List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3363166/publications.pdf Version: 2024-02-01



LAMIE ROSSIOHN

#	Article	IF	CITATIONS
1	MR1 presents microbial vitamin B metabolites to MAIT cells. Nature, 2012, 491, 717-723.	13.7	1,158
2	T-cell activation by transitory neo-antigens derived from distinct microbial pathways. Nature, 2014, 509, 361-365.	13.7	731
3	More than one reason to rethink the use of peptides in vaccine design. Nature Reviews Drug Discovery, 2007, 6, 404-414.	21.5	692
4	The burgeoning family of unconventional T cells. Nature Immunology, 2015, 16, 1114-1123.	7.0	655
5	Immune self-reactivity triggered by drug-modified HLA-peptide repertoire. Nature, 2012, 486, 554-558.	13.7	612
6	T Cell Antigen Receptor Recognition of Antigen-Presenting Molecules. Annual Review of Immunology, 2015, 33, 169-200.	9.5	603
7	CD1d–lipid-antigen recognition by the semi-invariant NKT T-cell receptor. Nature, 2007, 448, 44-49.	13.7	533
8	Antigen-loaded MR1 tetramers define T cell receptor heterogeneity in mucosal-associated invariant T cells. Journal of Experimental Medicine, 2013, 210, 2305-2320.	4.2	516
9	HLA variation and disease. Nature Reviews Immunology, 2018, 18, 325-339.	10.6	487
10	Structure of a Cholesterol-Binding, Thiol-Activated Cytolysin and a Model of Its Membrane Form. Cell, 1997, 89, 685-692.	13.5	457
11	Comprehensive, Quantitative Mapping of T Cell Epitopes in Gluten in Celiac Disease. Science Translational Medicine, 2010, 2, 41ra51.	5.8	393
12	Recognition of CD1d-restricted antigens by natural killer T cells. Nature Reviews Immunology, 2012, 12, 845-857.	10.6	382
13	Impairment of immunity to <i>Candida</i> and <i>Mycobacterium</i> in humans with bi-allelic <i>RORC</i> mutations. Science, 2015, 349, 606-613.	6.0	366
14	A molecular basis for the association of the <i>HLA-DRB1</i> locus, citrullination, and rheumatoid arthritis. Journal of Experimental Medicine, 2013, 210, 2569-2582.	4.2	354
15	AB5 subtilase cytotoxin inactivates the endoplasmic reticulum chaperone BiP. Nature, 2006, 443, 548-552.	13.7	351
16	Butyrophilin 3A1 binds phosphorylated antigens and stimulates human γδT cells. Nature Immunology, 2013, 14, 908-916.	7.0	351
17	Identification of phenotypically and functionally heterogeneous mouse mucosal-associated invariant T cells using MR1 tetramers. Journal of Experimental Medicine, 2015, 212, 1095-1108.	4.2	348
18	The Mechanism of Membrane Insertion for a Cholesterol-Dependent Cytolysin. Cell, 1999, 99, 293-299.	13.5	347

#	Article	IF	CITATIONS
19	Structural determinants of T-cell receptor bias in immunity. Nature Reviews Immunology, 2006, 6, 883-894.	10.6	322
20	A Structural Basis for the Selection of Dominant αβ T Cell Receptors in Antiviral Immunity. Immunity, 2003, 18, 53-64.	6.6	321
21	Identification of a Membrane-Spanning Domain of the Thiol-Activated Pore-Forming ToxinClostridium perfringensPerfringolysin O: An α-Helical to β-Sheet Transition Identified by Fluorescence Spectroscopyâ€. Biochemistry, 1998, 37, 14563-14574.	1.2	309
22	Human Leukocyte Antigen Class I-Restricted Activation of CD8+ T Cells Provides the Immunogenetic Basis of a Systemic Drug Hypersensitivity. Immunity, 2008, 28, 822-832.	6.6	309
23	A three-stage intrathymic development pathway for the mucosal-associated invariant T cell lineage. Nature Immunology, 2016, 17, 1300-1311.	7.0	288
24	T cell receptor recognition of a 'super-bulged' major histocompatibility complex class I–bound peptide. Nature Immunology, 2005, 6, 1114-1122.	7.0	280
25	Mucosal-associated invariant T cell alterations in obese and type 2 diabetic patients. Journal of Clinical Investigation, 2015, 125, 1752-1762.	3.9	272
26	A Common Fold Mediates Vertebrate Defense and Bacterial Attack. Science, 2007, 317, 1548-1551.	6.0	261
27	Recognition of vitamin B metabolites by mucosal-associated invariant T cells. Nature Communications, 2013, 4, 2142.	5.8	261
28	CD1d-lipid antigen recognition by the $\hat{I}^{3}\hat{I}$ TCR. Nature Immunology, 2013, 14, 1137-1145.	7.0	256
29	Structural basis of a unique interferon-β signaling axis mediated via the receptor IFNAR1. Nature Immunology, 2013, 14, 901-907.	7.0	255
30	Human mucosal-associated invariant T cells contribute to antiviral influenza immunity via IL-18–dependent activation. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 10133-10138.	3.3	246
31	A molecular basis underpinning the T cell receptor heterogeneity of mucosal-associated invariant T cells. Journal of Experimental Medicine, 2014, 211, 1585-1600.	4.2	245
32	The structural basis of Janus kinase 2 inhibition by a potent and specific pan-Janus kinase inhibitor. Blood, 2006, 107, 176-183.	0.6	243
33	Unconventional T Cell Targets for Cancer Immunotherapy. Immunity, 2018, 48, 453-473.	6.6	242
34	T Cell Allorecognition via Molecular Mimicry. Immunity, 2009, 31, 897-908.	6.6	232
35	Crystal structure of the N-terminal, growth factor-like domain of Alzheimer amyloid precursor protein. Nature Structural Biology, 1999, 6, 327-331.	9.7	229
36	Dissecting Specificity in the Janus Kinases: The Structures of JAK-Specific Inhibitors Complexed to the JAK1 and JAK2 Protein Tyrosine Kinase Domains. Journal of Molecular Biology, 2009, 387, 219-232.	2.0	225

#	Article	IF	CITATIONS
37	Incorporation of a non-human glycan mediates human susceptibility to a bacterial toxin. Nature, 2008, 456, 648-652.	13.7	217
38	Cytotoxic and regulatory roles of mucosal-associated invariant T cells in type 1 diabetes. Nature Immunology, 2017, 18, 1321-1331.	7.0	217
39	Mucosal-associated invariant T-cell activation and accumulation after in vivo infection depends on microbial riboflavin synthesis and co-stimulatory signals. Mucosal Immunology, 2017, 10, 58-68.	2.7	216
40	Understanding the drivers of MHC restriction of T cell receptors. Nature Reviews Immunology, 2018, 18, 467-478.	10.6	214
41	A T cell receptor flattens a bulged antigenic peptide presented by a major histocompatibility complex class I molecule. Nature Immunology, 2007, 8, 268-276.	7.0	206
42	Structure, biological functions and applications of the AB5 toxins. Trends in Biochemical Sciences, 2010, 35, 411-418.	3.7	204
43	Alternative cross-priming through CCL17-CCR4-mediated attraction of CTLs toward NKT cell–licensed DCs. Nature Immunology, 2010, 11, 313-320.	7.0	204
44	The insulin A-chain epitope recognized by human T cells is posttranslationally modified. Journal of Experimental Medicine, 2005, 202, 1191-1197.	4.2	201
45	CD94-NKG2A recognition of human leukocyte antigen (HLA)-E bound to an HLA class I leader sequence. Journal of Experimental Medicine, 2008, 205, 725-735.	4.2	198
46	Differential Recognition of CD1d-α-Galactosyl Ceramide by the Vβ8.2 and Vβ7 Semi-invariant NKT T Cell Receptors. Immunity, 2009, 31, 47-59.	6.6	198
47	Interferon-ε Protects the Female Reproductive Tract from Viral and Bacterial Infection. Science, 2013, 339, 1088-1092.	6.0	197
48	Functional role of T-cell receptor nanoclusters in signal initiation and antigen discrimination. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E5454-63.	3.3	194
49	A Naturally Selected Dimorphism within the HLA-B44 Supertype Alters Class I Structure, Peptide Repertoire, and T Cell Recognition. Journal of Experimental Medicine, 2003, 198, 679-691.	4.2	192
50	GABA production by glutamic acid decarboxylase is regulated by a dynamic catalytic loop. Nature Structural and Molecular Biology, 2007, 14, 280-286.	3.6	189
51	The CDR3 regions of an immunodominant T cell receptor dictate the 'energetic landscape' of peptide-MHC recognition. Nature Immunology, 2005, 6, 171-180.	7.0	187
52	A conserved human T cell population targets mycobacterial antigens presented by CD1b. Nature Immunology, 2013, 14, 706-713.	7.0	187
53	Genome-wide CRISPR–Cas9 screening reveals ubiquitous T cell cancer targeting via the monomorphic MHC class I-related protein MR1. Nature Immunology, 2020, 21, 178-185.	7.0	186
54	Dominant protection from HLA-linked autoimmunity by antigen-specific regulatory T cells. Nature, 2017, 545, 243-247.	13.7	181

