohistochemical and cytometric study

Radiofrequency radiation and the immune system. Part 3. In vitro effects on human immunoglobulin and on murine T- and B-lymphocytes.

Radiofrequency-induced carcinogenesis: cellular calcium homeostasis changes as a triggering factor.

Radioprotection of EGCG based on immunoregulatory effect and antioxidant activity against <sup>60</sup>Co gamma radiation-induced injury in mice

Radioprotective effect of green tea and grape seed extracts mixture on gamma irradiation induced immune suppression in male albino rats

Radioprotective effects of honeybee venom (*Apis mellifera*) against 915-MHz microwave radiation-induced DNA damage in wistar rat lymphocytes: in vitro study.

Randomized controlled study comparing reduced calcineurin inhibitors exposure versus standard cyclosporine-based immunosuppression

Randomized controlled trial of oral omega-3 PUFA in solar-simulated radiation-induced suppression of human cutaneous immune responses

Randomized trial of basiliximab induction versus steroid therapy in pediatric liver allograft recipients under tacrolimus immunosuppression

Rapamycin (Sirolimus) inhibits vascular smooth muscle DNA synthesis in vitro and suppresses narrowing in arterial allografts and in balloon-injured carotid arteries: evidence that rapamycin antagonizes growth factor action on immune and nonimmune cells

Rapid and durable response to intravenous immunoglobulin in delayed heparin-induced thrombocytopenia: a case report

Rapid cytoskeleton modification in thymocytes induced by the immunotoxicant tributyltin

Rapid granulomatosis with polyangiitis induced by immune checkpoint inhibition

Rapid immunoassays for diagnosis of heparin-induced thrombocytopenia: Comparison of diagnostic accuracy, reproducibility, and costs in clinical practice

Rat extracorporeal circulation model for evaluation of systemic immunotoxicity

Re: Biochemical evaluation of immunotoxic effects of cadmium in rats

Re: Nitrofurantoin-induced pulmonary hemorrhage in a renal transplant recipient receiving immunosuppressive therapy: case report and review of the literature

Reaction of the immune system to low-level RF/MW exposures.

Reactions to rubella vaccine and persistence of antibody in virgin-soil populations after vaccination and wild-virus-induced immunization

Reactive oxygen species (ROS) induced cytokine production and cytotoxicity of PAMAM dendrimers in J774A.1 cells

Reactive oxygen species sources and biomolecular oxidative damage induced by aflatoxin B1 and fumonisin B1 in rat spleen mononuclear cells

Recent advances in understanding the mechanisms of TCDD immunotoxicity

Recent studies on the effects of fetal alcohol exposure on the endocrine and immune systems

Recent US Food and Drug Administration warnings on hepatitis B reactivation with immune-suppressing and anticancer drugs: just the tip of the iceberg

Recommendations issued by the Spanish Society of Pediatric Infectious Diseases for the follow-up of the child exposed to the human immunodeficiency virus and to antiretroviral drugs during pregnancy and the neonatal period

Reconstitution of cyclophosphamide-induced, impaired function of the immune system in animal models

Recovery from 4-(methylnitrosamino)-1-(3-pyridyl)-1-butanone-induced immunosuppression in A/J mice by treatment with nonsteroidal anti-inflammatory drugs

Recovery of dimethylnitrosamine-induced immunosuppression by pargyline in the mixed cultures of murine hepatocytes and splenocytes

Recurrence of digoxin toxicity following treatment with digoxin immune fab in a patient with renal impairment

Recurrent ibuprofen-induced aseptic meningitis: evidence against an antigen-specific immune response

Reduced antibody responses to vaccinations in children exposed to polychlorinated biphenyls

Reduced birth defects caused by maternal immune stimulation in methylnitrosourea-exposed mice: association with placental improvement

Reduced expression of selective immune-related genes in silver catfish (*Rhamdia quelen*) monocytes exposed to atrazine

Reduction in peripheral lymphocytes and thymus atrophy induced by organotin compounds in vivo

Reduction in valproic acid-induced neural tube defects by maternal immune stimulation: role of apoptosis



Regression of arteriosclerotic lesions induced by immunization with heat shock protein 65-containing material in normocholesterolemic, but not hypercholesterolemic, rabbits

Regular consumption of a silicic acid-rich water prevents aluminium-induced alterations of nitroergic neurons in mouse brain: histochemical and immunohistochemical studies

Regulation of isocyanate-induced apoptosis, oxidative stress, and inflammation in cultured human neutrophils: isocyanate-induced neutrophils apoptosis

Regulation of prostaglandin endoperoxide H synthase-2 expression by 2,3,7,8-tetrachlorodibenzo-p-dioxin

Regulatory effects of interleukin-6 in immunoglobulin G immune-complex-induced lung injury

Regulatory T cells ameliorate acetaminophen-induced immune-mediated liver injury

Regulatory T cells in gamma irradiation-induced immune suppression

Regulatory T-cells promote pulmonary repair by modulating T helper cell immune responses in lipopolysaccharide-induced acute respiratory distress syndrome

Relating cytotoxicity, zinc ions, and reactive oxygen in ZnO nanoparticle-exposed human immune cells

Relationship between Ah receptor-mediated polychlorinated biphenyl (PCB)-induced humoral immunosuppression and thymic atrophy

Relationship of trichothecene structure to COX-2 induction in the macrophage: selective action of type B (8-keto) trichothecenes

Relationships between acrylamide and glycidamide hemoglobin adduct levels and allergy-related outcomes in general US population, NHANES 2005-2006

Relationships between polychlorinated biphenyls and health status in harbor porpoises (*Phocoena phocoena*) stranded in the United Kingdom

Relative potency based on hepatic enzyme induction predicts immunosuppressive effects of a mixture of PCDDS/PCDFS and PCBS

Relative sensitivities of 2,3,7,8-tetrachlorodibenzo-p-dioxin-induced Cyp1a-1 and Cyp1a-2 gene expression and immunotoxicity in female B6C3F1 mice

Relative sensitivity of developmental and immune parameters in juvenile versus adult male rats after exposure to di(2-ethylhexyl) phthalate

Release of Danger-Associated Molecular Patterns following Chemotherapy Does Not Induce Immunoparalysis in Leukemia Patients

Remission of severe disease induced by CMV and lymphoproliferative post-transplant disease after changing the immunosuppressive regime

Renal dysfunction following adult intestinal transplant under tacrolimus-based immunosuppression

Renin immunochemistry, sodium excretion and relative heart weight in cyclosporine- or alimentary-induced magnesium deficiency in rats

Repeated dose 90-day oral toxicity test of G-7% NANA in rats: An application of new criterion for toxicity determination to test article-induced changes

Repeated high dose oral exposure or continuous subcutaneous infusion of 2-methoxyacetic acid does not suppress humoral immunity in the mouse

Repeated ozone exposure exacerbates insulin resistance and activates innate immune response in genetically susceptible mice

Repeated-oral dose toxicity of polyethylene microplastics and the possible implications on reproduction and development of the next generation

Reply to the commentary by Onuigbo on the article "Exposure to inhibitors of the renin-angiotensin system is a major independent risk factor for acute renal failure induced by sucrose-containing intravenous immunoglobulins. A case-control study

Report of validation study of assessment of direct immunotoxicity in the rat. The ICICIS Group Investigators. International Collaborative Immunotoxicity Study

Reproductive and developmental toxicity of organotin compounds

Rescheduling the process of nanoparticle removal used for water mercury remediation can increase the risk to aquatic organism: evidence of innate immune functions modulation in European eel (*Anguilla anguilla* L

Resistance of a lizard (the green anole, *Anolis carolinensis*; Polychridae) to ultraviolet radiation-induced immunosuppression

Resistance of C57Bl/6 mice to immunosuppressive effects of aflatoxin B1 and relationship with neuroendocrine mechanisms

Resistance to acute silicosis in senescent rats: role of alveolar macrophages

Resistance to erythropoietin: immunohemolytic anemia induced by residual formaldehyde in dialyzers

Resistance to intestinal parasites during murine AIDS: role of alcohol and nutrition in immune dysfunction

Resolution of ipilimumab induced severe hepatotoxicity with triple immunosuppressants therapy

Respiratory and immune response to maximal physical exertion following exposure to secondhand smoke in healthy adults

Respiratory toxicity and immunotoxicity evaluations of microparticle and nanoparticle C60 fullerene aggregates in mice and rats following nose-only inhalation for 13 weeks

Restoration of cyclophosphamide-suppressed primary immune response in mice by Tolpa Peat Preparation (TPP)--comparison with calf thymus extract

Restorative effect of l-Dopa treatment against Ochratoxin A induced neurotoxicity

Results and long-term risks of immuno-suppressive treatment in chronic juvenile arthritis. Apropos of 40 cases

Resveratrol (3,5,4'-trihydroxystilbene) protects pregnant mother and fetus from the immunotoxic effects of 2,3,7,8-tetrachlorodibenzo-p-dioxin

Resveratrol Pretreatment Attenuates Concanavalin A-induced Hepatitis through Reverse of Aberration in the Immune Response and Regenerative Capacity in Aged Mice

Retinal laser burn-induced neuropathy leads to substance P-dependent loss of ocular immune privilege

RETRACTED: Composition and immunotoxicity activity of major essential oils from stems of *Allium victorialis* L. var. *platyphyllum* Makino against *Aedes aegypti* L

RETRACTED: Immunotoxicity activity from various organic solvents extract of *Allium* genus from South Korea against *Aedes aegypti* L

RETRACTED: Major essential oils composition and immunotoxicity activity from leaves of *Foeniculum vulgare* against *Aedes aegypti* L

Retrospective evaluation of the impact of functional immunotoxicity testing on pesticide hazard identification and risk assessment

Reversibility of alcohol-induced immune depression

Reversibility of thymulin production impairment by L-arginine supplementation in mice exposed to inorganic mercury

Review on the toxicity, occurrence, metabolism, detoxification, regulations and intake of zearalenone: an oestrogenic mycotoxin

Rhein Induces Oxidative Stress and Apoptosis in Mouse Blastocysts and Has Immunotoxic Effects during Embryonic Development

Rhinovirus exposure impairs immune responses to bacterial products in human alveolar macrophages

Rhinovirus impairs the immune response of alveolar macrophages to facilitate *Streptococcus pneumoniae* infection

Rifabutin-induced uveitis in an immunocompetent subject

Rifampicin-induced immune thrombocytopenia

Rifampicin-induced suppression of antitumor immunity

RIP1 and RIP3 contribute to Tributyltin-induced toxicity invitro and invivo

Risk assessment and immunotoxicology

Risk assessment for possible effects of 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD) and related substances on components and functions of the immune system

Risk assessment for the harmful effects of immunotoxic agents on the immunological resistance to infectious diseases: ultraviolet light as an example

Risk assessment for the harmful effects of UVB radiation on the immunological resistance to infectious diseases

Risk factors for immunologically mediated disease in workers with respiratory symptoms when exposed to hexahydrophthalic anhydride

Risk of developing certain malignancies is related to duration of immunosuppressive drug exposure in patients with rheumatic diseases

Risk of infections during the first year of life after in utero exposure to drugs acting on immunity: A population-based cohort study

Risks of novel therapeutics: gonococcemia in an immune-suppressed patient receiving eculizumab

Risperidone-induced immunoallergic hepatitis

Ritanserine counteracts both rat vacuous chewing movements and nigro-striatal tyrosine hydroxylase-immunostaining alterations induced by haloperidol

Rituximab impairs immunoglobulin (Ig)M and IgG (subclass) responses after influenza vaccination in rheumatoid arthritis patients

Rituximab treatment results in impaired secondary humoral immune responsiveness

Rituximab-induced accelerated cryoprecipitation in hepatitis C virus-associated mixed cryoglobulinemia has parallels with intravenous immunoglobulin-induced immune complex deposition in mixed cryoglobulinemia: comment on the article by Sene et al

Rituximab-induced immunodysregulatory ileocolitis in a patient with follicular lymphoma

Rituximab-induced interstitial lung disease in a patient with aquaporin-4 immunoglobulin G-positive neuromyelitis optica spectrum disorder

Rituximab-induced interstitial lung disease in a patient with immune thrombocytopenia purpura

Rituximab-induced serum sickness in refractory immune thrombocytopenic purpura

RNA-Seq revealed the impairment of immune defence of tilapia against the infection of *Streptococcus agalactiae* with simulated climate warming

Role of aflatoxin toxicity on transmissibility and pathogenicity of H9N2 avian influenza virus in turkeys

Role of  $\alpha 7$ -Nicotinic Acetylcholine Receptors of B Cells in the Immunotoxic Effect of Organophosphorus Compounds

Role of alterations in  $Ca^{2+}$ -associated signaling pathways in the immunotoxicity of polycyclic aromatic hydrocarbons

Role of altered arachidonic acid metabolism in 2,3,7, 8-tetrachlorodibenzo-p-dioxin-induced immune suppression in C57Bl/6 mice

Role of corticosteroids in cadmium induced immunotoxicity

Role of corticosterone in ethyl carbamate-induced immunosuppression in female BALB/c mice

Role of corticosterone in immunosuppressive effects of acute ethanol exposure on Toll-like receptor mediated cytokine production

Role of cyclooxygenase-2 in deoxynivalenol-induced immunoglobulin a nephropathy

Role of decomposition products in sodium methyldithiocarbamate-induced immunotoxicity

Role of Fas apoptosis and MHC genes in 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD)-induced immunotoxicity of T cells

Role of Fas-Fas ligand interactions in 2,3,7,8-tetrachlorodibenzo- p-dioxin (TCDD)-induced immunotoxicity: increased resistance of thymocytes from Fas-deficient (lpr) and Fas ligand-defective (gld) mice to TCDD-induced toxicity

Role of glutathione and reactive oxygen intermediates in 2,3,7,8-tetrachlorodibenzo-p-dioxin-induced immune suppression in C57Bl/6 mice

Role of Glutathione Conjugation in 1-Bromobutane-induced Immunotoxicity in Mice

Role of glutathione conjugation in the hepatotoxicity and immunotoxicity induced by 1-bromopropane in female BALB/c mice

Role of hepatoprotectors and immunomodulators in regulation of hepatocyte apoptosis induced by antituberculosis treatment

Role of IL-17 Pathways in Immune Privilege: A RNA Deep Sequencing Analysis of the Mice Testis Exposure to Fluoride

Role of Immune Status in Chemotherapy-Induced Transient Acantholytic Dermatitis

Role of metabolism by flavin-containing monooxygenase in thioacetamide-induced immunosuppression

Role of metabolism in cocaine-induced immunosuppression in splenocyte cultures from B6C3F1 female mice

Role of metabolism in dimethylnitrosamine-induced immunosuppression: a review

Role of metabolism in monocrotaline-induced immunotoxicity in C57BL/6 mice

Role of metabolism in parathion-induced hepatotoxicity and immunotoxicity

Role of nutrients in alcohol-induced immunomodulation

Role of oxidative stress in cadmium toxicity and carcinogenesis

Role of Oxidative Stress in the Suppression of Immune Responses in Peripheral Blood Mononuclear Cells Exposed to Combustible Tobacco Product Preparation

Role of PLC-PIP2 and cAMP-PKA signal pathways in radiation-induced immune-suppressing effect

Role of P-type ATPase metal transporters and plant immunity induced by jasmonic acid against Lead (Pb) toxicity in tomato

Role of quinones in toxicology

Role of regulatory T cells in the induction of atopic dermatitis by immunosuppressive chemicals

Role of RT6+ T lymphocytes in mercury-induced renal autoimmunity: experimental manipulations of "susceptible" and "resistant" rats

Role of Th1 and Th2 lymphocytes and cytokines produced by these cells in suppression of immune reactions during subacute poisoning with anticholinesterase toxicants

Role of the Ah locus in suppression of cytotoxic T lymphocyte activity by halogenated aromatic hydrocarbons (PCBs and TCDD): structure-activity relationships and effects in C57Bl/6 mice congenic at the Ah locus

Roles and mechanisms of human immunodeficiency virus protease inhibitor ritonavir and other anti-human immunodeficiency virus drugs in endothelial dysfunction of porcine pulmonary arteries and human pulmonary artery endothelial cells

Roles of dosage, pharmacokinetics, and cellular sensitivity to damage in the selective toxicity of cyclophosphamide towards B and T cells in development

Roles of ROS mediated oxidative stress and DNA damage in 3-methyl-2-quinoxalin benzenevinylketo-1, 4-dioxide-induced immunotoxicity of Sprague-Dawley rats

Rotavirus induces a biphasic enterotoxic and cytotoxic response in human-derived intestinal enterocytes, which is inhibited by human immunoglobulins

Royal jelly-induced asthma and anaphylaxis: clinical characteristics and immunologic correlations

RSV-specific anti-viral immunity is disrupted by chronic ethanol consumption

Safe administration of rotavirus vaccine in a cohort of infants exposed to immunosuppressive drugs during gestation

Safe Conversion From Tacrolimus to Belatacept in High Immunologic Risk Kidney Transplant Recipients With Allograft Dysfunction

Safe-by-Design of Glucan Nanoparticles: Size Matters When Assessing the Immunotoxicity

Safety and immunologic benefits of conversion to sirolimus in kidney transplant recipients with long-term exposure to calcineurin inhibitors

Safety of belatacept bridging immunosuppression in hepatitis C-positive liver transplant recipients with renal dysfunction

Safety of modified vaccinia virus Ankara (MVA) in immune-suppressed macaques

Salecan protected against concanavalin A-induced acute liver injury by modulating T cell immune responses and NMR-based metabolic profiles

Salinity change impairs pipefish immune defence

Sarcodon imbricatus polysaccharides protect against cyclophosphamide-induced immunosuppression via regulating Nrf2-mediated oxidative stress

Sarin (GB, O-isopropyl methylphosphonofluoridate) neurotoxicity: critical review

Schistosoma mansoni infection impairs antimalaria treatment and immune responses of rhesus macaques infected with mosquito-borne Plasmodium coatneyi

Search for biomarkers of asbestos exposure and asbestos-induced cancers in investigations of the immunological effects of asbestos

Seasonal variations in the level of immunoglobulins and serum proteins of children differing by exposure to air-borne lead

Secondhand Smoke Induces Inflammation and Impairs Immunity to Respiratory Infections

Secretion of interferon gamma from human immune cells is altered by exposure to tributyltin and dibutyltin

Secretion of interleukin-10 from murine colon carcinoma cells suppresses systemic antitumor immunity and impairs protective immunity induced against the tumors

Selected new developments in asbestos immunotoxicity

Selective immunolesioning of the basal forebrain cholinergic system disrupts short-term memory in rats

Selective immunotoxic effects in mice treated with the adenosine deaminase inhibitor 2'-deoxycoformycin

Selective increase in astrocytic elements in the rat dentate gyrus after chronic toluene exposure studied by GFAP immunocytochemistry and electron microscopy

Selective inhibition of B lymphocytes in TBTC-treated human bone marrow long-term culture

Selective modulation of immune function resulting from in vitro exposure to methylenedioxymethamphetamine (Ecstasy)

Selective myelotoxicity of propanil

Selective suppression of in vitro T-dependent humoral immunity by synthetic food additive antioxidants

Selective, prolonged alteration of complement-mediated immune clearance after acute exposure of mice to ethanol

Selenium Alleviates Aflatoxin B-Induced Immune Toxicity through Improving Glutathione Peroxidase 1 and Selenoprotein S Expression in Primary Porcine Splenocytes

Selenium Deficiency Aggravates Aflatoxin B1-Induced Immunotoxicity in Chick Spleen by Regulating 6 Selenoprotein Genes and Redox/Inflammation/Apoptotic Signaling

Selenium deficiency impaired immune function of the immune organs in young grass carp (*Ctenopharyngodon idella*)

Selenium deficiency impairs host innate immune response and induces susceptibility to *Listeria monocytogenes* infection

Selenium effects on gallium arsenide induced biochemical and immunotoxicological changes in rats

Selenium Rescues Aflatoxin B1-Inhibited T Cell Subsets and Cytokine Levels in Cecal Tonsil of Chickens

Selenium reverses *Pteridium aquilinum*-induced immunotoxic effects

Self-reported allergy, infection, and autoimmune diseases among men and women exposed in utero to diethylstilbestrol

Sensitive biomarkers identification for differentiating Cd and Pb induced toxicity on zebrafish embryos

Sensitive periods for 17 $\beta$ -estradiol exposure during immune system development in sea bass head kidney

Sensitivity of a standard host resistance assay using *Streptococcus agalactiae* for assessing exposure to immunotoxicants in wild cotton rats (*Sigmodon hispidus*)

Sensitivity of selected immunological, hematological, and reproductive parameters in the cotton rat (*Sigmodon hispidus*) to subchronic lead exposure

Sensitivity of the SRBC PFC assay versus ELISA for detection of immunosuppression by TCDD and TCDD-like congeners

Sensitivity of wild cotton rats (*Sigmodon hispidus*) to the immunotoxic effects of low-level arsenic exposure

Sensitivity to suppression of cytotoxic T cell generation by 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD) is dependent on the Ah genotype of the murine host

Sephadex induced bronchial hyperreactivity in the rat: hematology, histology, histochemistry and immunohistology of the lung

Sepsis and treatment-induced immunosuppression in the patient with cancer

Sepsis-Induced T Cell Immunoparalysis: The Ins and Outs of Impaired T Cell Immunity

Sequential respiratory, psychologic, and immunologic studies in relation to methyl isocyanate exposure over two years with model development

Sequestration of 2,3,7,8-tetrachlorodibenzo-p-dioxin by activated carbon eliminates bioavailability and the suppression of immune function in mice

Sera from patients with halothane hepatitis contain antibodies to halothane-induced liver antigens which are not detectable by immunoblotting

Serologic characteristics of ceftriaxone antibodies in 25 patients with drug-induced immune hemolytic anemia



Serologically documented loracarbef (Lorabid)-induced immune thrombocytopenia

Serum Bovine Immunoglobulin for Chemotherapy-Induced Gastrointestinal Mucositis

Serum concentrations of antibodies against vaccine toxoids in children exposed perinatally to immunotoxins

Serum immunoglobulin levels in persons exposed to mercury fumes for various periods of time

Serum immunoglobulins and lymphocyte subset distributions in children and adults living in communities assessed for lead and cadmium exposure

Serum immunoglobulins in chronic exposure to manganese

Serum proteomic profiling associated with immune system impaired by stress using ProteinChip technology

Serum-derived bovine immunoglobulin/protein isolate in the alleviation of chemotherapy-induced mucositis

Severe cytomegalovirus infection in a second kidney transplant recipient treated with ganciclovir, leflunomide, and immunoglobulins, with complications including seizures, acute HCV infection, drug-induced pancytopenia, diabetes, cholangitis, and multi-organ failure with fatal outcome: a case report

Severe Exacerbation of Myasthenia Gravis Associated with Checkpoint Inhibitor Immunotherapy

Severe exacerbation of psoriasis after intravenous immunoglobulin in patient with multiple sclerosis that started during biologic therapy

Severe immune dysfunction after lethal neutron irradiation in a JCO nuclear facility accident victim

Severe minocycline-induced eosinophilic pneumonia: extrapulmonary manifestations and the use of in vitro immunoassays

Severe pulmonary arterial hypertension induced by SU5416 and ovalbumin immunization

Severe stomatitis complicating immune-suppressive switch after cardiac transplantation

Severe stress, depressive symptoms, and changes in lymphocyte subsets in human immunodeficiency virus-infected men. A 2-year follow-up study

Severe thrombotic thrombocytopenic purpura (TTP) induced or exacerbated by the immunostimulatory herb Echinacea

Sex-based differences in lymphocyte proliferation in the spleen after vanadium inhalation

Sex-dimorphic adverse drug reactions to immune suppressive agents in inflammatory bowel disease

Sex-related differences in the immune response of weanling piglets exposed to low doses of fumonisin extract

Sex-specific effects of developmental lead exposure on the immune-neuroendocrine network

Sex-specific effects of neonatal exposures to low levels of cadmium through maternal milk on development and immune functions of juvenile and adult rats

Sex-specific immunomodulatory action of the environmental estrogen 17 $\alpha$ -ethynylestradiol alongside with reproductive impairment in fish

Shape matters: effects of silver nanospheres and wires on human alveolar epithelial cells

Shiga toxin-induced apoptosis is more efficiently inhibited by dimeric recombinant hybrid-IgG/IgA immunoglobulins than by the parental IgG monoclonal antibodies

Short course induction immunosuppression with thymoglobulin for renal transplant recipients

Short-term exposure of Chinook salmon (*Oncorhynchus tshawytscha*) to o,p-DDE or DMSO during early life-history stages causes long-term humoral immunosuppression

Short-term exposure of rodents to diesel exhausts: usefulness for studies of genotoxic and immunotoxic effects

Short-term exposure to JP-8 jet fuel results in long-term immunotoxicity

Short-term immunotherapy: a prospective, randomized, double-blind, placebo-controlled multicenter study of molecular standardized grass and rye allergens in patients with grass pollen-induced allergic rhinitis

SHP-1 is directly activated by the aryl hydrocarbon receptor and regulates BCL-6 in the presence of 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD)

Sialylation of 3-methylcholanthrene-induced fibrosarcoma determines antitumor immune responses during immunoediting

Signaling mechanisms in the restoration of impaired immune function due to diet-induced obesity

Significance of platelet and monocyte activation for therapeutic immunoglobulin-induced thromboembolism

Significant improvement in the outcome of HCV-infected transplant recipients by avoiding rapid steroid tapering and potent induction immunosuppression

Silicon and matrix macromolecules: new research opportunities for old diseases from analysis of potential mechanisms of breast implant toxicity

Silicone breast implants: immunotoxic and epidemiologic issues

Silver and fullerene nanoparticles' effect on interleukin-2-dependent proliferation of CD4 (+) T cells

Silver nanoparticles and dissolved silver activate contrasting immune responses and stress-induced heat shock protein expression in sea urchin

Silymarin attenuates benzo(a)pyrene induced toxicity by mitigating ROS production, DNA damage and calcium mediated apoptosis in peripheral blood mononuclear cells (PBMC)

Silymarin inhibits ultraviolet radiation-induced immune suppression through DNA repair-dependent activation of dendritic cells and stimulation of effector T cells

Silymarin protects PBMC against B(a)P induced toxicity by replenishing redox status and modulating glutathione metabolizing enzymes--an in vitro study

Similar prevalence of platelet factor 4/heparin immunoglobulin G antibodies in patients following cardiac surgery and other patients suspected of heparin-induced thrombocytopenia

Simulated climate change causes immune suppression and protein damage in the crustacean *Nephrops norvegicus*

Simultaneous administration of hesperidin or garlic oil modulates diazinon-induced hemato- and immunotoxicity in rats

Simultaneous geno- and immunotoxicological investigations for early detection of organophosphate toxicity in rats

Simultaneous islet and kidney transplantation in seven patients with type 1 diabetes and end-stage renal disease using a glucocorticoid-free immunosuppressive regimen with alemtuzumab induction

Simultaneous procainamide-induced immune thrombocytopenia and lupus anticoagulant

Single bolus antithymocyte globulin versus basiliximab induction in kidney transplantation with cyclosporine triple immunosuppression: efficacy and safety

Sinus bradycardia induced by darunavir-ritonavir in a patient with acquired immunodeficiency syndrome

Sirolimus addition to tacrolimus-based immunosuppression induces complete remission of post-transplant lymphoproliferative disorder in a liver transplant recipient

Sirolimus as an alternative treatment in patients with granulomatous-lymphocytic lung disease and humoral immunodeficiency with impaired regulatory T cells

Sirolimus in cardiac transplantation: use as a primary immunosuppressant in calcineurin inhibitor-induced nephrotoxicity

Sirolimus induced dyslipidemia in tacrolimus based vs. tacrolimus free immunosuppressive regimens in renal transplant recipients

Sirolimus-based immunosuppression after cardiac transplantation: predictors of recovery from calcineurin inhibitor-induced renal dysfunction

Size distribution effects of cadmium tellurium quantum dots (CdS/CdTe) immunotoxicity on aquatic organisms

Size-dependent effect of zinc oxide on toxicity and inflammatory potential of human monocytes

Sizing Up the Competition: Fc-Mediated Effector Functions of Immune Therapeutics Impaired During Chronic Viral Infection

SK&F 95654-induced acute cardiovascular toxicity in Sprague-Dawley rats--histopathologic, electron microscopic, and immunohistochemical studies

Skin cancer development in mice exposed chronically to immunosuppressive agents

Skin Exposure to Ultraviolet B Rapidly Activates Systemic Neuroendocrine and Immunosuppressive Responses

Skin toxicity of jet fuels: ultrastructural studies and the effects of substance P

Slow release cortisol implants result in impaired innate immune responses and higher infection prevalence following experimental challenge with infectious pancreatic necrosis virus in Atlantic salmon (*Salmo salar*) parr

Smoking habit, immune suppression, oral contraceptive use, and hormone replacement therapy use and cervical carcinogenesis: a review of the literature

Smoking Impairs the Immunomodulatory Capacity of Lung-Resident Mesenchymal Stem Cells in Chronic Obstructive Pulmonary Disease

Smoking-Induced Changes in the Maternal Immune, Endocrine, and Metabolic Pathways and Their Impact on Fetal Growth: A Topical Review

Social isolation during old age worsens cognitive, behavioral and immune impairment

Social stress in male mice impairs long-term antiviral immunity selectively in wounded subjects

Sodium fluoride exposure triggered the formation of neutrophil extracellular traps

Sodium fluoride impairs splenic innate immunity via inactivation of TLR2/MyD88 signaling pathway in mice

Sodium methyldithiocarbamate inhibits MAP kinase activation through toll-like receptor 4, alters cytokine production by mouse peritoneal macrophages, and suppresses innate immunity

Soman intoxication-induced changes in serum acute phase protein levels, corticosterone concentration and immunosuppressive potency of the serum

Somatic and autonomic small fiber neuropathy induced by bortezomib therapy: an immunofluorescence study

Some biochemical and immunological disorders caused by chronic exposure to nitrogen oxides

Some toxicological aspects of polycyclic aromatic hydrocarbons (PAHs) effects

Sources, distribution, and toxicity of polycyclic aromatic hydrocarbons

Species specificity of 2-aryl carbapenem-induced immune-mediated hemolytic anemia in primates

Specific and long-lasting protection from collagen-induced arthritis and oil-induced arthritis in DA rats by administration of immunogens

Specific immune responses in mice following subchronic exposure to acetamiprid

Specific non-coplanar PCB-mediated modulation of bottlenose dolphin and beluga whale phagocytosis upon in vitro exposure

Splenic immunotoxicity in developing cane toads (*Rhinella marina*) from Bermuda

Spuriously low IgG levels in lupus-associated mixed cryoglobulinaemia type II complicated by iatrogenic immunoglobulin-induced immune complex vasculitis

SSSSSSSSSSSS

Stability evaluation of reference genes for real-time PCR in zebrafish (*Danio rerio*) exposed to cadmium chloride and subsequently infected by bacteria *Aeromonas hydrophila*

Standard radiotherapy but not chemotherapy impairs systemic immunity in non-small cell lung cancer

Stannous chloride induces alterations in enzyme activities, lipid peroxidation and histopathology in male rabbit: antioxidant role of vitamin C

Staphylococcal Enterotoxin B (SEB) Induces Memory CD4 T Cell Anergy in vivo and Impairs Recall Immunity to Unrelated Antigens

Staphylococcus aureus proteases degrade lung surfactant protein A potentially impairing innate immunity of the lung

Statin-induced anti-HMGCR antibody-related immune-mediated necrotising myositis achieving complete remission with rituximab

Statin-triggered immune-mediated necrotizing myopathy

Steering carbon nanotubes to scavenger receptor recognition by nanotube surface chemistry modification partially alleviates NFkappaB activation and reduces its immunotoxicity

Stem Cell Transfusion Restores Immune Function in Radiation-Induced Lymphopenic C57BL/6 Mice

Stereoselective induction of developmental toxicity and immunotoxicity by acetochlor in the early life stage of zebrafish

Steroid-exacerbated HIV-associated cutaneous Kaposi's sarcoma immune reconstitution inflammatory syndrome: 'Where a good intention turns bad'

Steroid-Free Immune Suppression Impairs Glycemic Control in a Healthy Cynomolgus Monkey

Steroid-free immunosuppression after kidney transplantation with antithymocyte globulin induction and cyclosporine and mycophenolate mofetil maintenance therapy

Strain differences of cadmium-induced toxicity in rats: Insight from spleen and lung immune responses

Strain specific induction of pyometra and differences in immune responsiveness in mice exposed to 17alpha-ethinyl estradiol or the endocrine disrupting chemical bisphenol A

Strategy for selecting nanotechnology carriers to overcome immunological and hematological toxicities challenging clinical translation of nucleic acid-based therapeutics

Streptococcus pneumoniae disrupts pulmonary immune defence via elastase release following pneumolysin-dependent neutrophil lysis

Stress and demographic decline: a potential effect mediated by impairment of reproduction and immune function in cyclic vole populations

Stress and immune response. On appropriate stress management for prevention and treatment of diseases

Stress and immune responses of nestling tree swallows (*Tachycineta bicolor*) and eastern bluebirds (*Sialia sialis*) exposed to nonpersistent pesticides and p,p'-dichlorodiphenyldichloroethylene in apple orchards of southern Ontario, Canada

Stress and immunity in humans: a meta-analytic review

Stress impairs the efficacy of immune stimulation by CpG-C: Potential neuroendocrine mediating mechanisms and significance to tumor metastasis and the perioperative period

Stress response in Cu<sup>2+</sup> and Cd<sup>2+</sup> exposed oysters (*Crassostrea gigas*): an immunohistochemical approach

Stress response, biotransformation effort, and immunotoxicity in captive birds exposed to inhaled benzene, toluene, nitrogen dioxide, and sulfur dioxide

Stress, age, and immune function: toward a lifespan approach

Stressful life events are associated with low secretion rates of immunoglobulin A in saliva in the middle aged and elderly

Stressful life events of dental students and salivary immunoglobulin A

Striatal but not frontal cortical up-regulation of the epidermal growth factor receptor in rats exposed to immune activation in utero and cannabinoid treatment in adolescence

Stronger association of drug-induced progressive multifocal leukoencephalopathy (PML) with biological immunomodulating agents

Structural and immunohistochemical changes of conjunctiva induced by topical glaucoma medication

Structural equation modeling of immunotoxicity associated with exposure to perfluorinated alkylates

Structural skeletal impairment induced by immunosuppressive therapy in rats: cyclosporine A vs tacrolimus

Structure-activity relationship of phenolic compounds (phenol, pyrocatechol and hydroquinone) on natural lymphocytotoxicity of carp (*Cyprinus carpio*)

Studies of congenic lines in the Brown Norway rat model of Th2-mediated immunopathological disorders show that the aurothiopropanol sulfonate-induced immunological disorder (Aiid3) locus on chromosome 9 plays a major role compared to Aiid2 on chromosome 10

Studies on cell-mediated immunotoxicity of dimethylformamide

Study of biomaterial-induced macrophage activation, cell-mediated immune response and molecular oxidative damage in patients with dermal bioimplants

Study of immune reactivity of minocycline-induced chronic active hepatitis

Study of immunoglobulins, proinflammatory cytokines, lymphoproliferation and phagocytosis in peripheral blood of healthy young people exposed to different levels of atmospheric pollution

Study of immunological suppression induced by combined use of cyclophosphamide and immunopotentiator, with special reference to its adverse effects

Study of immunotoxic properties of formaldehyde

Study of low-intensity 2450-MHz microwave exposure enhancing the genotoxic effects of mitomycin C using micronucleus test and comet assay in vitro.