#	Article	IF	CITATIONS
55	Human CD8+ T cell cross-reactivity across influenza A, B and C viruses. Nature Immunology, 2019, 20, 613-625.	7.0	180
56	T-cell receptor recognition of HLA-DQ2–gliadin complexes associated with celiac disease. Nature Structural and Molecular Biology, 2014, 21, 480-488.	3.6	177
57	MAIT cells protect against pulmonary Legionella longbeachae infection. Nature Communications, 2018, 9, 3350.	5.8	177
58	SARS-CoV-2 mRNA vaccination elicits a robust and persistent T follicular helper cell response in humans. Cell, 2022, 185, 603-613.e15.	13.5	176
59	Drugs and drug-like molecules can modulate the function of mucosal-associated invariant T cells. Nature Immunology, 2017, 18, 402-411.	7.0	175
60	Two Structural Transitions in Membrane Pore Formation by Pneumolysin, the Pore-Forming Toxin of Streptococcus pneumoniae. Cell, 1999, 97, 647-655.	13.5	174
61	Killer cell immunoglobulin-like receptor 3DL1-mediated recognition of human leukocyte antigen B. Nature, 2011, 479, 401-405.	13.7	174
62	The structures of human glutathione transferase P1-1 in complex with glutathione and various inhibitors at high resolution. Journal of Molecular Biology, 1997, 274, 84-100.	2.0	172
63	Diversity of T Cells Restricted by the MHC Class I-Related Molecule MR1 Facilitates Differential Antigen Recognition. Immunity, 2016, 44, 32-45.	6.6	169
64	Suboptimal SARS-CoV-2â^'specific CD8 <sup>+</sup> T cell response associated with the prominent HLA-A*02:01 phenotype. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 24384-24391.	3.3	168
65	HLA-DQA1–HLA-DRB1 variants confer susceptibility to pancreatitis induced by thiopurine immunosuppressants. Nature Genetics, 2014, 46, 1131-1134.	9.4	165
66	A mixed disulfide bond in bacterial glutathione transferase: functional and evolutionary implications. Structure, 1998, 6, 721-734.	1.6	163
67	Cross-reactive CD8 <sup>+</sup> T-cell immunity between the pandemic H1N1-2009 and H1N1-1918 influenza A viruses. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 12599-12604.	3.3	163
68	High Resolution Structures of Highly Bulged Viral Epitopes Bound to Major Histocompatibility Complex Class I. Journal of Biological Chemistry, 2005, 280, 23900-23909.	1.6	162
69	Natural HLA Class I Polymorphism Controls the Pathway of Antigen Presentation and Susceptibility to Viral Evasion. Journal of Experimental Medicine, 2004, 200, 13-24.	4.2	159
70	Structural insight into MR1-mediated recognition of the mucosal associated invariant T cell receptor. Journal of Experimental Medicine, 2012, 209, 761-774.	4.2	159
71	Challenges, Progress, and Prospects of Developing Therapies to Treat Autoimmune Diseases. Cell, 2020, 181, 63-80.	13.5	159
72	The 2.0-Ã Crystal Structure of eqFP611, a Far Red Fluorescent Protein from the Sea Anemone Entacmaea quadricolor. Journal of Biological Chemistry, 2003, 278, 44626-44631.	1.6	158

#	Article	IF	CITATIONS
73	A Structural and Immunological Basis for the Role of Human Leukocyte Antigen DQ8 in Celiac Disease. Immunity, 2007, 27, 23-34.	6.6	157
74	DEC-205 is a cell surface receptor for CpG oligonucleotides. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 16270-16275.	3.3	155
75	Crystal Structure of the SARS-CoV-2 Non-structural Protein 9, Nsp9. IScience, 2020, 23, 101258.	1.9	155
76	Structural basis for the killing of human beta cells by CD8+ T cells in type 1 diabetes. Nature Immunology, 2012, 13, 283-289.	7.0	151
77	A Molecular Basis for the Control of Preimmune Escape Variants by HIV-Specific CD8+ T Cells. Immunity, 2013, 38, 425-436.	6.6	149
78	Crystal structure of the human T cell receptor CD3ÂÂ heterodimer complexed to the therapeutic mAb OKT3. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 7675-7680.	3.3	148
79	Human theta class glutathione transferase: the crystal structure reveals a sulfate-binding pocket within a buried active site. Structure, 1998, 6, 309-322.	1.6	147
80	Drug Hypersensitivity and Human Leukocyte Antigens of the Major Histocompatibility Complex. Annual Review of Pharmacology and Toxicology, 2012, 52, 401-431.	4.2	146
81	Preexisting CD8 <sup>+</sup> T-cell immunity to the H7N9 influenza A virus varies across ethnicities. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 1049-1054.	3.3	144
82	Germline-encoded recognition of diverse glycolipids by natural killer T cells. Nature Immunology, 2007, 8, 1105-1113.	7.0	143
83	Lack of prominent peptide–major histocompatibility complex features limits repertoire diversity in virus-specific CD8+ T cell populations. Nature Immunology, 2005, 6, 382-389.	7.0	142
84	A subset of HLA-I peptides are not genomically templated: Evidence for cis- and trans-spliced peptide ligands. Science Immunology, 2018, 3, .	5.6	142
85	CD1a-autoreactive T cells recognize natural skin oils that function as headless antigens. Nature Immunology, 2014, 15, 177-185.	7.0	141
86	Crystal structure of HLA-G: A nonclassical MHC class I molecule expressed at the fetal-maternal interface. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 3360-3365.	3.3	139
87	CD1a on Langerhans cells controls inflammatory skin disease. Nature Immunology, 2016, 17, 1159-1166.	7.0	134
88	Recognition of microbial and mammalian phospholipid antigens by NKT cells with diverse TCRs. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 1827-1832.	3.3	129
89	The 2.2 Ã Crystal Structure of a Pocilloporin Pigment Reveals a Nonplanar Chromophore Conformation. Structure, 2003, 11, 275-284.	1.6	127
90	The intracellular pathway for the presentation of vitamin B–related antigens by the antigen-presenting molecule MR1. Nature Immunology, 2016, 17, 531-537.	7.0	127

#	Article	IF	CITATIONS
91	A comprehensive review and performance evaluation of bioinformatics tools for HLA class I peptide-binding prediction. Briefings in Bioinformatics, 2020, 21, 1119-1135.	3.2	127
92	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. PLoS Biology, 2017, 15, e2001930.	2.6	126
93	Chronic Inflammation Permanently Reshapes Tissue-Resident Immunity in Celiac Disease. Cell, 2019, 176, 967-981.e19.	13.5	126
94	The Three-Dimensional Structure of the Human Pi Class Glutathione Transferase P1-1 in Complex with the Inhibitor Ethacrynic Acid and Its Glutathione Conjugate,. Biochemistry, 1997, 36, 576-585.	1.2	125
95	Have we cut ourselves too short in mapping CTL epitopes?. Trends in Immunology, 2006, 27, 11-16.	2.9	124
96	Recognition of Vitamin B Precursors and Byproducts by Mucosal Associated Invariant T Cells. Journal of Biological Chemistry, 2015, 290, 30204-30211.	1.6	123
97	Molecular basis for universal HLA-A*0201–restricted CD8 <sup>+</sup> T-cell immunity against influenza viruses. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 4440-4445.	3.3	122
98	The crystal structure of myelin oligodendrocyte glycoprotein, a key autoantigen in multiple sclerosis. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 11059-11064.	3.3	121
99	Biased T Cell Receptor Usage Directed against Human Leukocyte Antigen DQ8-Restricted Gliadin Peptides Is Associated with Celiac Disease. Immunity, 2012, 37, 611-621.	6.6	121
100	Functional Heterogeneity and Antimycobacterial Effects of Mouse Mucosal-Associated Invariant T Cells Specific for Riboflavin Metabolites. Journal of Immunology, 2015, 195, 587-601.	0.4	121
101	Lipid and small-molecule display by CD1 and MR1. Nature Reviews Immunology, 2015, 15, 643-654.	10.6	120
102	Arresting Pore Formation of a Cholesterol-dependent Cytolysin by Disulfide Trapping Synchronizes the Insertion of the Transmembrane β-Sheet from a Prepore Intermediate. Journal of Biological Chemistry, 2001, 276, 8261-8268.	1.6	118
103	The major histocompatibility complex class Ib molecule HLAâ€E at the interface between innate and adaptive immunity. Tissue Antigens, 2008, 72, 415-424.	1.0	118
104	A Molecular Basis for NKT Cell Recognition of CD1d-Self-Antigen. Immunity, 2011, 34, 315-326.	6.6	118
105	T cell receptor reversed polarity recognition of a self-antigen major histocompatibility complex. Nature Immunology, 2015, 16, 1153-1161.	7.0	115
106	β-Amino acid-containing hybrid peptides—new opportunities in peptidomimetics. Organic and Biomolecular Chemistry, 2007, 5, 2884.	1.5	114
107	Stabilizing short-lived Schiff base derivatives of 5-aminouracils that activate mucosal-associated invariant T cells. Nature Communications, 2017, 8, 14599.	5.8	113
108	Hotspot autoimmune T cell receptor binding underlies pathogen and insulin peptide cross-reactivity. Journal of Clinical Investigation, 2016, 126, 2191-2204.	3.9	113