Study of the inflammatory response of immunocytes to polymerized porcine hemoglobin (pPolyHb

Study of tyrosine hydroxylase immunoreactive neurons in neonate rats lactationally exposed to 2,4-dichlorophenoxyacetic Acid

Study on beta-adrenoceptors mechanism of cadmium immunotoxicity

Study on immunocyte proliferation and DNA damage induced by lead in mice

Study on mouse pulmonary acute injury induced by air-borne PM2.5

Study on reproductive and immune toxicity of male rats exposed to As2O3

Study on the effect of Part III from *Cynomorium songaricum* on immunosuppressive mice induced by cyclophosphamide

Study on the immunotoxicity of ribosome inhibiting protein of *Momordica charantia*

Study on the impact of lead acetate pollutant on immunotoxicity produced by thiamethoxam pesticide

Study on the relationship between cadmium immunotoxicity and corticotropin-releasing factor

Studying the synergistic damage effects induced by 1.8 GHz radiofrequency field radiation (RFR) with four chemical mutagens on human lymphocyte DNA using comet assay in vitro.

Subacute immunotoxicity of the marine phycotoxin yessotoxin in rats

Subacute immunotoxicity study of formaldehyde in male rats

Subacute oral exposure to dibromoacetic acid induced immunotoxicity and apoptosis in the spleen and thymus of the mice

Subchronic 10 day immunotoxicity of polydimethylsiloxane (silicone) fluid, gel and elastomer and polyurethane disks in female B6C3F1 mice

Subchronic administration of technical pentachlorophenol to lactating dairy cattle: immunotoxicologic evaluation

Subchronic arsenism-induced oxidative stress and inflammation contribute to apoptosis through mitochondrial and death receptor dependent pathways in chicken immune organs

Sub-chronic effect of perfluorooctanesulfonate (PFOS) on the balance of type 1 and type 2 cytokine in adult C57BL6 mice

Subchronic effects of perfluorooctanesulfonate exposure on inflammation in adult male C57BL/6 mice

Subchronic exposures to fungal bioaerosols promotes allergic pulmonary inflammation in naive mice

Subchronic immunotoxicity and screening of reproductive toxicity and developmental immunotoxicity following single instillation of HIPCO-single-walled carbon nanotubes: purity-based comparison

Subchronic inhalation toxicity of isobutyl nitrite in BALB/c mice. II. Immunotoxicity studies

Subchronic inhalation toxicity of the chlorinated propane 1,1,1,3,3,3-hexachloropropane (HCC-230fa

Subchronic lead exposure, immunotoxicology and increased disease resistance in Japanese quail (*Coturnix coturnix japonica*)

Subchronic Oral Cadmium Exposure Exerts both Stimulatory and Suppressive Effects on Pulmonary Inflammation/Immune Reactivity in Rats

Subchronic oral toxicity study of furan in Fischer-344 rats

Subchronic toxicity and immunotoxicity of MeO-PEG-poly(D,L-lactic-co-glycolic acid)-PEG-OMe triblock copolymer nanoparticles delivered intravenously into rats

Sub-chronic toxicity of low concentrations of industrial volatile organic pollutants in vitro

Subchronic treatment with mercuric chloride suppresses immune response, elicits behavioral deficits and increases brain serotonin and dopamine metabolism in rats

Subchronic, reproductive, and maternal toxicity studies with tertiary butyl acetate (TBAC

Subclinical doses of T-2 toxin impair acquired immune response and liver cytochrome P450 in pigs

Subclinical GvHD in non-irradiated F1 hybrids: severe lymphoid-tissue GvHD causing prolonged immune dysfunction

Subclinical Reactivation of Cytomegalovirus Drives CD4<sup>+</sup>CD28<sup>null</sup> T-Cell Expansion and Impaired Immune Response to Pneumococcal Vaccination in Antineutrophil Cytoplasmic Antibody-Associated Vasculitis

Sublethal hyperoxia impairs pulmonary innate immunity

Sublingual immunotherapy in children with grass pollen induced allergic rhinoconjunctivitis

Substance P as prophylaxis for JP-8 jet fuel-induced immunotoxicity

Successful combined treatment with rituximab and high dose immunoglobulin in a patient with chronic lymphocytic leukemia with fludarabine-induced severe immune thrombocytopenia



Successful intravenous human immunoglobulin treatment of drug-induced Stevens-Johnson syndrome in a dog

Successful Management by Immunoglobulin for Sunitinib-Induced Thrombocytopenia in a Patient with Advanced Metastatic Renal Cell Carcinoma

Successful management with intravenous immunoglobulins in alemtuzumab-induced acute inflammatory demyelinating neuropathy: clinical report of three patients

Successful treatment by voliconazole for pulmonary and adductor magnus muscle aspergillosis induced by immunosuppressive therapy for hypersensitivity pneumonia

Successful treatment of lymphoma with fludarabine combined with rituximab after immune thrombocytopenia induced by fludarabine

Successful treatment of recalcitrant penicillamine-induced pemphigus foliaceus by low-dose intravenous immunoglobulins

Successful Treatment of Severe Heparin-induced Thrombocytopenia with Intravenous Immunoglobulin, Platelet Transfusion and Rivaroxaban: A Case Report

Successfully treated sulphasalazine-induced fulminant hepatic failure, thrombocytopenia and erythroid hypoplasia with intravenous immunoglobulin

Sudden collapse during haemodialysis due to immune-mediated heparin-induced thrombocytopaenia

Sulfamethoxazole-trimethoprim-induced pneumonitis in a patient with hemophilia B who was infected with the human immunodeficiency virus

Sulfasalazine induced immune thrombocytopenia in a patient with rheumatoid arthritis

Sulfasalazine-induced linear immunoglobulin A bullous dermatosis with DRESS

Sulindac-induced immune hemolytic anemia

Sulphonylurea usage in melioidosis is associated with severe disease and suppressed immune response

Sunlight-induced immunosuppression in humans is initially because of UVB, then UVA, followed by interactive effects

Sunscreen applied at 2 mg cm<sup>-2</sup> during a sunny holiday prevents erythema, a biomarker of ultraviolet radiation-induced DNA damage and suppression of acquired immunity

Sunscreen lotions prevent ultraviolet radiation-induced suppression of antitumor immune responses

Sunset yellow FCF, a permitted food dye, alters functional responses of splenocytes at non-cytotoxic dose

Superoxide dismutase 3 controls adaptive immune responses and contributes to the inhibition of ovalbumin-induced allergic airway inflammation in mice

Supplementing an immunomodulatory feed ingredient to modulate thermoregulation, physiologic, and production responses in lactating dairy cows under heat stress conditions

Suppressed T helper 2 immunity and prolonged survival of a nematode parasite in protein-malnourished mice

Suppression by Delta(9)-tetrahydrocannabinol of the primary immunoglobulin M response by human peripheral blood B cells is associated with impaired STAT3 activation

Suppression of allergen-induced airway inflammation and immune response by the peroxisome proliferator-activated receptor-alpha agonist fenofibrate

Suppression of allergic immune responses to house dust mite (HDM) in rats exposed to 2,3,7,8-TCDD

Suppression of allograft immunity by 3,4,5,3',4',5'-hexachlorobiphenyl. I. Effects of exposure on tumor rejection and cytotoxic T cell activity in vivo

Suppression of allograft immunity by 3,4,5,3',4',5'-hexachlorobiphenyl. II. Effects of exposure on mixed lymphocyte reactivity and induction of suppressor cell activity in vitro

Suppression of B-cell mediated immunity in juvenile chinook salmon (*Oncorhynchus tshawytscha*) after exposure to either a polycyclic aromatic hydrocarbon or to polychlorinated biphenyls

Suppression of cell-mediated immune responses to listeria infection by repeated exposure to diesel exhaust particles in brown Norway rats

Suppression of cell-mediated immunity by a donor-transmitted lymphocytic choriomeningitis virus in a kidney transplant recipient

Suppression of cell-mediated immunity following oral feeding of mice with palmyrah (*Borassus flabellifer* L) flour

Suppression of cellular immune responses in BALB/c mice following oral exposure to permethrin

Suppression of cellular immunity by anesthesia and operation

Suppression of cellular immunity in men with a past history of posttraumatic stress disorder

Suppression of corneal graft rejection in rabbits by a new immunosuppressive agent, FK-506

Suppression of cyclosporine a nephrotoxicity in vivo by transforming growth factor beta receptor-immunoglobulin G chimeric protein

Suppression of human B cell activation by 2,3,7,8-tetrachlorodibenzo-p-dioxin involves altered regulation of B cell lymphoma-6

Suppression of humoral and cell-mediated immune responses by carbon tetrachloride

Suppression of humoral and cell-mediated immune responses in vitro by benzo(a)pyrene

Suppression of humoral immune responses by 2,3,7,8-tetrachlorodibenzo-p-dioxin intercalated in smectite clay

Suppression of humoral immune responses by dialkylnitrosamines: structure-activity relationships

Suppression of humoral immunity by perfluorooctanoic acid is independent of elevated serum corticosterone concentration in mice

Suppression of humoral immunity following exposure to the perfluorinated insecticide sulfluramid

Suppression of humoral immunity in mice following exposure to perfluorooctane sulfonate

Suppression of immune parameters in animal models of morphine dependence

Suppression of immune response associated with functions of B-cells, Th1- and Th2-lymphocytes and interleukins produced by these cells, and pharmacological correction of these disturbances upon acute intoxication with methanol

Suppression of immune response in the B6C3F1 mouse after dietary exposure to the Fusarium mycotoxins deoxynivalenol (vomitoxin) and zearalenone

Suppression of immunoglobulin production by a novel dihydroorotate dehydrogenase inhibitor, S-2678

Suppression of local gut-associated and splenic mitogen responsiveness of lymphoid cells following oral exposure of B6C3F1 mice to 7,12-dimethylbenz[a]anthracene

Suppression of natural killer cells after acute intoxication with alcohols and cholinotropic preparations and their reactivation with T-activin

Suppression of retrovirus-induced immunodeficiency disease (murine AIDS) by trimidox and didox: novel ribonucleotide reductase inhibitors with less bone marrow toxicity than hydroxyurea

Suppression of splenic lymphocyte function by 7,12-dimethylbenz[a]anthracene (DMBA) in vitro

Suppression of Th1- and Th2-type immune responses in infant mouse spleen after prenatal and postnatal exposure to low-level toluene and peptidoglycan

Suppression of the IgM Response by Aryl Hydrocarbon Receptor Activation in Human Primary B Cells Involves Impairment of Immunoglobulin Secretory Processes

Suppression of the immune system by oral glucocorticoid therapy in bronchial asthma

Suppression of the in vitro humoral immune response by chrysotile asbestos

Suppression of the in vitro humoral immune response of mouse splenocytes by 7,12-dimethylbenz[a]anthracene metabolites and inhibition of immunosuppression by alpha-naphthoflavone

Suppression of the in vitro immune response by 7,12-dimethylbenz[a]anthracene in mouse splenocytes co-cultured with rat hepatocytes

Suppression of the NF-kappaB pathway by diesel exhaust particles impairs human antimycobacterial immunity

Suppressive Effect of 1alpha,25-Dihydroxyvitamin D3 on Th17-Immune Responses in Kidney Transplant Recipients With Tacrolimus-Based Immunosuppression

Suppressive effect of O,O-dimethyl, S-ethyl phosphorothioate on immune response

Suppressive effect of ultraviolet B radiation on contact sensitization in mice. II. Systemic immunosuppression is modulated by ultraviolet irradiation and hapten application

Suppressive effects of benzo[a]pyrene upon fish immune function: evolutionarily conserved cellular mechanisms of immunotoxicity

Suppressive effects of thiopental and halothane on specific arms of the immune response

Suppressor T-cell levels are unreliable indicators of the impaired immune response following thermal injury

Surface charges and shell crosslinks each play significant roles in mediating degradation, biofouling, cytotoxicity and immunotoxicity for polyphosphoester-based nanoparticles

Surface coatings determine cytotoxicity and irritation potential of quantum dot nanoparticles in epidermal keratinocytes

Surface expression of the hRSV nucleoprotein impairs immunological synapse formation with T cells

Surface functionalized mesoporous silica nanoparticles with natural proteins for reduced immunotoxicity

Surgery-induced immunomodulation in breast cancer

Surgery-induced immunosuppression and postoperative pain management

Surgical damage to the lymphatic system promotes tumor growth via impaired adaptive immune response

Susceptibility to secondary bacterial infections in growing pigs as an early response in ochratoxicosis

Sustained 3-year efficacy of pre- and coseasonal 5-grass-pollen sublingual immunotherapy tablets in patients with grass pollen-induced rhinoconjunctivitis

Swainsonine exposure induces impairment of host immune response in pregnant BALB/c mice

Swiss mice CD1 fed on mussels contaminated by okadaic acid and yessotoxins: effects on thymus and spleen

Switching to sirolimus-based immune suppression after liver transplantation is safe and effective: a single-center experience

Symptoms and immunologic markers induced by exposure to methyltetrahydrophthalic anhydride

Symptoms of atopy in persons exposed to chronic immunosuppression of polycyclic aromatic hydrocarbons

Synaptophysin immunoreactive axonal swelling in p-bromophenylacetylurea-induced neuropathy

Syndrome of cocaine-levamisole-induced cutaneous vasculitis and immune-mediated leukopenia

Synergism in immunotoxicological effects due to repeated combined administration of arsenic and lead in mice

Synergistic effect of organic solvents and tobacco smoke on serum immunoglobulin levels in humans

Synthesis and characterization of deoxynivalenol glucuronide: its comparative immunotoxicity with deoxynivalenol

Synthesis of zanthoxylamide protoalkaloids and their in silico ADME-Tox screening and in vivo toxicity assessment in zebrafish embryos

Synthetic human chorionic gonadotropin-related oligopeptides impair early innate immune responses to *Listeria monocytogenes* in Mice

Synthetic predator cues impair immune function and make the biological pesticide Bti more lethal for vector mosquitoes

Systemic and immunotoxicity of pristine and PEGylated multi-walled carbon nanotubes in an intravenous 28 days repeated dose toxicity study

Systemic and immunotoxicity of silver nanoparticles in an intravenous 28 days repeated dose toxicity study in rats

Systemic immune cell response in rats after pulmonary exposure to manganese-containing particles collected from welding aerosols

Systemic immune challenges trigger and drive Alzheimer-like neuropathology in mice

Systemic immunosuppression following a single pharyngeal aspiration of 1,2:5,6-dibenzanthracene in female B6C3F1 mice

Systemic immunosuppression induced by photodynamic therapy (PDT) is adoptively transferred by macrophages

Systemic immunotoxicity in AJ mice following 6-month whole body inhalation exposure to diesel exhaust

Systemic immunotoxicity reactions induced by adjuvanted vaccines

Systemic toxicity of di-2-ethylhexyl terephthalate (DEHT) in rodents following four weeks of intravenous exposure

T cell immunoglobulin- and mucin-domain-containing molecule-4 attenuates concanavalin A-induced hepatitis by regulating macrophage

T lymphocyte-proliferative responses of harbor seal (*Phoca vitulina*) peripheral blood mononuclear cells (PBMCs) exposed to pharmaceuticals in vitro

T-2 toxin immunotoxicity on human B and T lymphoid cell lines

T-2 toxin impairment of enteric reovirus clearance in the mouse associated with suppressed immunoglobulin and IFN-gamma responses

T-2 toxin impairs murine immune response to respiratory reovirus and exacerbates viral bronchiolitis

T-2 toxin in the diet suppresses growth and induces immunotoxicity in juvenile Chinese mitten crab (*Eriocheir sinensis*)

T2 Toxin-Induced Changes in Cocaine- and Amphetamine-Regulated Transcript (CART)-Like Immunoreactivity in the Enteric Nervous System Within Selected Fragments of the Porcine Digestive Tract

Tamoxifen persistently disrupts the humoral adaptive immune response of gilthead seabream (*Sparus aurata* L)

Tamoxifen-induced immune-mediated platelet destruction. A case report

Targeted nanoparticles that mimic immune cells in pain control inducing analgesic and anti-inflammatory actions: a potential novel treatment of acute and chronic pain condition

Targeting TLR2 attenuates pulmonary inflammation and fibrosis by reversion of suppressive immune microenvironment

Taurine Administration Mitigates Cisplatin Induced Acute Nephrotoxicity by Decreasing DNA Damage and Inflammation: An Immunocytochemical Study

TCDD adsorbed on silica as a model for TCDD contaminated soils: Evidence for suppression of humoral immunity in mice

TCDD modulation of gut microbiome correlated with liver and immune toxicity in streptozotocin (STZ)-induced hyperglycemic mice

TCDD: an environmental immunotoxicant reveals a novel pathway of immunoregulation--a 30-year odyssey

Tear immunoglobulins in giant papillary conjunctivitis induced by contact lenses

Temporal clinical, proteomic, histological and cellular immune responses of dextran sulfate sodium-induced acute colitis

Temporal profiles of the in vitro phosphorylation rate and immunocontent of glial fibrillary acidic protein (GFAP) after kainic acid-induced lesions in area CA1 of the rat hippocampus: demonstration of a novel phosphoprotein associated with gliosis

Temporal relationships between HIV-1 Tat-induced neuronal degeneration, OX-42 immunoreactivity, reactive astrocytosis, and protein oxidation in the rat striatum

Terahertz radiation increases genomic instability in human lymphocytes.