#	Article	IF	CITATIONS
109	αβ T cell antigen receptor recognition of CD1a presenting self lipid ligands. Nature Immunology, 2015, 16, 258-266.	7.0	112
110	Antigen Ligation Triggers a Conformational Change within the Constant Domain of the $\hat{l}\pm\hat{l}^2$ T Cell Receptor. Immunity, 2009, 30, 777-788.	6.6	111
111	Recognition of β-linked self glycolipids mediated by natural killer T cell antigen receptors. Nature Immunology, 2011, 12, 827-833.	7.0	111
112	Polymorphism in Human Cytomegalovirus UL40 Impacts on Recognition of Human Leukocyte Antigen-E (HLA-E) by Natural Killer Cells. Journal of Biological Chemistry, 2013, 288, 8679-8690.	1.6	111
113	Structural basis for a major histocompatibility complex class Ib–restricted T cell response. Nature Immunology, 2006, 7, 256-264.	7.0	109
114	Human TRAV1-2-negative MR1-restricted T cells detect S. pyogenes and alternatives to MAIT riboflavin-based antigens. Nature Communications, 2016, 7, 12506.	5.8	108
115	A Molecular Basis for the Exquisite CD1d-Restricted Antigen Specificity and Functional Responses of Natural Killer T Cells. Immunity, 2011, 34, 327-339.	6.6	107
116	Molecular architecture of the αβ T cell receptor–CD3 complex. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 17576-17581.	3.3	107
117	Recognition of CD1d-sulfatide mediated by a type II natural killer T cell antigen receptor. Nature Immunology, 2012, 13, 857-863.	7.0	106
118	CD8+ TÂcells specific for an immunodominant SARS-CoV-2 nucleocapsid epitope display high naive precursor frequency and TCR promiscuity. Immunity, 2021, 54, 1066-1082.e5.	6.6	106
119	A structural basis for selection and cross-species reactivity of the semi-invariant NKT cell receptor in CD1d/glycolipid recognition. Journal of Experimental Medicine, 2006, 203, 661-673.	4.2	105
120	Escape from highly effective public CD8+ T-cell clonotypes by HIV. Blood, 2011, 118, 2138-2149.	0.6	103
121	Hard wiring of T cell receptor specificity for the major histocompatibility complex is underpinned by TCR adaptability. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 10608-10613.	3.3	101
122	The molecular mechanism of pneumolysin, a virulence factor from Streptococcus pneumoniae 1 1Edited by J. Thornton. Journal of Molecular Biology, 1998, 284, 449-461.	2.0	100
123	Antigen recognition by CD1d-restricted NKT T cell receptors. Seminars in Immunology, 2010, 22, 61-67.	2.7	100
124	Tc17 cells are a proinflammatory, plastic lineage of pathogenic CD8+ T cells that induce GVHD without antileukemic effects. Blood, 2015, 126, 1609-1620.	0.6	98
125	Structural and regulatory diversity shape HLA-C protein expression levels. Nature Communications, 2017, 8, 15924.	5.8	98
126	A class of Î <sup>3</sup> δT cell receptors recognize the underside of the antigen-presenting molecule MR1. Science, 2019, 366, 1522-1527.	6.0	98

#	Article	IF	CITATIONS
127	The Murine Orthologue of Human Antichymotrypsin. Journal of Biological Chemistry, 2005, 280, 43168-43178.	1.6	97
128	A semi-invariant Vα10+ T cell antigen receptor defines a population of natural killer T cells with distinct glycolipid antigen–recognition properties. Nature Immunology, 2011, 12, 616-623.	7.0	97
129	Mucosal-associated invariant T cells promote inflammation and intestinal dysbiosis leading to metabolic dysfunction during obesity. Nature Communications, 2020, 11, 3755.	5.8	97
130	A bird's eye view of <scp>NK</scp> cell receptor interactions with their <scp>MHC</scp> class I ligands. Immunological Reviews, 2015, 267, 148-166.	2.8	96
131	The Shaping of T Cell Receptor Recognition by Self-Tolerance. Immunity, 2009, 30, 193-203.	6.6	94
132	CTL Recognition of a Bulged Viral Peptide Involves Biased TCR Selection. Journal of Immunology, 2005, 175, 3826-3834.	0.4	93
133	Natural micropolymorphism in human leukocyte antigens provides a basis for genetic control of antigen recognition. Journal of Experimental Medicine, 2009, 206, 209-219.	4.2	93
134	The Fidelity, Occasional Promiscuity, and Versatility of T Cell Receptor Recognition. Immunity, 2008, 28, 304-314.	6.6	92
135	T cell receptor cross-reactivity between gliadin and bacterial peptides in celiac disease. Nature Structural and Molecular Biology, 2020, 27, 49-61.	3.6	91
136	Human Glutathione Transferase P1-1 and Nitric Oxide Carriers. Journal of Biological Chemistry, 2001, 276, 42138-42145.	1.6	90
137	T Cell Receptor CDR2β and CDR3β Loops Collaborate Functionally to Shape the iNKT Cell Repertoire. Immunity, 2009, 31, 60-71.	6.6	90
138	MAIT cells are depleted early but retain functional cytokine expression in HIV infection. Immunology and Cell Biology, 2015, 93, 177-188.	1.0	90
139	Peptide length determines the outcome of TCR/peptide-MHCI engagement. Blood, 2013, 121, 1112-1123.	0.6	89
140	Naturally Processed Non-canonical HLA-A*02:01 Presented Peptides. Journal of Biological Chemistry, 2015, 290, 2593-2603.	1.6	89
141	Molecular basis of glutathione synthetase deficiency and a rare gene permutation event. EMBO Journal, 1999, 18, 3204-3213.	3.5	88
142	Subtle Changes in Peptide Conformation Profoundly Affect Recognition of the Non-Classical MHC Class I Molecule HLA-E by the CD94–NKG2 Natural Killer Cell Receptors. Journal of Molecular Biology, 2008, 377, 1297-1303.	2.0	88
143	The Identification and Structure of the Membrane-spanning Domain of the Clostridium septicum Alpha Toxin. Journal of Biological Chemistry, 2004, 279, 14315-14322.	1.6	87
144	Structures of Perfringolysin O Suggest a Pathway for Activation of Cholesterol-dependent Cytolysins. Journal of Molecular Biology, 2007, 367, 1227-1236.	2.0	87

#	Article	IF	CITATIONS
145	The Heterodimeric Assembly of the CD94-NKG2 Receptor Family and Implications for Human Leukocyte Antigen-E Recognition. Immunity, 2007, 27, 900-911.	6.6	87
146	Human autoreactive T cells recognize CD1b and phospholipids. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 380-385.	3.3	85
147	Bimolecular Interaction of Insulin-Like Growth Factor (IGF) Binding Protein-2 with αvβ3 Negatively Modulates IGF-I-Mediated Migration and Tumor Growth 1. Cancer Research, 2004, 64, 977-984.	0.4	83
148	A minimal binding footprint on CD1d-glycolipid is a basis for selection of the unique human NKT TCR. Journal of Experimental Medicine, 2008, 205, 939-949.	4.2	83
149	MHC-I peptides get out of the groove and enable a novel mechanism of HIV-1 escape. Nature Structural and Molecular Biology, 2017, 24, 387-394.	3.6	83
150	The immunogenicity of a viral cytotoxic T cell epitope is controlled by its MHC-bound conformation. Journal of Experimental Medicine, 2005, 202, 1249-1260.	4.2	82
151	Natural Killer T cell obsession with self-antigens. Current Opinion in Immunology, 2013, 25, 168-173.	2.4	82
152	The 1.6ÂÃ Crystal Structure of the Catalytic Domain of PlyB, a Bacteriophage Lysin Active Against Bacillus anthracis. Journal of Molecular Biology, 2007, 366, 540-550.	2.0	81
153	Allelic polymorphism in the T cell receptor and its impact on immune responses. Journal of Experimental Medicine, 2010, 207, 1555-1567.	4.2	81
154	A structural voyage toward an understanding of the <scp>MHC</scp> â€lâ€restricted immune response: lessons learned and much to be learned. Immunological Reviews, 2012, 250, 61-81.	2.8	81
155	Killer cell immunoglobulin-like receptor 3DL1 polymorphism defines distinct hierarchies of HLA class I recognition. Journal of Experimental Medicine, 2016, 213, 791-807.	4.2	81
156	Modulation of innate and adaptive immunity by cytomegaloviruses. Nature Reviews Immunology, 2020, 20, 113-127.	10.6	80
157	The 2.1Ã Crystal Structure of the Far-red Fluorescent Protein HcRed: Inherent Conformational Flexibility of the Chromophore. Journal of Molecular Biology, 2005, 349, 223-237.	2.0	79
158	The 1.7ÂÃ Crystal Structure of Dronpa: A Photoswitchable Green Fluorescent Protein. Journal of Molecular Biology, 2006, 364, 213-224.	2.0	79
159	A BAFF antagonist suppresses experimental autoimmune encephalomyelitis by targeting cell-mediated and humoral immune responses. International Immunology, 2006, 18, 1473-1485.	1.8	79
160	Recipient mucosal-associated invariant T cells control GVHD within the colon. Journal of Clinical Investigation, 2018, 128, 1919-1936.	3.9	78
161	Human leukocyte antigen-associated drug hypersensitivity. Current Opinion in Immunology, 2013, 25, 81-89.	2.4	76
162	IL-23 costimulates antigen-specific MAIT cell activation and enables vaccination against bacterial infection. Science Immunology, 2019, 4, .	5.6	75

#	Article	IF	CITATIONS
163	Variations on the GFP Chromophore. Journal of Biological Chemistry, 2005, 280, 2401-2404.	1.6	74
164	Cleaved antitrypsin polymers at atomic resolution. Protein Science, 2000, 9, 417-420.	3.1	73
165	Reversed T Cell Receptor Docking on a Major Histocompatibility Class I Complex Limits Involvement in the Immune Response. Immunity, 2016, 45, 749-760.	6.6	73
166	The interplay between citrullination and HLA-DRB1 polymorphism in shaping peptide binding hierarchies in rheumatoid arthritis. Journal of Biological Chemistry, 2018, 293, 3236-3251.	1.6	73
167	Recognition of nectin-2 by the natural killer cell receptor T cell immunoglobulin and ITIM domain (TIGIT). Journal of Biological Chemistry, 2017, 292, 11413-11422.	1.6	72
168	The Structure of HLA-B8 Complexed to an Immunodominant Viral Determinant: Peptide-Induced Conformational Changes and a Mode of MHC Class I Dimerization. Journal of Immunology, 2002, 169, 5153-5160.	0.4	71
169	Phospholipid signaling in innate immune cells. Journal of Clinical Investigation, 2018, 128, 2670-2679.	3.9	71
170	The crystal structure of glucose dehydrogenase from Thermoplasma acidophilum. Structure, 1994, 2, 385-393.	1.6	70
171	Cutting Edge: CD1a Tetramers and Dextramers Identify Human Lipopeptide–Specific T Cells Ex Vivo. Journal of Immunology, 2013, 191, 4499-4503.	0.4	70
172	STAT3 is a critical cell-intrinsic regulator of human unconventional T cell numbers and function. Journal of Experimental Medicine, 2015, 212, 855-864.	4.2	70
173	The High Resolution Crystal Structure of the Human Tumor Suppressor Maspin Reveals a Novel Conformational Switch in the G-helix. Journal of Biological Chemistry, 2005, 280, 22356-22364.	1.6	69
174	Phosphorylated self-peptides alter human leukocyte antigen class I-restricted antigen presentation and generate tumor-specific epitopes. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 2776-2781.	3.3	69
175	New ways to turn on NKT cells. Journal of Experimental Medicine, 2011, 208, 1121-1125.	4.2	69
176	MAITs, MR1 and vitamin B metabolites. Current Opinion in Immunology, 2014, 26, 7-13.	2.4	69
177	Self-interaction of pneumolysin, the pore-forming protein toxin of Streptococcus pneumoniae. Journal of Molecular Biology, 1998, 284, 1223-1237.	2.0	68
178	T-cell receptor bias and immunity. Current Opinion in Immunology, 2008, 20, 119-125.	2.4	68
179	The structural basis for autonomous dimerization of the pre-T-cell antigen receptor. Nature, 2010, 467, 844-848.	13.7	68
180	The 1.5 Ã Crystal Structure of a Highly Selected Antiviral T Cell Receptor Provides Evidence for a Structural Basis of Immunodominance. Structure, 2002, 10, 1521-1532.	1.6	67

#	Article	IF	CITATIONS
181	Peptide-Dependent Recognition of HLA-B*57:01 by KIR3DS1. Journal of Virology, 2015, 89, 5213-5221.	1.5	67
182	MAIT cells and MR1-antigen recognition. Current Opinion in Immunology, 2017, 46, 66-74.	2.4	67
183	Structural and Sequence Comparisons of Quinone Oxidoreductase, ζ-Crystallin, and Glucose and Alcohol Dehydrogenases. Archives of Biochemistry and Biophysics, 1996, 328, 173-183.	1.4	65
184	Multifunctional Role of Tyr 108 in the Catalytic Mechanism of Human Glutathione Transferase P1-1. Crystallographic and Kinetic Studies on the Y108F Mutant Enzymeâ€,‡. Biochemistry, 1997, 36, 6207-6217.	1.2	65
185	Downregulation of MHC Class I Expression by Influenza A and B Viruses. Frontiers in Immunology, 2019, 10, 1158.	2.2	65
186	TCR Bias and Affinity Define Two Compartments of the CD1b–Glycolipid-Specific T Cell Repertoire. Journal of Immunology, 2014, 192, 4054-4060.	0.4	64
187	A multilayered immune system through the lens of unconventional T cells. Nature, 2021, 595, 501-510.	13.7	64
188	TCRα Genes Direct MHC Restriction in the Potent Human T Cell Response to a Class I-Bound Viral Epitope. Journal of Immunology, 2006, 177, 6804-6814.	0.4	63
189	The 2.7ÂÃ Crystal Structure of the Autoinhibited Human c-Fms Kinase Domain. Journal of Molecular Biology, 2007, 367, 839-847.	2.0	63
190	A Viral Immunoevasin Controls Innate Immunity by Targeting the Prototypical Natural Killer Cell Receptor Family. Cell, 2017, 169, 58-71.e14.	13.5	63
191	Circulating gluten-specific FOXP3 + CD39 + regulatory T cells have impaired suppressive function in patients with celiac disease. Journal of Allergy and Clinical Immunology, 2017, 140, 1592-1603.e8.	1.5	63
192	Single-Cell Approach to Influenza-Specific CD8+ T Cell Receptor Repertoires Across Different Age Groups, Tissues, and Following Influenza Virus Infection. Frontiers in Immunology, 2018, 9, 1453.	2.2	63
193	Public T cell receptors confer high-avidity CD4 responses to HIV controllers. Journal of Clinical Investigation, 2016, 126, 2093-2108.	3.9	63
194	The Structure of the Bacterial Oxidoreductase Enzyme DsbA in Complex with a Peptide Reveals a Basis for Substrate Specificity in the Catalytic Cycle of DsbA Enzymes. Journal of Biological Chemistry, 2009, 284, 17835-17845.	1.6	62
195	Post-translationally modified T cell epitopes: immune recognition and immunotherapy. Journal of Molecular Medicine, 2009, 87, 1045-51.	1.7	62
196	Protective Efficacy of Cross-Reactive CD8+ T Cells Recognising Mutant Viral Epitopes Depends on Peptide-MHC-I Structural Interactions and T Cell Activation Threshold. PLoS Pathogens, 2010, 6, e1001039.	2.1	62
197	NKT TCR Recognition of CD1d-α- <i>C</i> -Galactosylceramide. Journal of Immunology, 2011, 187, 4705-4713.	0.4	62
198	Functional and Structural Characteristics of NY-ESO-1-related HLA A2-restricted Epitopes and the Design of a Novel Immunogenic Analogue. Journal of Biological Chemistry, 2004, 279, 23438-23446.	1.6	61

#	Article	IF	CITATIONS
199	Divergent T-cell receptor recognition modes of a HLA-I restricted extended tumour-associated peptide. Nature Communications, 2018, 9, 1026.	5.8	61
200	Structural Studies on HLA-G: Implications for Ligand and Receptor Binding. Human Immunology, 2007, 68, 220-226.	1.2	60
201	Crystal Structures of the Lyn Protein Tyrosine Kinase Domain in Its Apo- and Inhibitor-bound State. Journal of Biological Chemistry, 2009, 284, 284-291.	1.6	60
202	An overview on the identification of <scp>MAIT</scp> cell antigens. Immunology and Cell Biology, 2018, 96, 573-587.	1.0	60
203	T cell receptor recognition of CD1b presenting a mycobacterial glycolipid. Nature Communications, 2016, 7, 13257.	5.8	59
204	A subset of HLA-DP molecules serve as ligands for the natural cytotoxicity receptor NKp44. Nature Immunology, 2019, 20, 1129-1137.	7.0	59
205	Constraints within major histocompatibility complex class I restricted peptides: Presentation and consequences for T-cell recognition. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 5534-5539.	3.3	58
206	Aerolysin—A Paradigm for Membrane Insertion of Beta-Sheet Protein Toxins?. Journal of Structural Biology, 1998, 121, 92-100.	1.3	57
207	HLA Peptide Length Preferences Control CD8+T Cell Responses. Journal of Immunology, 2013, 191, 561-571.	0.4	57
208	T Cell Determinants Incorporating β-Amino Acid Residues Are Protease Resistant and Remain Immunogenic In Vivo. Journal of Immunology, 2005, 175, 3810-3818.	0.4	56
209	Host immunomodulatory lipids created by symbionts from dietary amino acids. Nature, 2021, 600, 302-307.	13.7	56
210	The production, purification and crystallization of a soluble heterodimeric form of a highly selected T-cell receptor in its unliganded and liganded state. Acta Crystallographica Section D: Biological Crystallography, 2002, 58, 2131-2134.	2.5	55
211	Acute emergence and reversion of influenza A virus quasispecies within CD8+ T cell antigenic peptides. Nature Communications, 2013, 4, 2663.	5.8	55
212	HLA and kidney disease: from associations to mechanisms. Nature Reviews Nephrology, 2018, 14, 636-655.	4.1	55
213	A structural perspective on MHC class lb molecules in adaptive immunity. Trends in Immunology, 2006, 27, 413-420.	2.9	54
214	Enzymatic Properties of an Ecto-nucleoside Triphosphate Diphosphohydrolase from Legionella pneumophila. Journal of Biological Chemistry, 2008, 283, 12909-12918.	1.6	54
215	Impact of clonal competition for peptide-MHC complexes on the CD8+ T-cell repertoire selection in a persistent viral infection. Blood, 2008, 111, 4283-4292.	0.6	54
216	Mucosal-Associated Invariant T Cells Augment Immunopathology and Gastritis in Chronic <i>Helicobacter pylori</i> Infection. Journal of Immunology, 2018, 200, 1901-1916.	0.4	54

#	Article	IF	CITATIONS
217	CD4 <sup>+</sup> T cell–mediated HLA class II cross-restriction in HIV controllers. Science Immunology, 2018, 3, .	5.6	54
218	TAP genes and immunity. Current Opinion in Immunology, 2004, 16, 651-659.	2.4	53
219	Crystal Structure of LipL32, the Most Abundant Surface Protein of Pathogenic Leptospira spp Journal of Molecular Biology, 2009, 387, 1229-1238.	2.0	53
220	Canonical T cell receptor docking on peptide–MHC is essential for T cell signaling. Science, 2021, 372, .	6.0	53
221	Structure of the activation domain of the GM-CSF/IL-3/IL-5 receptor common β-chain bound to an antagonist. Blood, 2000, 95, 2491-2498.	0.6	52
222	Disparate thermodynamics governing T cell receptor-MHC-I interactions implicate extrinsic factors in guiding MHC restriction. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 6641-6646.	3.3	52
223	The molecular bases of δ/αβ T cell–mediated antigen recognition. Journal of Experimental Medicine, 2014, 211, 2599-2615.	4.2	52
224	The Cellular Redox Environment Alters Antigen Presentation. Journal of Biological Chemistry, 2014, 289, 27979-27991.	1.6	52
225	T cell autoreactivity directed toward CD1c itself rather than toward carried self lipids. Nature Immunology, 2018, 19, 397-406.	7.0	52
226	Killer cell immunoglobulin–like receptor 3DL1 variation modifies HLA-B*57 protection against HIV-1. Journal of Clinical Investigation, 2018, 128, 1903-1912.	3.9	52
227	Aerolysin and pertussis toxin share a common receptor-binding domain. EMBO Journal, 1997, 16, 3426-3434.	3.5	51
228	Remarkably low affinity of CD4/peptide-major histocompatibility complex class II protein interactions. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 5682-5687.	3.3	51
229	PD-L1– and calcitriol-dependent liposomal antigen-specific regulation of systemic inflammatory autoimmune disease. JCI Insight, 2019, 4, .	2.3	51
230	Alloreactivity between Disparate Cognate and Allogeneic pMHC-I Complexes Is the Result of Highly Focused, Peptide-dependent Structural Mimicry. Journal of Biological Chemistry, 2006, 281, 34324-34332.	1.6	50
231	Epitope-specific TCRÎ <sup>2</sup> repertoire diversity imparts no functional advantage on the CD8 <sup>+</sup> T cell response to cognate viral peptides. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 2034-2039.	3.3	50
232	Determinants of Gliadin-Specific T Cell Selection in Celiac Disease. Journal of Immunology, 2015, 194, 6112-6122.	0.4	50
233	Absence of mucosal-associated invariant T cells in a person with a homozygous point mutation in <i>MR1</i> . Science Immunology, 2020, 5, .	5.6	50
234	Crystallization, structural determination and analysis of a novel parasite vaccine candidate: Fasciola hepatica glutathione S-transferase 1 1Edited by R. Huber. Journal of Molecular Biology, 1997, 273, 857-872.	2.0	49