Teratocarcinomas Arising from Allogeneic Induced Pluripotent Stem Cell-Derived Cardiac Tissue Constructs Provoked Host Immune Rejection in Mice

Teratogenicity and immunotoxicity of 3,3',4,4',5-pentachlorobiphenyl in C57BL/6 mice

Teratogen-induced apoptosis may be affected by immunopotential

Testing of the system of biochemical and immunological indices of the state of population health in the survey of residents of Moscow, exposed to ambient air pollution

Testosterone-Mediated Endocrine Function and TH1/TH2 Cytokine Balance after Prenatal Exposure to Perfluorooctane Sulfonate: By Sex Status

Tetrabromobisphenol A: Disposition, kinetics and toxicity in animals and humans

Tetrabromobisphenol-A induces apoptotic death of auditory cells and hearing loss

Tetrabromoethylcyclohexane (TBECH) exhibits immunotoxicity in murine macrophages

Tetrahydrocannabinol suppresses immune function and enhances HIV replication in the huPBL-SCID mouse

Th17/Treg immunoregulation and implications in treatment of sulfur mustard gas-induced lung diseases

The action spectra for UV-induced suppression of MLR and MECLR show that immunosuppression is mediated by DNA damage

The alteration of immune reactions in inbred BALB/c mice following low-level sarin inhalation exposure

The alteration of ultrastructure and immunoreactivity of human embryonic organ of Corti tissue culture after exposure to aminoglycoside (neomycin) ototoxicity

The antibody response and immunity to challenge infection induced by whole, inactivated and tween-ether split influenza vaccines

The antiproliferative and immunotoxic effects of L-canavanine and L-canaline

The ascidian *Styela plicata* hemocytes as a potential biomarker of marine pollution: In vitro effects of seawater and organic mercury

The blood-brain barrier and immune function and dysfunction

The  $\text{Ca}^{2+}$ -activated  $\text{K}^{+}$  channel of intermediate conductance: a possible target for immune suppression

The capacity of foodstuffs to induce innate immune activation of human monocytes in vitro is dependent on food content of stimulants of Toll-like receptors 2 and 4

The carcinogenic potential of tacrolimus ointment beyond immune suppression: a hypothesis creating case report

The challenges of UV-induced immunomodulation for children's health

The changes of calretinin immunoreactivity in paraquat-induced nephrotoxic rats

The changing spectrum of drug-induced immune hemolytic anemia

The clinical (ir)relevance of opioid-induced immune suppression

The common immunogenic etiology of chronic fatigue syndrome: from infections to vaccines via adjuvants to the ASIA syndrome

The comparative immunotoxicity of five selected compounds following developmental or adult exposure

The comparative immunotoxicity of mesoporous silica nanoparticles and colloidal silica nanoparticles in mice

The consequences of UV-induced immunosuppression for human health

The cortical epithelium of the rat thymus after in vivo exposure to bis(tri-n-butyltin)oxide (TBTO). An (immuno)histological and ultrastructural study

The critical role of autophagy in cadmium-induced immunosuppression regulated by endoplasmic reticulum stress-mediated calpain activation in RAW264.7 mouse monocytes

The current understanding of immunotoxicity induced by silica nanoparticles

The dark and the sunny sides of UVR-induced immunosuppression: photoimmunology revisited

The decreased growth performance and impaired immune function and structural integrity by dietary iron deficiency or excess are associated with TOR, NF-kappaB, p38MAPK, Nrf2 and MLCK signaling in head kidney, spleen and skin of grass carp (*Ctenopharyngodon idella*)

The detoxification effect of vitamin C on zearalenone toxicity in piglets

The effect of aflatoxin on immunity in turkeys. I. Impairment of actively acquired resistance to bacterial challenge

The effect of benzo[alpha]pyrene on expression and signaling cross talk of aryl hydrocarbon receptor and NFATc1 in mouse lung tissue

The effect of brevenal on brevetoxin-induced DNA damage in human lymphocytes

The effect of chronic ammonia exposure on acute-phase proteins, immunoglobulin, and cytokines in laying hens

The effect of curriculum with voluntary activity or mid-term examination on self-esteem, health-promoting lifestyle and immunoglobulin in nursing college student

The effect of electromagnetic radiation with extremely high frequency and low intensity on cytotoxic activity of human natural killer cells.

The effect of ethephon on immune system in male offspring of mice

The effect of experimental epilepsy induced by penicillin administration during pregnancy on nestin expression in the immature rat cerebellum. A light, electron microscopic, and immunohistochemical study

The effect of HIV infection, immunodeficiency, and antiretroviral therapy on the risk of hepatic dysfunction

The effect of immune enhancement and suppression on the development of laparoscopic port site metastases



The effect of lead exposure on tracheal responsiveness to methacholine and ovalbumin, total and differential white blood cells count, and serum levels of immunoglobulin E, histamine, and cytokines in guinea pigs

The effect of Ochratoxin A on antimicrobial polypeptide expression and resistance to water mold infection in channel catfish (*Ictalurus punctatus*)

The effect of ozonization on furniture dust: microbial content and immunotoxicity in vitro

The effect of p-chloroaniline on leucocytes of sheep peripheral blood under the migration-inhibition test conditions

The effect of radiation-induced xerostomia on saliva and serum lysozyme and immunoglobulin levels

The effect of thymogen on the postintoxication immunodeficiency state induced by acute acetonitrile poisoning

The effects of 2100-MHz radiofrequency radiation on nasal mucosa and mucociliary clearance in rats.

The effects of acute exposure to 2,3,7,8-tetrachlorodibenzo-p-dioxin on glioma-specific cytotoxic T-cell activity in Fischer 344 rats

The effects of an in vitro exposure to 17beta-estradiol and nonylphenol on rainbow trout (*Oncorhynchus mykiss*) peripheral blood leukocytes

The effects of benzo[a]pyrene on leucocyte distribution and antibody response in rainbow trout (*Oncorhynchus mykiss*)

The effects of conjugate and light dose on photo-immunotherapy induced cytotoxicity

The effects of early life adversity on the immune system

The effects of estrogenic and androgenic endocrine disruptors on the immune system of fish: a review

The effects of in vitro pesticide exposures on the phagocytic function of four native Australian freshwater fish

The effects of low electromagnetic field and lead acetate combination on some hemato-biochemical and immunotoxicological parameters in mice

The effects of melamine on humoral immunity with or without cyanuric acid in mice

The effects of mycotoxins, fumonisin B1 and aflatoxin B1, on primary swine alveolar macrophages

The effects of ochratoxin A on lipid peroxidation and antioxidant enzymes: a protective role of melatonin

The effects of perinatal tebuconazole exposure on adult neurological, immunological, and reproductive function in rats

The effects of polycyclic aromatic hydrocarbons on the immune system of fish: a review

The effects of tributyltin (TBT) and 3,3',4,4',5-pentachlorobiphenyl (PCB-126) mixtures on antibody responses and phagocyte oxidative burst activity in channel catfish, *Ictalurus punctatus*

The efficacy of intravenous immunoglobulin on lipopolysaccharide-induced fetal brain inflammation in preterm rats

The entomopathogenic fungus *Nomuraea rileyi* impairs cellular immunity of its host *Helicoverpa armigera*

The evaluation of immunoregulatory markers in the course of neurointoxication by mercury over the post-exposure period

The evaluation of potential immunotoxicity induced by environmental pollutant ammonia in broilers

The first reported case of concurrent trimethoprim-sulfamethoxazole-induced immune hemolytic anemia and thrombocytopenia

The flame retardants tetrabromobisphenol A and tetrabromobisphenol A-bisallylether suppress the induction of interleukin-2 receptor alpha chain (CD25) in murine splenocytes

The flavonolignan-silymarin protects enzymatic, hematological, and immune system against gamma-radiation-induced toxicity

The glutathione disulfide mimetic NOV-002 inhibits cyclophosphamide-induced hematopoietic and immune suppression by reducing oxidative stress

The hair dyes PPD and PTD fail to induce a T(H)2 immune response following repeated topical application in BALB/c mice

The histone deacetylase inhibitor, romidepsin, suppresses cellular immune functions of cutaneous T-cell lymphoma patients

The human immunodeficiency virus type 1 Tat protein potentiates zidovudine-induced cellular toxicity in transgenic mice

The immune response of juvenile Atlantic cod (*Gadus morhua* L.) to chronic exposure to produced water

The immune response of peritoneal macrophages due to exposure to inorganic lead in the house mouse *Mus musculus*

The immune response of sheep to subclinical chronic exposure to the herbicide Bentazon TP

The immune response of women with prolonged exposure to electromagnetic fields produced by radiotelevision broadcasting stations.

The immune responses and expression of metallothionein (MT) gene and heat shock protein 70 (HSP 70) in juvenile rockfish, *Sebastes schlegelii*, exposed to waterborne arsenic (As<sub>3</sub>)

The immune status characteristics and kidney function in persons exposed to the effect of ionizing radiation

The immune system in dust-induced bronchitis

The immune system indices of hypertension patients exposed to ionizing radiation

The immune-modulating cytokine and endogenous Alarmin interleukin-33 is upregulated in skin exposed to inflammatory UVB radiation

The Immuno-Enhancement Effects of Tubiechong (*Eupolyphaga sinensis*) Lyophilized Powder in Cyclophosphamide-Induced Immunosuppressed Mice

The immunogenetic basis of collagen induced arthritis in mice: an experimental model for the rational design of immunomodulatory treatments of rheumatoid arthritis

The immunological and chemical detection of N-(hexanoyl)phosphatidylethanolamine and N-(hexanoyl)phosphatidylserine in an oxidative model induced by carbon tetrachloride

The immunology of nickel-induced allergic contact dermatitis

The immunology of the allergy epidemic and the hygiene hypothesis

The immunomodulating effect of microwaves in exposure of the sternum and thymus gland

The immunomodulatory effects of the herbicide propanil on murine macrophage interleukin-6 and tumor necrosis factor- $\alpha$  production

The immunophilin ligand FK506 protects against methamphetamine-induced dopaminergic neurotoxicity in mouse striatum

The immunostatus of T helper and T cytotoxic cells in the patients ten years after exposure to sulfur mustard

The immunosuppressant and hyperplasia-inducing drug cyclosporin A regulates the cell cycle and cyclin B1 gene expression in gingival fibroblasts in vitro

The immunosuppressant FTY720 prolongs survival in a mouse model of diet-induced coronary atherosclerosis and myocardial infarction

The immunosuppressive drug mycophenolate mofetil impairs the adhesion capacity of gastrointestinal tumour cells

The immunotoxic effects of dual exposure to PCP and TCDD

The immunotoxic effects of lead in lead-exposed laboratory animals

The immunotoxicity and neurobehavioral toxicity of zebrafish induced by famoxadone-cymoxanil

The immunotoxicity of aluminum trichloride on rat peritoneal macrophages via beta2-adrenoceptors/cAMP pathway acted by norepinephrine

The immunotoxicity of dibutyl phthalate on the macrophages in mice

The immunotoxicity of graphene oxides and the effect of PVP-coating

The immunotoxicity of three nickel compounds following 13-week inhalation exposure in the mouse

The immunotoxicity of tributyltin oxide (TBTO) does not increase the susceptibility of rats to experimental respiratory infection

The immunotoxicity of triphenyltin chloride in mice

The immunotoxicological basis for the revision of suspended particulates health standard

The immunotoxicological pattern of subchronic and chronic benzene exposure in rats

The Impact of Bisphenol A and Phthalates on Allergy, Asthma, and Immune Function: a Review of Latest Findings

The impact of nanoparticle protein corona on cytotoxicity, immunotoxicity and target drug delivery

The impact on T-regulatory cell related immune responses in rural women exposed to polycyclic aromatic hydrocarbons (PAHs) in household air pollution in Gansu, China: A pilot investigation

The impaired immune function and structural integrity by dietary iron deficiency or excess in gill of fish after infection with *Flavobacterium columnare*: Regulation of NF-kappaB, TOR, JNK, p38MAPK, Nrf2 and MLCK signalling

The impaired immune response to diphtheria vaccination in elderly chronic hemodialysis patients is related to zinc deficiency

The impaired intestinal mucosal immune system by valine deficiency for young grass carp (*Ctenopharyngodon idella*) is associated with decreasing immune status and regulating tight junction proteins transcript abundance in the intestine

The improvement effects of edible bird's nest on proliferation and activation of B lymphocyte and its antagonistic effects on immunosuppression induced by cyclophosphamide

The in vitro effects of mercury on peritoneal leukocytes (PMN and macrophages) from inbred brown Norway and Lewis rats

The increase in striatal neuropeptide Y immunoreactivity induced by neonatal dopamine-depleting lesions in rats is reversed by intrastriatal dopamine-rich transplants

The influence of aminostigmine on the parameters of nonspecific resistance and immune system of the organism upon acute poisoning of benzyl 3-quinuclidylate

The influence of calcium folinate on the immunotoxicity of methanol

The influence of exposure to immunosuppressive treatment during pregnancy on renal function and rate of apoptosis in native kidneys of female Wistar rats

The influence of hydrocarbon composition and exposure conditions on jet fuel-induced immunotoxicity

The influence of low-level sarin inhalation exposure on the host resistance and immune reaction of inbred BALB/c mice after their infection with *Francisella tularensis* LVS

The influence of petrochemicals and stress on the immune system of seabirds

The influence of physical activity on the profile of immune response cells and cytokine synthesis in mice with experimental breast tumors induced by 7,12-dimethylbenzanthracene

The influence of psychological factors on the immune system and immunological diseases

The Influence of Quercetin on Maternal Immunity, Oxidative Stress, and Inflammation in Mice with Exposure of Fine Particulate Matter during Gestation

The influence of single or repeated low-level sarin exposure on immune functions of inbred BALB/c mice

The inhibition of 2,3-dichloro-1-propanol on T cell in vitro and in vivo

The innate and adaptive immune response induced by alveolar macrophages exposed to ambient particulate matter

The Japanese medaka (*Oryzias latipes*) model: applicability for investigating the immunosuppressive effects of the aquatic pollutant benzo[a]pyrene (BaP)

The level of immunoglobulins in the blood serum of guinea pigs with epicutaneous exposure to petroleum refinery products

The long-term oral exposure to titanium dioxide impaired immune functions and triggered cytotoxic and genotoxic impacts in rats

The mechanism of dioxin toxicity: relationship to risk assessment

The mechanism of Mitogen-Activated Protein Kinases to mediate apoptosis and immunotoxicity induced by Benzo[a]pyrene on hemocytes of scallop *Chlamys farreri* in vitro

The mechanisms and consequences of ultraviolet-induced immunosuppression

The mechanisms of surface chemistry effects of mesoporous silicon nanoparticles on immunotoxicity and biocompatibility

The modulatory effect of lipids and glucose on the neonatal immune response induced by *Staphylococcus epidermidis*

The molecular mechanism of cell cycle arrest in the Bursa of Fabricius in chick exposed to Aflatoxin B<sub>1</sub>

The molecular mechanism of G2M cell cycle arrest induced by AFB<sub>1</sub> in the jejunum

The mycotoxin patulin: An updated short review on occurrence, toxicity and analytical challenges

The mycotoxin zearalenone enhances cell proliferation, colony formation and promotes cell migration in the human colon carcinoma cell line HCT116

The neonicotinoid insecticide Clothianidin adversely affects immune signaling in a human cell line

The nuclear factor-kappaB inhibitor pyrrolidine dithiocarbamate reduces polyinosinic-polycytidilic acid-induced immune response in pregnant rats and the behavioral defects of their adult offspring