#	Article	IF	CITATIONS
235	Affinity Thresholds for Naive CD8+ CTL Activation by Peptides and Engineered Influenza A Viruses. Journal of Immunology, 2011, 187, 5733-5744.	0.4	49
236	Recognition of the nonclassical MHC class I molecule H2-M3 by the receptor Ly49A regulates the licensing and activation of NK cells. Nature Immunology, 2012, 13, 1171-1177.	7.0	49
237	Diverse T Cell Receptor Gene Usage in HLA-DQ8-Associated Celiac Disease Converges into a Consensus Binding Solution. Structure, 2016, 24, 1643-1657.	1.6	49
238	The N terminus of the serpin, tengpin, functions to trap the metastable native state. EMBO Reports, 2007, 8, 658-663.	2.0	48
239	A Long, Naturally Presented Immunodominant Epitope from NY-ESO-1 Tumor Antigen: Implications for Cancer Vaccine Design. Cancer Research, 2009, 69, 1046-1054.	0.4	48
240	A Structural Basis for Antigen Presentation by the MHC Class Ib Molecule, Qa-1b. Journal of Immunology, 2012, 188, 302-310.	0.4	48
241	A Structural Basis for Varied αβ TCR Usage against an Immunodominant EBV Antigen Restricted to a HLA-B8 Molecule. Journal of Immunology, 2012, 188, 311-321.	0.4	48
242	Mutational and Structural Analysis of KIR3DL1 Reveals a Lineage-Defining Allotypic Dimorphism That Impacts Both HLA and Peptide Sensitivity. Journal of Immunology, 2014, 192, 2875-2884.	0.4	48
243	Coeliac disease and rheumatoid arthritis: similar mechanisms, different antigens. Nature Reviews Rheumatology, 2015, 11, 450-461.	3.5	48
244	Broad CD8+ T cell cross-recognition of distinct influenza A strains in humans. Nature Communications, 2018, 9, 5427.	5.8	48
245	Tissue Specificity of Cross-Reactive Allogeneic Responses by EBV EBNA3A-Specific Memory T Cells. Transplantation, 2011, 91, 494-500.	0.5	47
246	A molecular basis for the T cell response in HLA-DQ2.2 mediated celiac disease. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 3063-3073.	3.3	47
247	The impact of HLA-B micropolymorphism outside primary peptide anchor pockets on the CTL response to CMV. European Journal of Immunology, 2007, 37, 946-953.	1.6	46
248	A mortise–tenon joint in the transmembrane domain modulates autotransporter assembly into bacterial outer membranes. Nature Communications, 2014, 5, 4239.	5.8	46
249	MR1 presentation of vitamin B-based metabolite ligands. Current Opinion in Immunology, 2015, 34, 28-34.	2.4	46
250	Adaptability of the semi-invariant natural killer T-cell receptor towards structurally diverse CD1d-restricted ligands. EMBO Journal, 2009, 28, 3579-3590.	3.5	45
251	The 1.5 Ã Crystal Structure of a Prokaryote Serpin. Structure, 2003, 11, 387-397.	1.6	44
252	Specificity on a knife-edge: the αβ T cell receptor. Current Opinion in Structural Biology, 2006, 16, 787-795.	2.6	44

#	Article	IF	CITATIONS
253	T-cell allorecognition: a case of mistaken identity or déjà vu?. Trends in Immunology, 2008, 29, 220-226.	2.9	44
254	The structural bases of direct Tâ€cell allorecognition: implications for Tâ€cellâ€mediated transplant rejection. Immunology and Cell Biology, 2011, 89, 388-395.	1.0	44
255	Understanding the complexity and malleability of Tâ€cell recognition. Immunology and Cell Biology, 2015, 93, 433-441.	1.0	44
256	Rapid screening for the detection of HLAâ€B57 and HLAâ€B58 in prevention of drug hypersensitivity. Tissue Antigens, 2011, 78, 11-20.	1.0	43
257	Structural basis for enabling T-cell receptor diversity within biased virus-specific CD8 <sup>+</sup> T-cell responses. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 9536-9541.	3.3	43
258	Ligand-dependent downregulation of MR1 cell surface expression. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 10465-10475.	3.3	43
259	CD1d antigen presentation: treats for NKT cells. Nature Immunology, 2005, 6, 754-756.	7.0	42
260	The impact of human leukocyte antigen (HLA) micropolymorphism on ligand specificity within the HLA-B*41 allotypic family. Haematologica, 2011, 96, 110-118.	1.7	42
261	Human T cell response to CD1a and contact dermatitis allergens in botanical extracts and commercial skin care products. Science Immunology, 2020, 5, .	5.6	42
262	Studies on the structure and mechanism of a bacterial protein toxin by analytical ultracentrifugation and small-angle neutron scattering 1 1Edited by M. F. Moody. Journal of Molecular Biology, 1999, 293, 1145-1160.	2.0	41
263	X-ray crystal structure of MENT: evidence for functional loop–sheet polymers in chromatin condensation. EMBO Journal, 2006, 25, 3144-3155.	3.5	41
264	Targeting of a natural killer cell receptor family by a viral immunoevasin. Nature Immunology, 2013, 14, 699-705.	7.0	41
265	Structural Basis for CD96 Immune Receptor Recognition of Nectin-like Protein-5, CD155. Structure, 2019, 27, 219-228.e3.	1.6	41
266	The molecular basis underpinning the potency and specificity of MAIT cell antigens. Nature Immunology, 2020, 21, 400-411.	7.0	41
267	Identification of a Potent Microbial Lipid Antigen for Diverse NKT Cells. Journal of Immunology, 2015, 195, 2540-2551.	0.4	40
268	Structural determination of lipid antigens captured at the CD1d–T-cell receptor interface. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 8348-8353.	3.3	40
269	A plasmid-encoded peptide from Staphylococcus aureus induces anti-myeloperoxidase nephritogenic autoimmunity. Nature Communications, 2019, 10, 3392.	5.8	40
270	Crystal Structure of a Legionella pneumophila Ecto -Triphosphate Diphosphohydrolase, A Structural and Functional Homolog of the Eukaryotic NTPDases. Structure, 2010, 18, 228-238.	1.6	39

#	Article	IF	CITATIONS
271	A molecular basis of human T cell receptor autoreactivity toward self-phospholipids. Science Immunology, 2017, 2, .	5.6	39
272	The Structure of the Atypical Killer Cell Immunoglobulin-like Receptor, KIR2DL4. Journal of Biological Chemistry, 2015, 290, 10460-10471.	1.6	38
273	Towards identification of immune and genetic correlates of severe influenza disease in Indigenous Australians. Immunology and Cell Biology, 2016, 94, 367-377.	1.0	38
274	Discriminative T-cell receptor recognition of highly homologous HLA-DQ2–bound gluten epitopes. Journal of Biological Chemistry, 2019, 294, 941-952.	1.6	38
275	Site-directed mutagenesis of theProteus mirabilisglutathione transferase B1-1 G-site. FEBS Letters, 1998, 423, 122-124.	1.3	37
276	Movement of a Loop in Domain 3 of Aerolysin Is Required for Channel Formationâ€. Biochemistry, 1998, 37, 741-746.	1.2	37
277	Mucosa-Associated Invariant T Cells Are Systemically Depleted in Simian Immunodeficiency Virus-Infected Rhesus Macaques. Journal of Virology, 2016, 90, 4520-4529.	1.5	37
278	Anthem: a user customised tool for fast and accurate prediction of binding between peptides and HLA class I molecules. Briefings in Bioinformatics, 2021, 22, .	3.2	37
279	T cell allorecognition and MHC restriction—A case of Jekyll and Hyde?. Molecular Immunology, 2008, 45, 583-598.	1.0	36
280	Structure and Function of the Oxidoreductase DsbA1 from Neisseria meningitidis. Journal of Molecular Biology, 2009, 394, 931-943.	2.0	36
281	Vβ2 natural killer T cell antigen receptor-mediated recognition of CD1d-glycolipid antigen. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 19007-19012.	3.3	36
282	The Impact of a Large and Frequent Deletion in the Human TCR Î <sup>2</sup> Locus on Antiviral Immunity. Journal of Immunology, 2012, 188, 2742-2748.	0.4	36
283	Highly Divergent T-cell Receptor Binding Modes Underlie Specific Recognition of a Bulged Viral Peptide bound to a Human Leukocyte Antigen Class I Molecule. Journal of Biological Chemistry, 2013, 288, 15442-15454.	1.6	36
284	Antigen Specificity of Type I NKT Cells Is Governed by TCR β-Chain Diversity. Journal of Immunology, 2015, 195, 4604-4614.	0.4	36
285	Germline bias dictates cross-serotype reactivity in a common dengue-virus-specific CD8+ T cell response. Nature Immunology, 2017, 18, 1228-1237.	7.0	36
286	Molecular basis for increased susceptibility of Indigenous North Americans to seropositive rheumatoid arthritis. Annals of the Rheumatic Diseases, 2017, 76, 1915-1923.	0.5	36
287	Endoplasmic reticulum chaperones stabilize ligand-receptive MR1 molecules for efficient presentation of metabolite antigens. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 24974-24985.	3.3	36
288	Human Mucosal-Associated Invariant T Cells in Older Individuals Display Expanded TCRαβ Clonotypes with Potent Antimicrobial Responses. Journal of Immunology, 2020, 204, 1119-1133.	0.4	36