The occurrence, potential toxicity, and toxicity mechanism of bisphenol S, a substitute of bisphenol A: A critical review of recent progress

The osmolyte taurine protects against ultraviolet B radiation-induced immunosuppression

The pathology induced by highly active antiretroviral therapy against human immunodeficiency virus: an update

The pathophysiology of immune-mediated heparin-induced thrombocytopenia

The PBDE-209 exposure during pregnancy and lactation impairs immune function in rats

The peptide toxin amylosin of *Bacillus amyloliquefaciens* from moisture-damaged buildings is immunotoxic, induces potassium efflux from mammalian cells, and has antimicrobial activity

The physiological performance and immune response of juvenile turbot (*Scophthalmus maximus*) to nitrite exposure

The phytoestrogen genistein suppresses cell-mediated immunity in mice

The possible alleviating effect of saffron on chlorpyrifos experimentally induced cardiotoxicity: Histological, immunohistochemical and biochemical study

The potential immune modulatory effect of chronic bisphenol A exposure on gene regulation in male medaka (*Oryzias latipes*) liver

The potential immunotoxicity of fine particulate matter based on SD rat spleen

The potential of using enzyme-linked immunospot to diagnose cephalosporin-induced maculopapular exanthems

The potential protective role of Akropower against Atrazine- induced humoral immunotoxicity in rabbits

The protective effect of gamma-aminobutyric acid on the development of immune function in chickens under heat stress

The protective effect of hydrated sodium calcium aluminosilicate against haematological, biochemical and pathological changes induced by Zearalenone in mice

The protective effect of N-acetylcysteine on UVB-induced immunosuppression by inhibition of the action of cis-urocanic acid

The protective effect of *Rubia cordifolia* against lead nitrate-induced immune response impairment and kidney oxidative damage

The protective role of the immunomodulator AS101 against chemotherapy-induced alopecia studies on human and animal models

The rat as a model in developmental immunotoxicology

The recombinant 23-kDa N-terminal fragment of bactericidal/permeability-increasing protein (rBPI23) decreases *Escherichia coli*-induced mortality and organ injury during immunosuppression-related neutropenia

The Relationship Between Noncoplanar PCB-Induced Immunotoxicity and Hepatic CYP1A Induction in a Fish Model

The remedial effect of *Thymus vulgaris* extract against lead toxicity-induced oxidative stress, hepatorenal damage, immunosuppression, and hematological disorders in rats

The role of adherent cells in the immunosuppressed state of mouse progeny transplacentally exposed to benzo(alpha)pyrene

The role of altered cutaneous immune responses in the induction and persistence of rosacea

The role of B7 costimulation in benzene immunotoxicity and its potential association with cancer risk

The role of calcitonin gene-related peptide in cutaneous immunosuppression induced by repeated subinflammatory ultraviolet irradiation exposure

The role of CCAAT enhancer-binding protein homologous protein in human immunodeficiency virus protease-inhibitor-induced hepatic lipotoxicity in mice

The role of circulating immune complexes and biocompatibility of staphylococcal protein A immunoadsorption in mitomycin C-induced hemolytic uremic syndrome

The role of cis-urocanic acid in UVB-induced immunosuppression

The role of cytokines in ultraviolet-B induced immunosuppression

The role of cytokines in UV-induced systemic immune suppression

The role of DNA damage and repair in ultraviolet B radiation-induced immunomodulation: relevance for human photocarcinogenesis

The role of epidermal cytokines in the generation of cutaneous immune reactions and ultraviolet radiation-induced immune suppression

The role of fluorescence polarization immuno-assay in the diagnosis of plant-induced cardiac glycoside poisoning livestock in South Africa

The role of IL-4, IL-10, and TNF-alpha in the immune suppression induced by ultraviolet radiation

The role of immunosuppressive drugs in aggravating renal ischemia and reperfusion injury

The role of interleukin family in perfluorooctanoic acid (PFOA)-induced immunotoxicity

The role of leuprolide acetate therapy in triggering auto-immune thyroiditis

The role of metabolism in carbon tetrachloride-mediated immunosuppression. In vitro studies

The role of oxidative stress and innate immunity in O(3) and endotoxin-induced human allergic airway disease

The role of PERK and IRE1 signaling pathways in excessive fluoride mediated impairment of lymphocytes in rats' spleen *in vivo* and *in vitro*

The role of the immune system in hexachlorobenzene-induced toxicity

The role of urocanic acid in UVB-induced suppression of immunity to *Trichinella spiralis* infection in the rat

The role of urocanic acid in UV-induced immunosuppression: recent advances (1992-1994)

The scaffold protein RACK1 is a target of endocrine disrupting chemicals (EDCs) with important implication in immunity

The SCID-hu mouse as a tool in immunotoxicological risk assessment: effects of 2-acetyl-4(5)-tetrahydroxybutyl-imidazole (THI) and di-n-butyltin dichloride (DBTC) on the human thymus in SCID-hu mice

The semiconductor elements arsenic and indium induce apoptosis in rat thymocytes

The severity of trauma determines the immune response to PF4/heparin and the frequency of heparin-induced thrombocytopenia

The significance of epidermal cytokines in UV-induced immune suppression

The skin: target organ in immunotoxicology of small-molecular-weight compounds

The state of health of oncology nurses characterized by genetic and immunotoxicologic biomarkers

The structure-dependent effects of heptachlorodibenzofuran isomers in male C57BL/6 mice: immunotoxicity and monooxygenase enzyme induction

The suitability of different cellular in vitro immunotoxicity and genotoxicity methods for the analysis of nanoparticle-induced events

The suppression of immune activation during enfuvirtide-based salvage therapy is associated with reduced CCR5 expression and decreased concentrations of circulating interleukin-12 and IP-10 during 48 weeks of longitudinal follow-up

The suppression of peritoneal cellular immunity in women with endometriosis could be restored after gonadotropin releasing hormone agonist treatment

The suppressive effects of ultraviolet radiation on immunity in the skin and internal organs: implications for autoimmunity

The teleostean liver as an immunological organ: Intrahepatic immune cells (IHICs) in healthy and benzo[a]pyrene challenged rainbow trout (*Oncorhynchus mykiss*)

The therapy of the heparin-induced thrombosis-thrombocytopenia syndrome with immunoglobulins

The time course and characterization of mercuric chloride-induced immunopathology in the brown Norway rat

The Toxic Effects and Mechanisms of Nano-Cu on the Spleen of Rats

The toxic effects of ammonia exposure on antioxidant and immune responses in Rockfish, *Sebastes schlegelii* during thermal stress

The toxic effects of chlorophenols and associated mechanisms in fish

The toxic effects of multiple persistent organic pollutant exposures on the post-hatch immunity maturation of glaucous gulls



The toxic effects on the stress and immune responses in juvenile rockfish, *Sebastes schlegelii* exposed to hexavalent chromium

The toxic oil syndrome: a perspective on immunotoxicological mechanisms

The toxicity of acute exposure to T-2 toxin evaluated by the metabonomics technique

The toxicity of chlorpyrifos on the early life stage of zebrafish: a survey on the endpoints at development, locomotor behavior, oxidative stress and immunotoxicity

The toxicity of deglycosylated ricin A chain-containing immunotoxins in patients with non-Hodgkin's lymphoma is exacerbated by prior radiotherapy: a retrospective analysis of patients in five clinical trials

The toxicity of silica nanoparticles to the immune system

The toxicity of the N-hydroxy and 6-hydroxy metabolites of 3,4-dichloropropionanilide does not depend on calcium release-activated calcium channel inhibition

The toxicity potential of pharmaceuticals found in the Douro River estuary (Portugal): evaluation of impacts on fish liver, by histopathology, stereology, vitellogenin and CYP1A immunohistochemistry, after sub-acute exposures of the zebrafish model

The toxicological impacts of the *Fusarium* mycotoxin, deoxynivalenol, in poultry flocks with special reference to immunotoxicity

The toxicology of mercury: Current research and emerging trends

The transferability from rat subacute 4-week oral toxicity study to translational research exemplified by two pharmaceutical immunosuppressants and two environmental pollutants with immunomodulating properties

The unique immunological features of heparin-induced thrombocytopenia

The upregulation of pro-inflammatory cytokines in the rabbit uterus under the lipopolysaccharide-induced reversible immunoresponse state

The uptake kinetics and immunotoxic effects of microcystin-LR in human and chicken peripheral blood lymphocytes in vitro

The urokinase receptor can be induced by *Borrelia burgdorferi* through receptors of the innate immune system

The use of FK506 in simultaneous pancreas/kidney transplantation: rescue, induction, and maintenance immunosuppression

The use of tacrolimus as induction and maintenance immunosuppression in renal cadaveric transplant recipients over the age of 60

Thearubigins protect against acetaminophen-induced hepatic and renal injury in mice: biochemical, histopathological, immunohistochemical, and flow cytometry study

T-helper type 1-T-helper type 2 shift and nasal remodeling after fine particulate matter exposure in a rat model of allergic rhinitis

Therapeutic and immune effects of 5-aminolevulinic acid photodynamic therapy on UVB-induced squamous cell carcinomas in hairless mice

Therapeutic effects of antibodies to tumor necrosis factor- $\alpha$ , interleukin-6 and cytotoxic T-lymphocyte antigen 4 immunoglobulin in mice with glucose-6-phosphate isomerase induced arthritis

Therapy of heparin-induced thrombosis-thrombocytopenia syndrome using immunoglobulins

Therapy options for chronic lung allograft dysfunction-bronchiolitis obliterans syndrome following first-line immunosuppressive strategies: A systematic review

Thiamethoxam-mediated alteration in multi-biomarkers of a model organism, *Galleria mellonella* L. (Lepidoptera: Pyralidae)

Thiamin deficiency induces impaired fish gill immune responses, tight junction protein expression and antioxidant capacity: Roles of the NF- $\kappa$ B, TOR, p38 MAPK and Nrf2 signaling molecules

Thiol compounds inhibit mercury-induced immunological and immunopathological alterations in susceptible mice

Thiopronine-induced herpetiform pemphigus: report of a case studied by immunoelectron microscopy and immunoblot analysis

Thrombotic micro-angiopathy with sirolimus-based immunosuppression: potentiation of calcineurin-inhibitor-induced endothelial damage

Thymocyte apoptosis as a mechanism for tributyltin-induced thymic atrophy in vivo

Thymoglobulin induction in liver transplant recipients with a tacrolimus, mycophenolate mofetil, and steroid immunosuppressive regimen: a five-year randomized prospective study

Thymoquinone ameliorates the immunological and histological changes induced by exposure to imidacloprid insecticide

Thymus atrophy and changes in thymocyte subpopulations of BN rats with mercury-induced renal autoimmune disease

Thymus atrophy and immunosuppression induced by organotin compounds

Thymus-directed immunotoxicity of airborne dust particles from Upper Silesia (Poland) under acute extrapulmonary studies in mice

Thyroid Dysfunction, Neurological Disorder and Immunosuppression as the Consequences of Long-term Combined Stress

Thyroid hormone suppression and cell-mediated immunomodulation in American kestrels (*Falco sparverius*) exposed to PCBs

Thyroxine modulation of immune toxicity induced by mixture pesticides mancozeb and fipronil in mice

Ticlopidine-induced aplastic anemia: development of chromosomal abnormalities and response to immunosuppressive therapy

Tier-2 studies on monocrotaline immunotoxicity in C57BL/6 mice

Tilapia (*Oreochromis niloticus*) and rodents exhibit similar patterns of inhibited antibody production following exposure to immunotoxic chemicals

Time course of viremia and antibody seroconversion following human immunodeficiency virus exposure

Time-course study of the immunotoxic effects of the anticancer drug chlorambucil in the rat

Time-dependent inhibition of immune complex-induced lung injury by catalase: relationship to alterations in macrophage and neutrophil matrix metalloproteinase elaboration

Time-response characteristic and potential biomarker identification of heavy metal induced toxicity in zebrafish

Tissue distribution of <sup>14</sup>C-labelled perfluorooctanoic acid in adult mice after 1-5 days of dietary exposure to an experimental dose or a lower dose that resulted in blood levels similar to those detected in exposed humans

TNF Drives Monocyte Dysfunction with Age and Results in Impaired Anti-pneumococcal Immunity

TNF-mediated toxicity after massive induction of specific CD8+ T cells following immunization of mice with a tumor-specific peptide

Tobacco smoke exposure and multiplexed immunoglobulin E sensitization in children: a population-based study

Tobacco smoking increases immune activation and impairs T-cell function in HIV infected patients on antiretrovirals: a cross-sectional pilot study

Tolerability of cyclophosphamide and methotrexate induction immunosuppression in nonhuman primates

Tolerance to *Bacillus thuringiensis* endotoxin in immune-suppressed larvae of the flour moth *Ephestia kuehniella*

Toll-like receptor 4 mediates chronic restraint stress-induced immune suppression

Toll-like receptor 9 is required for chronic stress-induced immune suppression

Topical and systemic immunoreaction triggered by intravesical chemotherapy in an N-butyl-N-(4-hydroxybutyl) nitrosamine induced bladder cancer mouse model

Topical application of a novel immunomodulatory peptide, RDP58, reduces skin inflammation in the phorbol ester-induced dermatitis model

Topical application of the anti-microbial chemical triclosan induces immunomodulatory responses through the S100A8/A9-TLR4 pathway

Topical cathelicidin (LL-37) an innate immune peptide induces acute olfactory epithelium inflammation in a mouse model

Topical cidofovir-induced acute kidney injury in two severely immunocompromised patients with refractory multidrug-resistant herpes simplex virus infections

Topical imiquimod treatment prevents UV-light induced loss of contact hypersensitivity and immune tolerance

Topical nicotinamide modulates cellular energy metabolism and provides broad-spectrum protection against ultraviolet radiation-induced immunosuppression in humans

Topical riboflavin attenuates ultraviolet B- and ultraviolet A-induced immunosuppression in humans

Topical vitamin E inhibition of immunosuppression and tumorigenesis induced by ultraviolet irradiation

Topically applied eicosapentaenoic acid protects against local immunosuppression induced by UVB irradiation, cis-urocanic acid and thymidine dinucleotides

Torque Teno Virus Is Associated With the State of Immune Suppression Early After Liver Transplantation

Total and differential white blood cell counts in *Caiman latirostris* after in ovo and in vivo exposure to insecticides

Toward establishment of temperature thresholds for immunological impact of heat exposure in humans

Towards a nanospecific approach for risk assessment

Toxic and immuno-allergic nephropathies induced by antibiotics

Toxic effects of benzene and toluene

Toxic effects of chemical warfare agent mixtures on the mussel *Mytilus trossulus* in the Baltic Sea: A laboratory exposure study

Toxic effects of chlorpyrifos on lysozyme activities, the contents of complement C3 and IgM, and IgM and complement C3 expressions in common carp (*Cyprinus carpio* L)

Toxic effects of D-galactose on thymus and spleen that resemble aging

Toxic effects of dietary methylmercury on immune function and hematology in American kestrels (*Falco sparverius*)

Toxic effects of lead exposure on bioaccumulation, oxidative stress, neurotoxicity, and immune responses in fish: A review

Toxic effects of maternal zearalenone exposure on intestinal oxidative stress, barrier function, immunological and morphological changes in rats

Toxic effects of methyl methanesulfonate (MMS) on activated macrophages from chickens

Toxic effects of paraquat on cytokine expression in common carp, *Cyprinus carpio* L

Toxic effects of sarin in rats at three months following single or repeated low-level inhalation exposure

Toxic effects of various pollutants in 11B7501 lymphoma B cell line from harbour seal (*Phoca vitulina*)

Toxic effects of waterborne nitrite exposure on antioxidant responses, acetylcholinesterase inhibition, and immune responses in olive flounders, *Paralichthys olivaceus*, reared in bio-floc and seawater

Toxicity data relevant for hazard characterization

Toxicity identification evaluations for the investigation of fish kills: a case study

Toxicity of atrazine and nonylphenol in juvenile rainbow trout (*Oncorhynchus mykiss*): effects on general health, disease susceptibility and gene expression

Toxicity of bis(tri-n-butyltin)oxide in the rat. II. Suppression of thymus-dependent immune responses and of parameters of nonspecific resistance after short-term exposure

Toxicity of carbon-based nanomaterials: Reviewing recent reports in medical and biological systems

Toxicity of copper oxide and basic copper carbonate nanoparticles after short-term oral exposure in rats

Toxicity of dermally applied alpha-cypermethrin in rats

Toxicity of domoic acid in the marine mussel *Mytilus edulis*

Toxicity of environmentally-relevant concentrations of arsenic on developing T lymphocyte

Toxicity of inhaled isobutyl nitrite in BALB/c mice: systemic and immunotoxic studies

Toxicity of methyl tertiary-butyl ether on human blood lymphocytes

Toxicity of Nanoparticles and an Overview of Current Experimental Models

Toxicity of organotin compounds: shared and unshared biochemical targets and mechanisms in animal cells

Toxicity of polybrominated biphenyls (PBBs) in Domestic and laboratory animals

Toxicity of the antimalarial artemisinin and its derivatives

Toxicity of the immune suppressant cyclosporin A in the rat

Toxicity of tributyltin in the marine mollusc *Mytilus edulis*

Toxicity of triorganotin compounds: comparative in vivo studies with a series of trialkyltin compounds and triphenyltin chloride in male rats

Toxicity profile of the GATA-3-specific DNase hgd40 after inhalation exposure

Toxicity, histopathological alterations and immunohistochemical CYP1A induction in the early life stages of the seabream, *Sparus aurata*, following waterborne exposure to B(a)P and TCDD

Toxicogenetic evaluation of dichlorophene in peripheral blood and in the cells of the immune system using molecular and flow cytometric approaches

Toxicogenomic analysis of immune system-related genes in Japanese flounder (*Paralichthys olivaceus*) exposed to heavy oil

Toxicogenomic profiles in relation to maternal immunotoxic exposure and immune functionality in newborns

Toxicokinetics and toxicodynamics of ochratoxin A, an update

Toxicologic and immunologic effects of perinatal exposure to the brominated diphenyl ether (BDE) mixture DE-71 in the Sprague-Dawley rat

Toxicologic, pathologic, and immunotoxic effects of 2,4-dichlorophenol in rats

Toxicological and immune findings in workers exposed to pentachlorophenol (PCP)

Toxicological assessments of rats exposed prenatally to inhaled vapors of gasoline and gasoline-ethanol blends

Toxicological characteristics of Ochratoxin A and its impact on male reproduction

Toxicological consequences of Aroclor 1254 ingestion by female rhesus (*Macaca mulatta*) monkeys. Part 1B. Prebreeding phase: clinical and analytical laboratory findings

Toxicological effects of nickel chloride on IgA+ B Cells and sIgA, IgA, IgG, IgM in the intestinal mucosal immunity in broilers

Toxicological effects of trichloroethylene exposure on immune disorders

Toxicological effects of waterborne Zn on the proximal and distal intestines of large yellow croaker *Larimichthys crocea*

Toxicological impact of inhaled electric mosquito-repellent liquid on the rat: a hematological, cytokine indications, oxidative stress and tumor markers

Toxicological in vitro effects of heavy metals on gilthead seabream (*Sparus aurata* L.) head-kidney leucocytes

Toxicological safety evaluation of gadobutrol

Toxicology and immunotoxicology of mercury: a comparative review in fish and humans

Toxicology of deoxynivalenol (vomitoxin)

Toxicopathological and immunological studies on different concentrations of chitosan-coated silver nanoparticles in rats

Traditional preparation of *Phaleria nisidai*, a Palauan tea, reduces exposure to toxic daphnane-type diterpene esters while maintaining immunomodulatory activity

Trans-10, cis-12 conjugated linoleic acid and the PPAR-gamma agonist rosiglitazone attenuate lipopolysaccharide-induced TNF-alpha production by bovine immune cells

Transcriptional networks associated with the immune system are disrupted by organochlorine pesticides in largemouth bass (*Micropterus salmoides*) ovary

Transcriptional Profiling of Dibenzo[def,p]chrysene-induced Spleen Atrophy Provides Mechanistic Insights into its Immunotoxicity in MutaMouse

Transcriptome analysis of different sizes of 3-mercaptopropionic acid-modified cadmium telluride quantum dot-induced toxic effects reveals immune response in rat hippocampus

Transcriptome analysis provides insights into the immunity function of venom glands in *Pardosa pseudoannulata* in responses to cadmium toxicity

Transcriptome Profiling Reveals Disruption of Innate Immunity in Chronic Heavy Ethanol Consuming Female Rhesus Macaques

Transcriptomic analyses of human bronchial epithelial cells BEAS-2B exposed to brominated flame retardant (tetrabromobisphenol A

Transcriptomic and proteomic profiling reveals the intestinal immunotoxicity induced by aflatoxin M1 and ochratoxin A

Transcriptomic profile indicative of immunotoxic exposure: in vitro studies in peripheral blood mononuclear cells

Transcriptomic study of the toxic mechanism triggered by beauvericin in Jurkat cells

Transfusion medicine illustrated. Profound piperacillin-mediated drug-induced immune hemolysis in a patient with cystic fibrosis

Transgenerational disrupting impacts of atrazine in zebrafish: Beneficial effects of dietary spirulina

Transient immune impairment after a simulated long-haul flight

Translational immunotoxicology of immunomodulatory monoclonal antibodies

Transport of immunoglobulins in the gingival epithelium. I. In diphenylhydantoin-induced hyperplasia

Treatment of drug-induced toxic epidermal necrolysis (Lyell's syndrome) with intravenous human immunoglobulins

Treatment of gold-induced aplastic anaemia with immunosuppressive therapy

Treatment of pemphigus vulgaris with high-dose intravenous immunoglobulins in a patient with steroid-induced osteonecrosis of the femoral head

Treatment of silymarin, a plant flavonoid, prevents ultraviolet light-induced immune suppression and oxidative stress in mouse skin

Treatment of virus-induced myocardial injury with a novel immunomodulating agent, vesnarinone. Suppression of natural killer cell activity and tumor necrosis factor-alpha production

Trehalose suppresses cadmium-activated Nrf2 signaling pathway to protect against spleen injury

Tremor induced by Calcineurin inhibitor immunosuppression: a single-centre observational study in kidney transplanted patients

Tributyltin chloride-induced immunotoxicity and thymocyte apoptosis are related to abnormal Fas expression

Tributyltin induces mitochondrial fission through Mfn1 degradation in human induced pluripotent stem cells

Tributyltin potentiates 3,3',4,4',5-pentachlorobiphenyl-induced cytochrome P-4501A-related activity

Tributyltin-sulphydryl interaction as a cause of immunotoxicity in phagocytes of tunicates

Trichloroethylene Exposure Reduces Liver Injury in a Mouse Model of Primary Biliary Cholangitis

Trichosanthin treatment of HIV-induced immune dysregulation

Trimethoprim-induced immune hemolytic anemia in a pediatric oncology patient presenting as an acute hemolytic transfusion reaction

Trimethoprim-sulfamethoxazole-induced cholestatic hepatitis. Clinico-immunological demonstration of its allergic origin

Trimethyltin-induced neurotoxicity: gene expression pathway analysis, q-RT-PCR and immunoblotting reveal early effects associated with hippocampal damage and gliosis

Triorganotin-induced cytotoxicity to rat thymus, bone marrow and red blood cells as determined by several in vitro assays

Triphenyltin acetate-induced cytotoxicity and CD4(+) and CD8(+) depletion in mouse thymocyte primary cultures

Triple drug immunosuppression significantly reduces immune activation and allograft arteriosclerosis in cytomegalovirus-infected rat aortic allografts and induces early latency of viral infection

Triptolide protects against 1-methyl-4-phenyl pyridinium-induced dopaminergic neurotoxicity in rats: implication for immunosuppressive therapy in Parkinson's disease

Trivalent arsenic species induce changes in expression and levels of proinflammatory cytokines in intestinal epithelial cells

Trophic transfer and in vivo immunotoxicological effects of tributyltin (TBT) in polar seastar *Leptasterias polaris*

Trypanosoma cruzi extracts elicit protective immune response against chemically induced colon and mammary cancers

Tryptophan-induced lung disease: an immunophenotypic, immunofluorescent, and electron microscopic study

Tungsten-induced denaturation and aggregation of epoetin alfa during primary packaging as a cause of immunogenicity

Tunisian radish (*Raphanus sativus*) extract prevents cadmium-induced immunotoxic and biochemical alterations in rats



## Two Cases of Sinusitis Induced by Immune Checkpoint Inhibition

Two immunotoxicity ring studies according to OECD TG 407-comparison of data on cyclosporin A and hexachlorobenzene

Two patients with heparin-induced thrombocytopenia followed by idiopathic (immune) thrombocytopenic purpura: case report

Two types of nomifensine-induced immune haemolytic anaemias: drug-dependent sensitization and/or autoimmunization

Type 1 and Type 2 cytokines imbalance in adult male C57BL/6 mice following a 7-day oral exposure to perfluorooctanesulfonate (PFOS)

Type I IFN signaling in CD8<sup>+</sup> DCs impairs Th1-dependent malaria immunity

Tyrosine kinase inhibitors impair B-cell immune responses in CML through off-target inhibition of kinases important for cell signaling

Ultrafine particulate matter impairs mitochondrial redox homeostasis and activates phosphatidylinositol 3-kinase mediated DNA damage responses in lymphocytes

Ultrastructural and immunocytochemical studies of macrophages in an excitotoxin induced lesion in the rat brain

Ultrastructural immunohistochemical study of L-type amino acid transporter 1-4F2 heavy chain in tumor microvasculatures of N-butyl-N-(4-hydroxybutyl) nitrosamine (BBN) induced rat bladder carcinoma

Ultraviolet A radiation (320-400 nm) protects hairless mice from immunosuppression induced by ultraviolet B radiation (280-320 nm) or cis-urocanic acid

Ultraviolet B (UVB)-induced immunosuppression: biologic, cellular, and molecular effects

Ultraviolet radiation-induced immune modulation: potential consequences for infectious, allergic, and autoimmune disease

Ultraviolet Radiation-Induced Immunosuppression: Induction of Regulatory T Cells

Ultraviolet spectral energy differences affect the ability of sunscreen lotions to prevent ultraviolet-radiation-induced immunosuppression

Ultraviolet-A (UVA-1) radiation suppresses immunoglobulin production of activated B lymphocytes in vitro

Ultraviolet-induced immunosuppression: implications for photocarcinogenesis

Umbilical cord mesenchymal stem cells modulate dextran sulfate sodium induced acute colitis in immunodeficient mice

Under ambient UVA exposure, pefloxacin exhibits both immunomodulatory and genotoxic effects via multiple mechanisms

Understanding the cytotoxicity or cytoprotective effects of biological and synthetic quinone derivatives by redox mechanism

Unique pulmonary immunotoxicological effects of urban PM are not recapitulated solely by carbon black, diesel exhaust or coal fly ash

Unravelling the mechanisms of PFOS toxicity by combining morphological and transcriptomic analyses in zebrafish embryos

Unusual case report of thrombotic microangiopathy of the small bowel following liver transplantation, a possible immunosuppressant-induced disease with histological and ultrastructural findings

Unusual serological findings associated with ceftriaxone-induced immune hemolytic anemia in a child with disseminated low-grade glioma

Uptake, distribution and immunotoxicological effects of mercury in mice

Use of a suite of biomarkers to assess the effects of carbamazepine, bisphenol A, atrazine, and their mixtures on green mussels, *Perna viridis*

Use of acetylcysteine and polyoxydonium for the correction of immune homeostasis and antioxidant system impaired by combined action of 1,2-dichloroethane and heavy mechanical trauma

Use of basiliximab induction in low-immunological risk renal transplant recipients receiving tacrolimus-based immunosuppression

Use of bed rest and head-down tilt to simulate spaceflight-induce immune system changes

Use of genetic toxicology data in U.S. EPA risk assessment: the mercury study report as an example

Use of intravenous gammaglobulin as an immune replacement and an immune suppressant

Use of intravenous immunoglobulin G to treat spontaneous heparin-induced thrombocytopenia

Use of intravenous immunoglobulin to treat chronic bilateral otomastoiditis in the setting of rituximab induced hypogammaglobulinemia

Use of Mishell-Dutton culture for the detection of the immunosuppressive effect of iron-containing compounds

Use of *Trichosporon mycotoxinivorans* to suppress the effects of ochratoxicosis on the immune system of broiler chicks

Usefulness of immunologic and protein determinations for assessment of exposure to mercury

Usefulness of specific immunotherapy in patients with severe perennial allergic rhinitis induced by house dust mite: a double-blind, randomized, placebo-controlled trial

Using cell apoptosis, micronuclei and immune alternations as biomarkers of phenanthrene exposure in yellowfin seabream (*Acanthopagrus latus*)

Using immunotoxicity information to improve cancer risk assessment for polycyclic aromatic hydrocarbon mixtures

Utility of immune monitoring in heart transplant recipients on everolimus-based immune suppression

UV irradiation of immunized mice induces type 1 regulatory T cells that suppress tumor antigen specific cytotoxic T lymphocyte responses

UV radiation-induced immunosuppression and skin cancer

UV radiation-induced immunosuppression is greater in men and prevented by topical nicotinamide

UVA exposure affects UVB and cis-urocanic acid-induced systemic suppression of immune responses in *Listeria monocytogenes*-infected Balb/c mice

UVB exposure impairs immune responses after hepatitis B vaccination in two different mouse strains

UV-induced immune suppression and sunscreen

UV-induced immunosuppression and skin cancers

UV-induced immunosuppression and the efficacy of vaccination

UV-induced immunosuppression in the balance

UV-induced production of immunosuppressive mediators in human skin: prevention by a broadspectrum sunscreen

Vaccine-elicited CD4 T cells induce immunopathology after chronic LCMV infection

Vaccine-induced autoimmunity: the role of molecular mimicry and immune crossreaction

Vacuolar myelopathy and vacuolar cerebellar leukoencephalopathy: a late complication of AIDS after highly active antiretroviral therapy-induced immune reconstitution

Validation of immune function testing during a 4-week oral toxicity study with FK506

Validation of immunotoxicology techniques in passerine chicks exposed to Oil Sands tailings water

Validation of the *Candida albicans* delayed-type hypersensitivity (DTH) model in the female BCF mouse for use in immunotoxicological investigations

Value of antibody level in diagnosing anhydride-induced immunologic respiratory disease

Value of phagocyte function screening for immunotoxicity of nanoparticles in vivo

Vanadium (V)-induced neurotoxicity in the rat central nervous system: a histo-immunohistochemical study

Vanadium and tungsten derivatives as antidiabetic agents: a review of their toxic effects

Vanadium carcinogenic, immunotoxic and neurotoxic effects: a review of in vitro studies

Vanadium pentoxide inhalation provokes germinal center hyperplasia and suppressed humoral immune responses

Vanadium toxicity in the thymic development

Vancomycin-induced Immune Thrombocytopenia Proven by the Detection of Vancomycin-dependent Anti-platelet Antibody with Flow Cytometry

Varicella exposure in a child at risk of being immunosuppressed

Varicella zoster reactivation after hematopoietic stem cell transplant in children is strongly correlated with leukemia treatment and suppression of host T-lymphocyte immunity

Varied immuno-related adverse events induced by immune-check point inhibitors - Nivolumab-associated psoriasiform dermatitis related with increased serum level of interleukin-6

Venom components of *Asobara japonica* impair cellular immune responses of host *Drosophila melanogaster*

Venom of parasitoid, *Pteromalus puparum*, suppresses host, *Pieris rapae*, immune promotion by decreasing host C-type lectin gene expression

Vigabatrin-induced modification of Ki-67 expression in gingival epithelium: immunohistochemical study of a short series

Vitamin A deficiency impairs the immune response to intranasal vaccination and RSV infection in neonatal calves

Vitamin A deficiency impairs vaccine-elicited gastrointestinal immunity

Vitamin C Modulates the Immunotoxic Effect of 17alpha-Methyltestosterone in Nile Tilapia

Vitamin E pretreatment prevents the immunotoxicity of dithiocarbamate pesticide mancozeb in vitro: A comparative age-related assessment in mice and chick

Vomitoxin-induced cyclooxygenase-2 gene expression in macrophages mediated by activation of ERK and p38 but not JNK mitogen-activated protein kinases

Vulnerability, distress, and immune response to vaccination in older adults

Walnut Polyphenol Extract Attenuates Immunotoxicity Induced by 4-Pentylphenol and 3-methyl-4-nitrophenol in Murine Splenic Lymphocyte

Walnut Polyphenol Extract Protects against Fenitrothion-Induced Immunotoxicity in Murine Splenic Lymphocytes

Walnut Polyphenol Extract Protects against Malathion- and Chlorpyrifos-Induced Immunotoxicity by Modulating TLRx-NOX-ROS

Water Contaminants Associated With Unconventional Oil and Gas Extraction Cause Immunotoxicity to Amphibian Tadpoles

Waterborne microcystin-LR exposure induced chronic inflammatory response via MyD88-dependent toll-like receptor signaling pathway in male zebrafish

Water-soluble polysaccharides from *Grifola Frondosa* fruiting bodies protect against immunosuppression in cyclophosphamide-induced mice via JAK2/STAT3/SOCS signal transduction pathways

Wavelength dependency for UVA-induced suppression of recall immunity in humans

Well-being and immune response: a multi-system perspective

Western-style diets induce oxidative stress and dysregulate immune responses in the colon in a mouse model of sporadic colon cancer

What is the main source of human exposure to higher alcohols and is there a link to immunotoxicity

Whole blood TNF- $\alpha$  production as a sensitive measure for immunotoxicity of anticancer drugs

Whole-body inhalation exposure to 1-bromopropane suppresses the IgM response to sheep red blood cells in female B6C3F1 mice and Fisher 344/N rats

Wild *Panax ginseng* (*Panax ginseng* C.A. Meyer) protects against methotrexate-induced cell regression by enhancing the immune response in RAW 264.7 macrophages

Wogonin prevents immunosuppressive action but not anti-inflammatory effect induced by glucocorticoid

Work stress and innate immune response

Workshop on perinatal exposure to dioxin-like compounds. V. Immunologic effects

Yulangsan polysaccharide improves redox homeostasis and immune impairment in D-galactose-induced mimetic aging

Yupingfeng Powder relieves the immune suppression induced by dexamethasone in mice

Zearalenone induces immunotoxicity in mice: possible protective effects of radish extract (*Raphanus sativus*)

Zearalenone inhibits T cell chemotaxis by inhibiting cell adhesion and migration related proteins

Zearalenone, an estrogenic mycotoxin, is an immunotoxic compound

Zebrafish: A complete animal model to enumerate the nanoparticle toxicity

Zidovudine as an immunomodulatory agent

Zidovudine impairs immunological recovery on first-line antiretroviral therapy: collaborative analysis of cohort studies in southern Africa

Zidovudine-induced fatal lactic acidosis and hepatic failure in patients with acquired immunodeficiency syndrome: report of two patients and review of the literature

Zinc- and tin-induced apoptotic mechanisms in immune system and cranial nerve system

Zinc deficiency impairs immune responses against parasitic nematode infections at intestinal and systemic sites

Zinc oxide nanoparticle induced age dependent immunotoxicity in BALB/c mice

Zinc protects HepG2 cells against the oxidative damage and DNA damage induced by ochratoxin A

Ziram induces apoptosis and necrosis in human immune cells

ZnO nanoparticles induced inflammatory response and genotoxicity in human blood cells: A mechanistic approach

#### A4-C. Bibliography – Low-Tech Factors that Strengthen Immune System

In the main text, two types of strategic treatments for a weakened immune system were defined: vaccine-based and non-vaccine-based. The vaccine-based treatments were addressed in Appendix 3. The non-vaccine-based treatments were divided into two categories: identifying and eliminating factors that contribute to weakening the immune system, and adding factors that contribute to strengthening a weak immune system. Many factors of the first category were presented in Table A4-1. In the present section, some factors of the second category (especially low-tech factors that strengthen the immune system) will be presented in the form of a brief bibliography. Many more factors of this nature exist, and could be identified with a much larger study. The bibliography follows.