#	Article	IF	CITATIONS
289	Complete modification of TCR specificity and repertoire selection does not perturb a CD8 <sup>+</sup> T cell immunodominance hierarchy. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 19408-19413.	3.3	35
290	Virus-Mediated Suppression of the Antigen Presentation Molecule MR1. Cell Reports, 2020, 30, 2948-2962.e4.	2.9	35
291	A natural product compound inhibits coronaviral replication inÂvitro by binding to the conserved Nsp9 SARS-CoV-2 protein. Journal of Biological Chemistry, 2021, 297, 101362.	1.6	35
292	The peptide length specificity of some HLA class I alleles is very broad and includes peptides of up to 25 amino acids in length. Molecular Immunology, 2009, 46, 1911-1917.	1.0	34
293	Human and Mouse Type I Natural Killer T Cell Antigen Receptors Exhibit Different Fine Specificities for CD1d-Antigen Complex. Journal of Biological Chemistry, 2012, 287, 39139-39148.	1.6	34
294	Effective functional maturation of invariant natural killer T cells is constrained by negative selection and T-cell antigen receptor affinity. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E119-28.	3.3	34
295	Atypical natural killer T-cell receptor recognition of CD1d–lipid antigens. Nature Communications, 2016, 7, 10570.	5.8	34
296	The molecular basis for peptide repertoire selection in the human leukocyte antigen (HLA) C*06:02 molecule. Journal of Biological Chemistry, 2017, 292, 17203-17215.	1.6	34
297	Human Î <sup>3</sup> δT cells recognize CD1b by two distinct mechanisms. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 22944-22952.	3.3	34
298	Immune cellular networks underlying recovery from influenza virus infection in acute hospitalized patients. Nature Communications, 2021, 12, 2691.	5.8	34
299	T-cells behaving badly: structural insights into alloreactivity and autoimmunity. Current Opinion in Immunology, 2008, 20, 575-580.	2.4	33
300	A structurally derived consensus pattern for theta class glutathione transferases. Protein Engineering, Design and Selection, 1996, 9, 327-332.	1.0	32
301	Identification of a dominant self-ligand bound to three HLA B44 alleles and the preliminary crystallographic analysis of recombinant forms of each complex. FEBS Letters, 2002, 527, 27-32.	1.3	32
302	The Structure of H-2Kb and Kbm8 Complexed to a Herpes Simplex Virus Determinant: Evidence for a Conformational Switch That Governs T Cell Repertoire Selection and Viral Resistance. Journal of Immunology, 2004, 173, 402-409.	0.4	31
303	HIV-1–Specific CD8 T Cells Exhibit Limited Cross-Reactivity during Acute Infection. Journal of Immunology, 2016, 196, 3276-3286.	0.4	31
304	The Diverse Family of MR1-Restricted T Cells. Journal of Immunology, 2018, 201, 2862-2871.	0.4	31
305	HLA-B57 micropolymorphism defines the sequence and conformational breadth of the immunopeptidome. Nature Communications, 2018, 9, 4693.	5.8	31
306	A T-cell receptor escape channel allows broad T-cell response to CD1b and membrane phospholipids. Nature Communications, 2019, 10, 56.	5.8	31

#	Article	IF	CITATIONS
307	Human skin is colonized by T cells that recognize CD1a independently of lipid. Journal of Clinical Investigation, 2021, 131, .	3.9	31
308	Antagonism of Antiviral and Allogeneic Activity of a Human Public CTL Clonotype by a Single Altered Peptide Ligand: Implications for Allograft Rejection. Journal of Immunology, 2005, 174, 5593-5601.	0.4	30
309	Tetrahydrolipstatin Inhibition, Functional Analyses, and Three-dimensional Structure of a Lipase Essential for Mycobacterial Viability. Journal of Biological Chemistry, 2010, 285, 30050-30060.	1.6	30
310	The High Resolution Crystal Structure of a Native Thermostable Serpin Reveals the Complex Mechanism Underpinning the Stressed to Relaxed Transition. Journal of Biological Chemistry, 2005, 280, 8435-8442.	1.6	29
311	Tollâ€like receptors: structural pieces of a curveâ€shaped puzzle. Immunology and Cell Biology, 2007, 85, 406-410.	1.0	29
312	Loss of Anti-Viral Immunity by Infection with a Virus Encoding a Cross-Reactive Pathogenic Epitope. PLoS Pathogens, 2012, 8, e1002633.	2.1	29
313	A Structural Basis for the pH-dependent Increase in Fluorescence Efficiency of Chromoproteins. Journal of Molecular Biology, 2007, 368, 998-1010.	2.0	28
314	The Energetic Basis Underpinning T-cell Receptor Recognition of a Super-bulged Peptide Bound to a Major Histocompatibility Complex Class I Molecule. Journal of Biological Chemistry, 2012, 287, 12267-12276.	1.6	28
315	SARSâ€CoVâ€2â€specific CD8 <sup>+</sup> Tâ€cell responses and TCR signatures in the context of a prominent HLAâ€A*24:02 allomorph. Immunology and Cell Biology, 2021, 99, 990-1000.	1.0	28
316	Francisella tularensis induces Th1 like MAIT cells conferring protection against systemic and local infection. Nature Communications, 2021, 12, 4355.	5.8	28
317	Targeted suppression of autoreactive CD8+ T-cell activation using blocking anti-CD8 antibodies. Scientific Reports, 2016, 6, 35332.	1.6	27
318	Dual Modifications of α-Galactosylceramide Synergize to Promote Activation of Human Invariant Natural Killer T Cells and Stimulate Anti-tumor Immunity. Cell Chemical Biology, 2018, 25, 571-584.e8.	2.5	27
319	Peptide mimic for influenza vaccination using nonnatural combinatorial chemistry. Journal of Clinical Investigation, 2018, 128, 1569-1580.	3.9	27
320	Antigen-Driven Patterns of TCR Bias Are Shared across Diverse Outcomes of Human Hepatitis C Virus Infection. Journal of Immunology, 2011, 186, 901-912.	0.4	26
321	Endogenous antigen presentation impacts on T-box transcription factor expression and functional maturation of CD8+ T cells. Blood, 2012, 120, 3237-3245.	0.6	25
322	The Interaction of KIR3DL1*001 with HLA Class I Molecules Is Dependent upon Molecular Microarchitecture within the Bw4 Epitope. Journal of Immunology, 2015, 194, 781-789.	0.4	25
323	A hot spot on interferon α/β receptor subunit 1 (IFNAR1) underpins its interaction with interferon-β and dictates signaling. Journal of Biological Chemistry, 2017, 292, 7554-7565.	1.6	25
324	A Biophysical Analysis of the Tetratricopeptide Repeat-rich Mitochondrial Import Receptor, Tom70, Reveals an Elongated Monomer That Is Inherently Flexible, Unstable, and Unfolds via a Multistate Pathway. Journal of Biological Chemistry, 2004, 279, 46448-46454.	1.6	24

#	Article	IF	CITATIONS
325	<scp>CD</scp> 1d protein structure determines speciesâ€selective antigenicity of isoglobotrihexosylceramide (i <scp>G</scp> b3) to invariant <scp>NKT</scp> cells. European Journal of Immunology, 2013, 43, 815-825.	1.6	24
326	Mucosalâ€associated invariant T cell receptor recognition of small molecules presented by <scp>MR</scp> 1. Immunology and Cell Biology, 2018, 96, 588-597.	1.0	24
327	Growth Hormone Stops Excessive Inflammation After Partial Hepatectomy, Allowing Liver Regeneration and Survival Through Induction of H2â€Bl/HLAâ€G. Hepatology, 2021, 73, 759-775.	3.6	24
328	CD1a selectively captures endogenous cellular lipids that broadly block T cell response. Journal of Experimental Medicine, 2021, 218, .	4.2	24
329	Hijacking of a Substrate-binding Protein Scaffold for use in Mycobacterial Cell Wall Biosynthesis. Journal of Molecular Biology, 2006, 359, 983-997.	2.0	23
330	Crystal Structure of a UDP-glucose-specific Glycosyltransferase from a Mycobacterium Species. Journal of Biological Chemistry, 2008, 283, 27881-27890.	1.6	23
331	Structural and Biological Basis of CTL Escape in Coronavirus-Infected Mice. Journal of Immunology, 2008, 180, 3926-3937.	0.4	23
332	Structural and Biochemical Characterization of the Oxidoreductase NmDsbA3 from Neisseria meningitidis. Journal of Biological Chemistry, 2008, 283, 32452-32461.	1.6	23
333	Lack of Heterologous Cross-reactivity toward HLA-A*02:01 Restricted Viral Epitopes Is Underpinned by Distinct αβT Cell Receptor Signatures. Journal of Biological Chemistry, 2016, 291, 24335-24351.	1.6	23
334	Structure–function analyses of a pertussis-like toxin from pathogenic Escherichia coli reveal a distinct mechanism of inhibition of trimeric G-proteins. Journal of Biological Chemistry, 2017, 292, 15143-15158.	1.6	23
335	Identification of Native and Posttranslationally Modified HLAâ€B*57:01â€Restricted HIV Envelope Derived Epitopes Using Immunoproteomics. Proteomics, 2018, 18, e1700253.	1.3	23
336	SARS-CoV-2-specific TÂcell memory with common TCRαβ motifs is established in unvaccinated children who seroconvert after infection. Immunity, 2022, 55, 1299-1315.e4.	6.6	23
337	Shifting Substrate Specificity of Human Glutathione Transferase (from Class Pi to Class Alpha) by a Single Point Mutation. Biochemical and Biophysical Research Communications, 1998, 252, 184-189.	1.0	22
338	EcxAB Is a Founding Member of a New Family of Metalloprotease AB5 Toxins with a Hybrid Cholera-like B Subunit. Structure, 2013, 21, 2003-2013.	1.6	22
339	Lipids hide or step aside for CD1-autoreactive T cell receptors. Current Opinion in Immunology, 2018, 52, 93-99.	2.4	22
340	T cell receptor recognition of hybrid insulin peptides bound to HLA-DQ8. Nature Communications, 2021, 12, 5110.	5.8	22
341	Recognition of the antigen-presenting molecule MR1 by a Vδ3 <sup>+</sup> γδT cell receptor. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	22
342	Structural Basis of Subtilase Cytotoxin SubAB Assembly. Journal of Biological Chemistry, 2013, 288, 27505-27516.	1.6	21

#	Article	IF	CITATIONS
343	An Extensive Antigenic Footprint Underpins Immunodominant TCR Adaptability against a Hypervariable Viral Determinant. Journal of Immunology, 2014, 193, 5402-5413.	0.4	21
344	Differing roles of CD1d2 and CD1d1 proteins in type I natural killer T cell development and function. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E1204-E1213.	3.3	21
345	The early proximal $\hat{I}\pm\hat{I}^2$ TCR signalosome specifies thymic selection outcome through a quantitative protein interaction network. Science Immunology, 2019, 4, .	5.6	21
346	A single-domain bispecific antibody targeting CD1d and the NKT T-cell receptor induces a potent antitumor response. Nature Cancer, 2020, 1, 1054-1065.	5.7	21
347	Structural plasticity of KIR2DL2 and KIR2DL3 enables altered docking geometries atop HLA-C. Nature Communications, 2021, 12, 2173.	5.8	21
348	Evaluation of the role of two conserved active-site residues in Beta class glutathione S-transferases. Biochemical Journal, 2000, 351, 341-346.	1.7	20
349	Identification of the Structural and Functional Domains of the Large Serine Recombinase TnpX from Clostridium perfringens. Journal of Biological Chemistry, 2005, 280, 2503-2511.	1.6	20
350	The 2.1Ã Crystal Structure of copGFP, a Representative Member of the Copepod Clade Within the Green Fluorescent Protein Superfamily. Journal of Molecular Biology, 2006, 359, 890-900.	2.0	20
351	A Molecular Basis for the Interplay between T Cells, Viral Mutants, and Human Leukocyte Antigen Micropolymorphism. Journal of Biological Chemistry, 2014, 289, 16688-16698.	1.6	20
352	The versatility of the $\hat{I} \pm \hat{I}^2$ Tâ $\in c$ ell antigen receptor. Protein Science, 2014, 23, 260-272.	3.1	20
353	A T Cell Receptor Locus Harbors a Malaria-Specific Immune Response Gene. Immunity, 2017, 47, 835-847.e4.	6.6	20
354	A conserved energetic footprint underpins recognition of human leukocyte antigen-E by two distinct αβ T cell receptors. Journal of Biological Chemistry, 2017, 292, 21149-21158.	1.6	20
355	Recognition of host Clr-b by the inhibitory NKR-P1B receptor provides a basis for missing-self recognition. Nature Communications, 2018, 9, 4623.	5.8	20
356	Inability To Detect Cross-Reactive Memory T Cells Challenges the Frequency of Heterologous Immunity among Common Viruses. Journal of Immunology, 2018, 200, 3993-4003.	0.4	20
357	CD8+ T cell landscape in Indigenous and non-Indigenous people restricted by influenza mortality-associated HLA-A*24:02 allomorph. Nature Communications, 2021, 12, 2931.	5.8	20
358	Structural basis of biased T cell receptor recognition of an immunodominant HLA-A2 epitope of the SARS-CoV-2 spike protein. Journal of Biological Chemistry, 2021, 297, 101065.	1.6	20
359	The production, purification and crystallization of a pocilloporin pigment from a reef-forming coral. Acta Crystallographica Section D: Biological Crystallography, 2003, 59, 597-599.	2.5	19
360	Engineering a New C-terminal Tail in the H-site of Human Glutathione Transferase P1-1: Structural and Functional Consequences. Journal of Molecular Biology, 2003, 325, 111-122.	2.0	19

#	Article	IF	CITATIONS
361	Degenerate Recognition of MHC Class I Molecules with Bw4 and Bw6 Motifs by a Killer Cell Ig-like Receptor 3DL Expressed by Macaque NK Cells. Journal of Immunology, 2012, 189, 4338-4348.	0.4	19
362	Molecular Imprint of Exposure to Naturally Occurring Genetic Variants of Human Cytomegalovirus on the T cell Repertoire. Scientific Reports, 2014, 4, 3993.	1.6	19
363	Recognition of the Major Histocompatibility Complex (MHC) Class Ib Molecule H2-Q10 by the Natural Killer Cell Receptor Ly49C. Journal of Biological Chemistry, 2016, 291, 18740-18752.	1.6	19
364	Peripheral Blood Mucosal-Associated Invariant T Cells in Tuberculosis Patients and Healthy Mycobacterium tuberculosis-Exposed Controls. Journal of Infectious Diseases, 2020, 222, 995-1007.	1.9	19
365	Crystallization and preliminary X-ray analysis of a thiol-activated cytolysin. FEBS Letters, 1996, 397, 290-292.	1.3	18
366	The Structure and Stability of the Monomorphic HLA-G Are Influenced by the Nature of the Bound Peptide. Journal of Molecular Biology, 2010, 397, 467-480.	2.0	18
367	Preemptive priming readily overcomes structure-based mechanisms of virus escape. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 5570-5575.	3.3	18
368	Structure of the activation domain of the GM-CSF/IL-3/IL-5 receptor common beta-chain bound to an antagonist. Blood, 2000, 95, 2491-8.	0.6	18
369	Structures of thermolabile mutants of human glutathione transferase P1-1 1 1Edited by R. Huber. Journal of Molecular Biology, 2000, 302, 295-302.	2.0	17
370	Distinct CD1d docking strategies exhibited by diverse Type II NKT cell receptors. Nature Communications, 2019, 10, 5242.	5.8	17
371	A homology model for the human theta-class glutathione transferase T1–1. , 1998, 33, 444-454.		16
372	Determination of chromophore charge states in the low pH color transition of the fluorescent protein Rtms5H146S via time-dependent DFT. Chemical Physics Letters, 2006, 420, 507-511.	1.2	16
373	The production and crystallization of the human leukocyte antigen class II molecules HLA-DQ2 and HLA-DQ8 complexed with deamidated gliadin peptides implicated in coeliac disease. Acta Crystallographica Section F: Structural Biology Communications, 2007, 63, 1021-1025.	0.7	16
374	The Ly49 natural killer cell receptors: a versatile tool for viral selfâ€discrimination. Immunology and Cell Biology, 2014, 92, 214-220.	1.0	16
375	The molecular basis of how buried human leukocyte antigen polymorphism modulates natural killer cell function. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 11636-11647.	3.3	16
376	The molecular assembly of the marsupial γμ T cell receptor defines a third T cell lineage. Science, 2021, 371, 1383-1388.	6.0	16
377	Our evolving understanding of the role of the î³Î´T cell receptor in î³Î´T cell mediated immunity. Biochemical Society Transactions, 2021, 49, 1985-1995.	1.6	16
378	<i>Nfkb2</i> variants reveal a p100-degradation threshold that defines autoimmune susceptibility. Journal of Experimental Medicine, 2021, 218, .	4.2	16