Anon: Eat more fruits and vegetables to improve your immune system. Five or more portions of fruits and vegetables daily significantly increase antibody response. *DukeMedicine healthnews*. 2013;19(2):5-6.

Asyary A, Veruswati M. Sunlight exposure increased Covid-19 recovery rates: A study in the central pandemic area of Indonesia. *The Science of the total environment*. 2020;729:139016.

Bogden JD, Louria DB. Micronutrients and Immunity in Older People. Bendich A, Deckelbaum RJ, editors 2009. 545-65 p.

Briguglio M, Pregliasco FE, Lombardi G, Perazzo P and Banfi G: The Malnutritional Status of the Host as a Virulence Factor for New Coronavirus SARS-CoV-2. *Frontiers in medicine* 7: 146, 2020.

Briguglio M, Pregliasco FE, Lombardi G, Perazzo P, Banfi G. The Malnutritional Status of the Host as a Virulence Factor for New Coronavirus SARS-CoV-2. *Frontiers in Medicine*. 2020;7.

Calder PC, Carr AC, Gombart AF, Eggersdorfer M. Optimal Nutritional Status for a Well-Functioning Immune System Is an Important Factor to Protect against Viral Infections. *Nutrients*. 2020;12(4).

Chandra RK: Nutrition, immunity and infection: from basic knowledge of dietary manipulation of immune responses to practical application of ameliorating suffering and improving survival. *Proceedings of the National Academy of Sciences of the United States of America* 93: 14304-14307, 1996.

Cotter SC, Reavey CE, Tummala Y, Randall JL, Holdbrook R, Ponton F, et al. Diet modulates the relationship between immune gene expression and functional immune responses. *Insect Biochemistry and Molecular Biology*. 2019;109:128-41.

Cunningham-Rundles S, McNeeley DF and Moon A: Mechanisms of nutrient modulation of the immune response. *The Journal of allergy and clinical immunology* 115: 1119-1128; quiz 1129, 2005.

Davison G, Kehaya C, Jones AW. Nutritional and Physical Activity Interventions to Improve Immunity. *American Journal of Lifestyle Medicine*. 2016;10(3):152-69.

El-Kadiki A. The Role of Micronutrients in Preventing Infections in the Elderly. Watson RR, editor 2015. 277-82 p.

Elmadfa I, Meyer AL. The Role of the Status of Selected Micronutrients in Shaping the Immune Function. *Endocrine Metabolic & Immune Disorders-Drug Targets*. 2019;19(8):1100-15.

Field CJ, Johnson IR, Schley PD. Nutrients and their role in host resistance to infection. *Journal of Leukocyte Biology*. 2002;71(1):16-32.

Forster SE, Powers HJ, Foulds GA, Flower DJ, Hopkinson K, Parker SG, et al. Improvement in Nutritional Status Reduces the Clinical Impact of Infections in Older Adults. *Journal of the American Geriatrics Society*. 2012;60(9):1645-54.

Gaertner C. The importance of nutrition in immunity. *Agro Food Industry Hi-Tech*. 2007;18(2):30-2.

Gariballa S. Vitamin and mineral supplements for preventing infections in older people - May have a place for some, but improved diet and physical activity will do more good. *British Medical Journal*. 2005;331(7512):304-5.

Gombart AF, Pierre A, Maggini S. A Review of Micronutrients and the Immune System-Working in Harmony to Reduce the Risk of Infection. *Nutrients*. 2020;12(1).

Grant WB, Lahore H, McDonnell SL, Baggerly CA, French CB, Aliano JL, et al. Evidence that Vitamin D Supplementation Could Reduce Risk of Influenza and COVID-19 Infections and Deaths. *Nutrients*. 2020;12(4).

Hemila H. Vitamin E administration may decrease the incidence of pneumonia in elderly males. *Clinical Interventions in Aging*. 2016;11:1379-85.

High KP. Micronutrient supplementation and immune function in the elderly. *Clinical Infectious Diseases*. 1999;28(4):717-22.

High KP. Nutrition and susceptibility to infection in the elderly. *Infectious Diseases in Clinical Practice*. 1999;8(1):34-40.

High KP. Nutritional strategies to boost immunity and prevent infection in elderly individuals. *Clinical Infectious Diseases*. 2001;33(11):1892-900.

Ibrahim KS, El-Sayed EM. Potential role of nutrients on immunity. *International Food Research Journal*. 2016;23(2):464-74.

Ibrahim MK, Zambruni M, Melby CL, Melby PC. Impact of Childhood Malnutrition on Host Defense and Infection. *Clinical Microbiology Reviews*. 2017;30(4):919-71.

Iddir M, Brito A, Dingeo G, et al.: Strengthening the Immune System and Reducing Inflammation and Oxidative Stress through Diet and Nutrition: Considerations during the COVID-19 Crisis. *Nutrients* 12, 2020.

Iliakis D, Kressig RW. The relationship between malnutrition and immune function in the elderly. *Therapeutische Umschau*. 2014;71(1):55-61.

Jahns L, Conrad Z, Johnson LK, Whigham LD, Wu D and Claycombe-Larson KJ: A diet high in carotenoid-rich vegetables and fruits favorably impacts inflammation status by increasing plasma concentrations of IFN-alpha2 and decreasing MIP-1beta and TNF-alpha in healthy individuals during a controlled feeding trial. *Nutrition research* (New York, N.Y.) 52: 98-104, 2018.

Leandro CG, Ferreira E Silva WT, Lima-Silva AE. Covid-19 and Exercise-Induced Immunomodulation. *Neuroimmunomodulation*. 2020:1-3.

Lesourd B, Ferry M. Nutrition and immune function in the elderly. In: Raats M, DeGroot L, VanStaveren W, editors. *Food for the Ageing Population*. Woodhead Publishing in Food Science Technology and Nutrition 2009. p. 271-87.

Maggini S, Pierre A, Calder PC. Immune Function and Micronutrient Requirements Change over the Life Course. *Nutrients*. 2018;10(10).

Maggini S, Wishart K, Wintergerst ES. Micronutrients and Ginseng for Immune Support in Older Adults. *Watson RR, editor* 2015. 265-75 p.

Maijo M, Clements SJ, Ivory K, Nicoletti C, Carding SR. Nutrition, diet and immunosenescence. *Mechanisms of Ageing and Development*. 2014;136:116-28.

Majde JA and Krueger JM: Links between the innate immune system and sleep. *The Journal of allergy and clinical immunology* 116: 1188-1198, 2005.

Marcos A, Nova E and Montero A: Changes in the immune system are conditioned by nutrition. *European journal of clinical nutrition* 57 Suppl 1: S66-69, 2003.

Messina G, Polito R, Monda V, Cipolloni L, Di Nunno N, Di Mizio G, et al. Functional Role of Dietary Intervention to Improve the Outcome of COVID-19: A Hypothesis of Work. *International journal of molecular sciences*. 2020;21(9).

Moazzen N, Imani B, Aelami MH, Haghi NSM, Kianifar HR, Khoushkhui M, et al. How to Boost your Immune System Against Coronavirus Infection? *Archives of Bone and Joint Surgery-Abjs*. 2020;8:220-5.

Nilashi M, Samad S, Yusuf SYM and Akbari E: Can complementary and alternative medicines be beneficial in the treatment of COVID-19 through improving immune system function? *Journal of infection and public health*, 2020.

Pae M, Meydani SN, Wu DY. The Role of Nutrition in Enhancing Immunity in Aging. *Aging and Disease*. 2012;3(1):91-129.

Pae M, Wu DY. Nutritional modulation of age-related changes in the immune system and risk of infection. *Nutrition Research*. 2017;41:14-35.



Ponton F, Wilson K, Holmes AJ, Cotter SC, Raubenheimer D, Simpson SJ. Integrating nutrition and immunology: A new frontier. *Journal of Insect Physiology*. 2013;59(2):130-7.

Prentice S. They Are what You eat: Can Nutritional Factors during Gestation and early infancy Modulate the Neonatal immune Response? *Frontiers in Immunology*. 2017;8.

Ribeiro KDD, Garcia LRS, Dametto JFD, Assuncao DGF, Maciel BLL. COVID-19 and Nutrition: The Need for Initiatives to Promote Healthy Eating and Prevent Obesity in Childhood. *Childhood Obesity*.

Rocha FAC, de Assis MR. Curcumin as a potential treatment for COVID-19. *Phytotherapy research : PTR*. 2020.

Roy A, Sarkar B, Celik C, Ghosh A, Basu U, Jana M, et al. Can concomitant use of zinc and curcumin with other immunity-boosting nutraceuticals be the arsenal against COVID-19? *Phytotherapy research : PTR*. 2020.

Saeed F, Nadeem M, Ahmed RS, Nadeem MT, Arshad MS and Ullah A: Studying the impact of nutritional immunology underlying the modulation of immune responses by nutritional compounds - a review. *Food and Agricultural Immunology* 27: 205-229, 2016.

Saeed F, Nadeem M, Ahmed RS, Nadeem MT, Arshad MS, Ullah A. Studying the impact of nutritional immunology underlying the modulation of immune responses by nutritional compounds - a review. *Food and Agricultural Immunology*. 2016;27(2):205-29.

Skalny AV, Rink L, Ajsuvakova OP, et al.: Zinc and respiratory tract infections: Perspectives for COVID-19 (Review). *International journal of molecular medicine*, 2020.

Suchner U, Kuhn KS, Furst P. The scientific basis of immunonutrition. *Proceedings of the Nutrition Society*. 2000;59(4):553-63.

Wu DY, Lewis ED, Pae MY, Meydani SN. Nutritional Modulation of Immune Function: Analysis of Evidence, Mechanisms, and Clinical Relevance. *Frontiers in Immunology*. 2019;9.

Abd El-Salam HS, Hassan AA. Phytochemicals Boost Anti-inflammatory Effect Against Gamma Radiation: Activities of Ginger and Coriander Extracts. *Arab Journal of Nuclear Sciences and Applications*. 2017;50(2):278-91.

Aibana O, Franke MF, Huang CC, Galea JT, Calderon R, Zhang ZB, et al. Vitamin E Status Is Inversely Associated with Risk of Incident Tuberculosis Disease among Household Contacts. *Journal of Nutrition*. 2018;148(1):56-62.

Al-Biltagi MAM, Abo-Elezz AAE, Abd-Elhafez MA, Mabrouk MM, Suliman GA. Beneficial Effects of Omega-3 Supplement to the Enteral Feeding in Children With Mild to Moderate Sepsis. *Journal of Intensive Care Medicine*. 2017;32(3):212-7.

Aryaeian N, Shahram F, Mahmoudi M, Tavakoli H, Yousefi B, Arablou T, et al. The effect of ginger supplementation on some immunity and inflammation intermediate genes expression in patients with active Rheumatoid Arthritis. *Gene*. 2019;698:179-85.

Aslam MM, John P, Bhatti A, Jahangir S, Kamboh MI. Vitamin D as a Principal Factor in Mediating Rheumatoid Arthritis-Derived Immune Response. *BioMed research international*. 2019;2019:3494937.

Bartosik-Psujek H, Psujek M. Vitamin D as an immune modulator in multiple sclerosis. *Neurologia i neurochirurgia polska*. 2019;53(2):113-22.

Batista TH, Alves Freire Ribeiro AC, Kalil B, Giusti-Paiva A, Vilela FC. Maternal protein malnutrition prolongs sickness behavior in male offspring. *Journal of Neuroimmunology*. 2020;341.

Bauer SR, Kapoor A, Rath M, Thomas SA. What is the role of supplementation with ascorbic acid, zinc, vitamin D, or N-acetylcysteine for prevention or treatment of COVID-19? *Cleveland Clinic journal of medicine*. 2020.

Besedovsky L, Lange T, Haack M. THE SLEEP-IMMUNE CROSSTALK IN HEALTH AND DISEASE. *Physiological Reviews*. 2019;99(3):1325-80.

Bessler H, Djaldetti M. Broccoli and human health: immunomodulatory effect of sulforaphane in a model of colon cancer. *International journal of food sciences and nutrition*. 2018;69(8):946-53.

Bobeck EA. NUTRITION AND HEALTH: COMPANION ANIMAL APPLICATIONS: Functional nutrition in livestock and companion animals to modulate the immune response. *Journal of Animal Science*. 2020;98(3).

Calder PC. Feeding the immune system. *Proceedings of the Nutrition Society*. 2013;72(3):299-309.

Campbell NK, Fitzgerald HK, Fletcher JM, Dunne A. Plant-Derived Polyphenols Modulate Human Dendritic Cell Metabolism and Immune Function via AMPK-Dependent Induction of Heme Oxygenase-1. *Frontiers in immunology*. 2019;10:345.

Cantorna MT, Snyder L, Arora J. Vitamin A and vitamin D regulate the microbial complexity, barrier function, and the mucosal immune responses to ensure intestinal homeostasis. *Critical reviews in biochemistry and molecular biology*. 2019;54(2):184-92.

Carr AC, Maggini S. Vitamin C and Immune Function. *Nutrients*. 2017;9(11).

Catanzaro M, Corsini E, Rosini M, Racchi M, Lanni C. Immunomodulators Inspired by Nature: A Review on Curcumin and Echinacea. *Molecules (Basel, Switzerland)*. 2018;23(11).

Chandra RK. Impact of nutritional status and nutrient supplements on immune responses and incidence of infection in older individuals. *Ageing Research Reviews*. 2004;3(1):91-104.

Choi J-W, Lee J, Lee J-H, Park B-J, Lee EJ, Shin S, et al. Omega-3 Polyunsaturated Fatty Acids Prevent *Toxoplasma gondii* Infection by Inducing Autophagy via AMPK Activation. *Nutrients*. 2019;11(9).

Colotta F, Jansson B, Bonelli F. Modulation of inflammatory and immune responses by vitamin D. *Journal of Autoimmunity*. 2017;85:78-97.

Craddock JC, Neale EP, Peoples GE, Probst YC. Vegetarian-Based Dietary Patterns and their Relation with Inflammatory and Immune Biomarkers: A Systematic Review and Meta-Analysis. *Advances in Nutrition*. 2019;10(3):433-51.

Darnton-Hill I, Ahmed F. Micronutrients: Immunological and Infection Effects on Nutritional Status and Impact on Health in Developing Countries. Bendich A, Deckelbaum RJ, editors 2009. 567-609 p.

Davison G, Kehaya C and Wyn Jones A: Nutritional and Physical Activity Interventions to Improve Immunity. *American journal of lifestyle medicine* 10: 152-169, 2016.

Dawson DR, Branch-Mays G, Gonzalez OA, Ebersole JL. Dietary modulation of the inflammatory cascade. *Periodontology* 2000. 2014;64(1):161-97.

Decoeur F, Benmamar-Badel A, Leyrolle Q, Persillet M, Laye S, Nadjar A. Dietary N-3 PUFA deficiency affects sleep-wake activity in basal condition and in response to an inflammatory challenge in mice. *Brain Behavior and Immunity*. 2020;85:162-9.

Del Giudice MM, Maiello N, Capristo C, Alterio E, Capasso M, Perrone L, et al. Resveratrol plus carboxymethyl-beta-glucan reduces nasal symptoms in children with pollen-induced allergic rhinitis. *Current Medical Research and Opinion*. 2014;30(10):1931-5.

Deshpande R, Raina P, Shinde K, Mansara P, Karandikar M, Kaul-Ghanekar R. Flax seed oil reduced tumor growth, modulated immune responses and decreased HPV E6 and E7 oncoprotein expression in a murine model of ectopic cervical cancer. *Prostaglandins & other lipid mediators*. 2019;143:106332.

Doh KC, Kim B-M, Kim KW, Chung BH, Yang CW. Effects of resveratrol on Th17 cell-related immune responses under tacrolimus-based immunosuppression. *BMC complementary and alternative medicine*. 2019;19(1):54.

Eda N, Ito H, Shimizu K, Suzuki S, Lee E, Akama T. Yoga stretching for improving salivary immune function and mental stress in middle-aged and older adults. *Journal of women & aging*. 2018;30(3):227-41.

Eom JS, Song J, Choi HS. Protective Effects of a Novel Probiotic Strain of *Lactobacillus plantarum* JSA22 from Traditional Fermented Soybean Food Against Infection by *Salmonella enterica* Serovar Typhimurium. *Journal of microbiology and biotechnology*. 2015;25(4):479-91.

Fan S, Li J, Bai B. Purification, structural elucidation and in vivo immunity-enhancing activity of polysaccharides from quinoa (*Chenopodium quinoa* Willd.) seeds. *Bioscience, biotechnology, and biochemistry*. 2019;83(12):2334-44.

Fernandez RCG, Perez AD, Saidler LM, Gomez JMQ. Vitamin D as immunity element against infection. *Medicina Clinica*. 2009;133(9):344-8.

Fernandez-Ruiz I. Exercise protects against cardiovascular disease by modulating immune cell supply. *Nature reviews Cardiology*. 2020;17(1):5.

Ferreira MJ, Irigoyen MC, Consolim-Colombo F, Saraiva JFK, Angelis KD. Physically Active Lifestyle as an Approach to Confronting COVID-19. *Arquivos brasileiros de cardiologia*. 2020;114(4):601-2.

Fox GJ, Lee RS, Lucas M, Khan FA, Proulx J-F, Hornby K, et al. Inadequate Diet Is Associated with Acquiring *Mycobacterium tuberculosis* Infection in an Inuit Community. A Case-Control Study. *Annals of the American Thoracic Society*. 2015;12(8):1153-62.

Gonzalez SM, Aguilar-Jimenez W, Trujillo-Gil E, Zapata W, Su R-C, Ball TB, et al. Vitamin D treatment of peripheral blood mononuclear cells modulated immune activation and reduced susceptibility to HIV-1 infection of CD4+ T lymphocytes. *PloS one*. 2019;14(9):e0222878.

Greiller CL, Martineau AR. Modulation of the Immune Response to Respiratory Viruses by Vitamin D. *Nutrients*. 2015;7(6):4240-70.

Guo N-H, Fu X, Zi F-M, Song Y, Wang S, Cheng J. The potential therapeutic benefit of resveratrol on Th17/Treg imbalance in immune thrombocytopenic purpura. *International immunopharmacology*. 2019;73:181-92.

Gutierrez S, Svahn SL, Johansson ME. Effects of Omega-3 Fatty Acids on Immune Cells. *International journal of molecular sciences*. 2019;20(20).

Han Y, Song M, Gu M, Ren D, Zhu X, Cao X, et al. Dietary Intake of Whole Strawberry Inhibited Colonic Inflammation in Dextran-Sulfate-Sodium-Treated Mice via Restoring Immune Homeostasis and Alleviating Gut Microbiota Dysbiosis. *Journal of agricultural and food chemistry*. 2019;67(33):9168-77.

Hussain MI, Ahmed W, Nasir M, Mushtaq MH, Sheikh AA, Shaheen AY, et al. Immune boosting role of vitamin E against pulmonary tuberculosis. *Pakistan journal of pharmaceutical sciences*. 2019;32(1(Supplementary)):269-76.

Husson M-O, Ley D, Portal C, Gottrand M, Hueso T, Desseyn J-L, et al. Modulation of host defence against bacterial and viral infections by omega-3 polyunsaturated fatty acids. *The Journal of infection*. 2016;73(6):523-35.

Hwang JH, McGovern J, Minett GM, Della Gatta PA, Roberts L, Harris JM, et al. Mobilizing serum factors and immune cells through exercise to counteract age-related changes in cancer risk. *Exercise immunology review*. 2020;26:80-99.

Jiang S, Pan ZY, Li H, Li FL, Song YY, Qiu Y. Meta-Analysis: Low-Dose Intake of Vitamin E Combined with Other Vitamins or Minerals May Decrease All-Cause Mortality. *Journal of Nutritional Science and Vitaminology*. 2014;60(3):194-205.

Jo S, Kim S, Shin DH, Kim M-S. Inhibition of SARS-CoV 3CL protease by flavonoids. *Journal of enzyme inhibition and medicinal chemistry*. 2020;35(1):145-51.

Kahkhaie KR, Mirhosseini A, Aliabadi A, Mohammadi A, Mousavi MJ, Haftcheshmeh SM, et al. Curcumin: a modulator of inflammatory signaling pathways in the immune system. *Inflammopharmacology*. 2019;27(5):885-900.

Karim T, Muhit M, Khandaker G. Interventions to prevent respiratory diseases - Nutrition and the developing world. *Paediatric Respiratory Reviews*. 2017;22:31-7.

Khammassi M, Ouerghi N, Said M, Feki M, Khammassi Y, Pereira B, et al. Continuous Moderate-Intensity but Not High-Intensity Interval Training Improves Immune Function Biomarkers in Healthy Young Men. *Journal of strength and conditioning research*. 2020;34(1):249-56.

Kohut ML, Lee W, Martin A, Arnston B, Russell DW, Ekkekakis P, et al. The exercise-induced enhancement of influenza immunity is mediated in part by improvements in psychosocial factors in older adults. *Brain, behavior, and immunity*. 2005;19(4):357-66.

Kroner JD, Sommer A, Fabri M. Vitamin D Every Day to Keep the Infection Away? *Nutrients*. 2015;7(6):4170-88.

Kubena KS, McMurray DN. Nutrition and the immune system: A review of nutrient-nutrient interactions. *Journal of the American Dietetic Association*. 1996;96(11):1156-64.

Kunisawa J. Immunity and Nutrition: the Role of Vitamins. Ratcliffe MJH, Cavazzana M, Cooke A, editors 2016. 120-6 p.

Langkamp-Henken B, Bender BS, Gardner EM, Herrlinger-Garcia KA, Kelley MJ, Murasko DM, et al. Nutritional formula enhanced immune function and reduced days of symptoms of upper respiratory tract infection in seniors. *Journal of the American Geriatrics Society*. 2004;52(1):3-12.

Langkamp-Henken B, Wood SM, Herlinger-Garcia KA, Thomas DJ, Stechmiller JK, Bender BS, et al. Nutritional formula improved immune profiles of seniors living in nursing homes. *Journal of the American Geriatrics Society*. 2006;54(12):1861-70.

Langley-Evans SC and Carrington LJ: Diet and the developing immune system. *Lupus* 15: 746-752, 2006.

Larbi A, Cexus O, Bosco N. Nutrition as a Tool to Reverse Immunosenescence? Chatterjee S, Jungraithmayr W, Bagchi D, editors 2018. 319-37 p.

Lesourd B. Nutritional factors and immunological ageing. *Proceedings of the Nutrition Society*. 2006;65(3):319-25.

Lewis ED, Meydani SN, Wu DY. Regulatory role of vitamin E in the immune system and inflammation. *Immunology Letters*. 2019;71(4):487-94.

Lobo de Sa FD, Butkevych E, Natramparasu PK, Fromm A, Mousavi S, Moos V, et al. Curcumin Mitigates Immune-Induced Epithelial Barrier Dysfunction by *Campylobacter jejuni*. *International journal of molecular sciences*. 2019;20(19).

Maggini S, Wintergerst ES, Beveridge S, Hornig DH. Selected vitamins and trace elements support immune function by strengthening epithelial barriers and cellular and humoral immune responses. *British Journal of Nutrition*. 2007;98:S29-S35.

Maggini S. Vitamins and Minerals: Contribution to Immune Function and Health. Watson RR, Zibadi S, Preedy VR, editors 2010. 227-52 p.

Mainardi T, Kapoor S and Bielory L: Complementary and alternative medicine: herbs, phytochemicals and vitamins and their immunologic effects. *The Journal of allergy and clinical immunology* 123: 283-294; quiz 295-286, 2009.

Maki KC, Kaspar KL, Khoo C, Derrig LH, Schild AL, Gupta K. Consumption of a cranberry juice beverage lowered the number of clinical urinary tract infection episodes in women with a recent history of urinary tract infection. *The American journal of clinical nutrition*. 2016;103(6):1434-42.

Malaguarnera L. Influence of Resveratrol on the Immune Response. *Nutrients*. 2019;11(5).

Mantzorou M, Giaginis C. Cranberry Consumption Against Urinary Tract Infections: Clinical State of the Art and Future Perspectives. *Current pharmaceutical biotechnology*. 2018;19(13):1049-63.

Martel SI, Riquelme SA, Kalergis AM, Bozinovic F. Dietary effect on immunological energetics in mice. *Journal of Comparative Physiology B-Biochemical Systems and Environmental Physiology*. 2014;184(7):937-44.

McCartney DM, Byrne DG. Optimisation of Vitamin D Status for Enhanced Immuno-protection Against Covid-19. *Irish medical journal*. 2020;113(4):58.

McCullough PJ, Lehrer DS. Vitamin D, cod liver oil, sunshine, and phototherapy: Safe, effective and forgotten tools for treating and curing tuberculosis infections - A comprehensive review. *The Journal of steroid biochemistry and molecular biology*. 2018;177:21-9.

Meydani SN. DIETARY MODULATION OF THE IMMUNE-RESPONSE IN THE AGED. *Age*. 1991;14(4):108-15.

Mil-Homens D, Ferreira-Dias S, Fialho AM. Fish oils against *Burkholderia* and *Pseudomonas aeruginosa*: invitro efficacy and their therapeutic and prophylactic effects on infected *Galleria mellonella* larvae. *Journal of applied microbiology*. 2016;120(6):1509-19.

Mocchegiani E, Romeo J, Malavolta M, Costarelli L, Giacconi R, Diaz L-E, et al. Zinc: dietary intake and impact of supplementation on immune function in elderly. *Age*. 2013;35(3):839-60.

Mollazadeh H, Cicero AFG, Blesso CN, Pirro M, Majeed M, Sahebkar A. Immune modulation by curcumin: The role of interleukin-10. *Critical reviews in food science and nutrition*. 2019;59(1):89-101.

Mukherjee S, Hussaini R, White R, Atwi D, Fried A, Sampat S, et al. TriCurin, a synergistic formulation of curcumin, resveratrol, and epicatechin gallate, repolarizes tumor-associated macrophages and triggers an immune response to cause suppression of HPV+ tumors. *Cancer immunology, immunotherapy : CII*. 2018;67(5):761-74.

Nantz MP, Rowe CA, Nieves C, Jr., Percival SS. Immunity and antioxidant capacity in humans is enhanced by consumption of a dried, encapsulated fruit and vegetable juice concentrate. *The Journal of nutrition*. 2006;136(10):2606-10.

Passos GS, Poyares D, Santana MG, Teixeira AAdS, Lira FS, Youngstedt SD, et al. Exercise improves immune function, antidepressive response, and sleep quality in patients with chronic primary insomnia. *BioMed research international*. 2014;2014:498961.

Puertollano MA, Puertollano E, Contreras-Moreno J, Ceron JM, de Cienfuegos GA, de Pablo MA. Natural Antioxidants and Resistance to Infection. Watson RR, Preedy VR, editors 2013. 157-74 p.

Puertollano MA, Puertollano E, de Cienfuegos GA, de Pablo MA. Dietary Antioxidants: Immunity and Host Defense. *Current Topics in Medicinal Chemistry*. 2011;11(14):1752-66.

Qin X, Liu J, Du Y, Li Y, Zheng L, Chen G, et al. Different doses of vitamin C supplementation enhances the Th1 immune response to early *Plasmodium yoelii* 17XL infection in BALB/c mice. *International immunopharmacology*. 2019;70:387-95.

Rak K, Bronkowska M. Immunomodulatory Effect of Vitamin D and Its Potential Role in the Prevention and Treatment of Type 1 Diabetes Mellitus-A Narrative Review. *Molecules (Basel, Switzerland)*. 2018;24(1).

Rivas E, Crandall CG, Suman OE, Moustaid-Moussa N, Ben-Ezra V. Exercise heat acclimation causes post-exercise hypotension and favorable improvements in lipid and immune profiles: A crossover randomized controlled trial. *Journal of thermal biology*. 2019;84:266-73.

Roth DE, Caulfield LE, Ezzati M, Black RE. Acute lower respiratory infections in childhood: opportunities for reducing the global burden through nutritional interventions. *Bulletin of the World Health Organization*. 2008;86(5):356-64.

Sanudo B, Seixas A, Gloeckl R, Rittweger J, Rawer R, Taiar R, et al. Potential Application of Whole Body Vibration Exercise For Improving The Clinical Conditions of COVID-19 Infected Individuals: A Narrative Review From the World Association of Vibration Exercise Experts (WAVex) Panel. *International journal of environmental research and public health*. 2020;17(10).

Sarkar S, Hewison M, Studzinski GP, Li YC, Kalia V. Role of vitamin D in cytotoxic T lymphocyte immunity to pathogens and cancer. *Critical Reviews in Clinical Laboratory Sciences*. 2016;53(2):132-45.

Sassi F, Tamone C, D'Amelio P. Vitamin D: Nutrient, Hormone, and Immunomodulator. *Nutrients*. 2018;10(11).

Senna SM, Torres MK, Pereira Lopes DA, Alheiros-Lira MC, de Moura DB, Alves Pereira VR, et al. Moderate physical training attenuates perinatal low-protein-induced spleen lymphocyte apoptosis in endotoxemic adult offspring rats. *European Journal of Nutrition*. 2016;55(3):1113-22.

Sharma S, Naura AS. Potential of phytochemicals as immune-regulatory compounds in atopic diseases: A review. *Biochemical Pharmacology*. 2020;173.

Skinner SJM, Hunter D, Cho S, Skinner M. The Potential Health Benefits of the Subtropical Fruits Kiwifruit, Feijoa and Tamarillo. Skinner M, Hunter D, editors 2013. 169-95 p.

Solomons NW. Malnutrition, immunity and infection. In: Calder PC, Yaqoob P, editors. *Diet, Immunity and Inflammation*. Woodhead Publishing Series in Food Science Technology and Nutrition. 2322013. p. 686-717.

Sreelatha A, Sam NV, Raghavan RP. Protective Effects of Vitamin D -A Review Article. *Journal of Young Pharmacists*. 2018;10(4):388-91.

Teter B, Morihara T, Lim GP, Chu T, Jones MR, Zuo X, et al. Curcumin restores innate immune Alzheimer's disease risk gene expression to ameliorate Alzheimer pathogenesis. *Neurobiology of disease*. 2019;127:432-48.

Vanherwegen AS, Gysemans C, Mathieu C. Regulation of Immune Function by Vitamin D and Its Use in Diseases of Immunity. *Endocrinology and Metabolism Clinics of North America*. 2017;46(4):1061-+.

Wentz LM, Ward MD, Potter C, Oliver SJ, Jackson S, Izard RM, et al. Increased Risk of Upper Respiratory Infection in Military Recruits Who Report Sleeping Less Than 6 h per night. *Military medicine*. 2018;183(11-12):e699-e704.

Wessels I, Rink L. Micronutrients in autoimmune diseases: possible therapeutic benefits of zinc and vitamin D. *Journal of Nutritional Biochemistry*. 2020;77.

Williams NC, Killer SC, Svendsen IS, Jones AW. Immune nutrition and exercise: Narrative review and practical recommendations. *European journal of sport science*. 2019;19(1):49-61.

Wintergerst ES, Maggini S, Hornig DH. Contribution of selected vitamins and trace elements to immune function. *Annals of Nutrition and Metabolism*. 2007;51(4):301-23.

Witek Janusek L, Tell D, Mathews HL. Mindfulness based stress reduction provides psychological benefit and restores immune function of women newly diagnosed with breast cancer: A randomized trial with active control. *Brain, behavior, and immunity*. 2019;80:358-73.

Yang H, Sun Y, Cai R, Chen Y and Gu B: The impact of dietary fiber and probiotics in infectious diseases. *Microbial pathogenesis* 140: 103931, 2020.

You J, Chang Y, Zhao D, Zhuang J, Zhuang W. A Mixture of Functional Complex Extracts from *Lycium barbarum* and Grape Seed Enhances Immunity Synergistically In Vitro and In Vivo. *Journal of food science*. 2019;84(6):1577-85.

Zapatera B, Prados A, Gomez-Martinez S, Marcos A. Immunonutrition: methodology and applications. *Nutricion Hospitalaria*. 2015;31:145-54.

Zhou R, Wang X, Liu H, Guo L, Su Q, Wang H, et al. GalNAc-Specific Soybean Lectin Inhibits HIV Infection of Macrophages through Induction of Antiviral Factors. *Journal of virology*. 2018;92(6)."