#	Article	IF	CITATIONS
379	Repeated <i>Plasmodium falciparum</i> infection in humans drives the clonal expansion of an adaptive γδT cell repertoire. Science Translational Medicine, 2021, 13, eabe7430.	5.8	16
380	Mutagenic analysis of conserved arginine residues in and around the novel sulfate binding pocket of the human Theta class glutathione transferase T2â $\in 2$ . Protein Science, 1999, 8, 2205-2212.	3.1	15
381	Purification and biological characterization of soluble, recombinant mouse IFNÎ <sup>2</sup> expressed in insect cells. Protein Expression and Purification, 2014, 94, 7-14.	0.6	15
382	The Structure of the Cytomegalovirus-Encoded m04 Glycoprotein, a Prototypical Member of the m02 Family of Immunoevasins. Journal of Biological Chemistry, 2014, 289, 23753-23763.	1.6	15
383	H1N1 hemagglutinin-specific HLA-DQ6-restricted CD4+ T cells can be readily detected in narcolepsy type 1 patients and healthy controls. Journal of Neuroimmunology, 2019, 332, 167-175.	1.1	15
384	A high-affinity human TCR-like antibody detects celiac disease gluten peptide–MHC complexes and inhibits T cell activation. Science Immunology, 2021, 6, .	5.6	15
385	Recent advances in all-protein chromophore technology. Biotechnology Annual Review, 2006, 12, 31-66.	2.1	14
386	How a Home-Grown T Cell Receptor Interacts with a Foreign Landscape. Cell, 2007, 129, 19-20.	13.5	14
387	Prevention of Cytotoxic T Cell Escape Using a Heteroclitic Subdominant Viral T Cell Determinant. PLoS Pathogens, 2008, 4, e1000186.	2.1	14
388	T Cell Cross-Reactivity between a Highly Immunogenic EBV Epitope and a Self-Peptide Naturally Presented by HLA-B*18:01+ Cells. Journal of Immunology, 2015, 194, 4668-4675.	0.4	14
389	Disrupting the Allosteric Interaction between the Plasmodium falciparum cAMP-dependent Kinase and Its Regulatory Subunit. Journal of Biological Chemistry, 2016, 291, 25375-25386.	1.6	14
390	Challenging immunodominance of influenza-specific CD8+ T cell responses restricted by the risk-associated HLA-A*68:01 allomorph. Nature Communications, 2019, 10, 5579.	5.8	14
391	Carbamazepine Induces Focused T Cell Responses in Resolved Stevens-Johnson Syndrome and Toxic Epidermal Necrolysis Cases But Does Not Perturb the Immunopeptidome for T Cell Recognition. Frontiers in Immunology, 2021, 12, 653710.	2.2	14
392	The shared susceptibility epitope of HLA-DR4 binds citrullinated self-antigens and the TCR. Science Immunology, 2021, 6, .	5.6	14
393	Predisposed αβ T cell antigen receptor recognition of MHC and MHC-I like molecules?. Current Opinion in Immunology, 2013, 25, 653-659.	2.4	13
394	Structure of the Chicken CD3ϵδ/γ Heterodimer and Its Assembly with the αβT Cell Receptor. Journal of Biological Chemistry, 2014, 289, 8240-8251.	1.6	13
395	Total Synthesis of <i>Mycobacterium tuberculosis</i> Dideoxymycobactinâ€838 and Stereoisomers: Diverse CD1aâ€Restricted T Cells Display a Common Hierarchy of Lipopeptide Recognition. Chemistry - A European Journal, 2017, 23, 1694-1701.	1.7	13
396	Structural basis for the recognition of nectin-like protein-5 by the human-activating immune receptor, DNAM-1. Journal of Biological Chemistry, 2019, 294, 12534-12546.	1.6	13

#	Article	IF	CITATIONS
397	Atypical TRAV1-2â^' T cell receptor recognition of the antigen-presenting molecule MR1. Journal of Biological Chemistry, 2020, 295, 14445-14457.	1.6	13
398	Transcriptional profiling of human Vδ1 TÂcells reveals a pathogen-driven adaptive differentiation program. Cell Reports, 2022, 39, 110858.	2.9	13
399	Two distinct regions of the large serine recombinase TnpX are required for DNA binding and biological function. Molecular Microbiology, 2006, 60, 591-601.	1.2	12
400	Crystal Structure and Comparative Functional Analyses of a Mycobacterium Aldo-Keto Reductase. Journal of Molecular Biology, 2010, 398, 26-39.	2.0	12
401	Identification of Self-antigen–specific T Cells Reflecting Loss of Tolerance in Autoimmune Disease Underpins Preventative Immunotherapeutic Strategies in Rheumatoid Arthritis. Rheumatic Disease Clinics of North America, 2014, 40, 735-752.	0.8	12
402	Characterization and Purification of Mouse Mucosalâ€Associated Invariant T (MAIT) Cells. Current Protocols in Immunology, 2019, 127, e89.	3.6	12
403	Cross-Reactive Donor-Specific CD8+ Tregs Efficiently Prevent Transplant Rejection. Cell Reports, 2019, 29, 4245-4255.e6.	2.9	12
404	Allelic association with ankylosing spondylitis fails to correlate with human leukocyte antigen B27 homodimer formation. Journal of Biological Chemistry, 2019, 294, 20185-20195.	1.6	12
405	A Shared TCR Bias toward an Immunogenic EBV Epitope Dominates in HLA-B*07:02–Expressing Individuals. Journal of Immunology, 2020, 205, 1524-1534.	0.4	12
406	The Role of the HLA Class I α2 Helix in Determining Ligand Hierarchy for the Killer Cell Ig-like Receptor 3DL1. Journal of Immunology, 2021, 206, 849-860.	0.4	12
407	Atypical sideways recognition of CD1a by autoreactive γδT cell receptors. Nature Communications, 2022, 13, .	5.8	12
408	Tumors reveal their secrets to cytotoxic T cells. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 14649-14650.	3.3	11
409	Flow Cytometric Clinical Immunomonitoring Using Peptide–MHC Class II Tetramers: Optimization of Methods and Protocol Development. Frontiers in Immunology, 2018, 9, 8.	2.2	11
410	Characterization of Human Mucosalâ€associated Invariant T (MAIT) Cells. Current Protocols in Immunology, 2019, 127, e90.	3.6	11
411	α-Glucuronosyl and α-glucosyl diacylglycerides, natural killer T cell-activating lipids from bacteria and fungi. Chemical Science, 2020, 11, 2161-2168.	3.7	11
412	The cryo-EM structure of the endocytic receptor DEC-205. Journal of Biological Chemistry, 2021, 296, 100127.	1.6	11
413	Varicella Zoster Virus Impairs Expression of the Nonclassical Major Histocompatibility Complex Class I–Related Gene Protein (MR1). Journal of Infectious Diseases, 2023, 227, 391-401.	1.9	11
414	HLA-A*11:01-restricted CD8+ T cell immunity against influenza A and influenza B viruses in Indigenous and non-Indigenous people. PLoS Pathogens, 2022, 18, e1010337.	2.1	11

#	Article	IF	CITATIONS
415	Fighting infection with your MAITs. Nature Immunology, 2010, 11, 693-695.	7.0	10
416	Cloning, expression, purification and crystallographic studies of galectin-11 from domestic sheep (Ovis aries). Acta Crystallographica Section F, Structural Biology Communications, 2015, 71, 993-997.	0.4	10
417	A TCR β-Chain Motif Biases toward Recognition of Human CD1 Proteins. Journal of Immunology, 2019, 203, 3395-3406.	0.4	10
418	A microfluidic-SERSplatform for isolation and immuno-phenotyping of antigen specific T-cells. Sensors and Actuators B: Chemical, 2019, 284, 281-288.	4.0	10
419	Binding of a pyrimidine RNA base-mimic to SARS-CoV-2 nonstructural protein 9. Journal of Biological Chemistry, 2021, 297, 101018.	1.6	10
420	Binding Studies of the Prodrug HAO472 to SARS-Cov-2 Nsp9 and Variants. ACS Omega, 2022, 7, 7327-7332.	1.6	10
421	Amino acid substitutions around the chromophore of the chromoprotein Rtms5 influence polypeptide cleavage. Biochemical and Biophysical Research Communications, 2006, 340, 1139-1143.	1.0	9
422	Tracking the Unfolding Pathway of a Multirepeat Protein via Tryptophan Scanning. Journal of Biological Chemistry, 2006, 281, 24345-24350.	1.6	9
423	Structural and Functional Correlates of Enhanced Antiviral Immunity Generated by Heteroclitic CD8 T Cell Epitopes. Journal of Immunology, 2014, 192, 5245-5256.	0.4	9
424	Overlapping Peptides Elicit Distinct CD8+ T Cell Responses following Influenza A Virus Infection. Journal of Immunology, 2020, 205, 1731-1742.	0.4	9
425	Mucosal-Associated Invariant T Cell Effector Function Is an Intrinsic Cell Property That Can Be Augmented by the Metabolic Cofactor α-Ketoglutarate. Journal of Immunology, 2021, 206, 1425-1435.	0.4	9
426	Structural Insight into Natural Killer T Cell Receptor Recognition of CD1d. Advances in Experimental Medicine and Biology, 2007, 598, 20-34.	0.8	9
427	Evaluation of the role of two conserved active-site residues in beta class glutathione S-transferases. Biochemical Journal, 2000, 351 Pt 2, 341-6.	1.7	9
428	Evaluation of the role of two conserved active-site residues in Beta class glutathione S-transferases. Biochemical Journal, 2000, 351, 341.	1.7	8
429	The production, purification and crystallization of a soluble form of the nonclassical MHC HLA-G: the essential role of cobalt. Acta Crystallographica Section F: Structural Biology Communications, 2006, 62, 70-73.	0.7	8
430	The 2.0 Ã Crystal Structure of a Pocilloporin at pH 3.5: The Structural Basis for the Linkage Between Color Transition and Halide Binding. Photochemistry and Photobiology, 2006, 82, 359.	1.3	8
431	αβ T Cell Receptors Come Out Swinging. Immunity, 2011, 35, 660-662.	6.6	8
432	Benzofuran sulfonates and small self-lipid antigens activate type II NKT cells via CD1d. Proceedings of the United States of America, 2021, 118, .	3.3	8

#	Article	IF	CITATIONS
433	Expression, purification, crystallization and preliminary X-ray analysis of eCGP123, an extremely stable monomeric green fluorescent protein with reversible photoswitching properties. Acta Crystallographica Section F: Structural Biology Communications, 2011, 67, 1266-1268.	0.7	7
434	A Green Fluorescent Protein Containing a QFG Tri-Peptide Chromophore: Optical Properties and X-Ray Crystal Structure. PLoS ONE, 2012, 7, e47331.	1.1	7
435	Crystal structure of fuculose aldolase from the Antarctic psychrophilic yeast <i>Glaciozyma antarctica</i> PI12. Acta Crystallographica Section F, Structural Biology Communications, 2016, 72, 831-839.	0.4	7
436	A biased view toward celiac disease. Mucosal Immunology, 2016, 9, 583-586.	2.7	7
437	Cytomegalovirus replication is associated with enrichment of distinct γδT cell subsets following lung transplantation: A novel therapeutic approach?. Journal of Heart and Lung Transplantation, 2020, 39, 1300-1312.	0.3	7
438	Preferential HLA-B27 Allorecognition Displayed by Multiple Cross-Reactive Antiviral CD8+ T Cell Receptors. Frontiers in Immunology, 2020, 11, 248.	2.2	7
439	Effects of a local auxiliary protein on the two-dimensional affinity of a TCR-peptide MHC interaction. Journal of Cell Science, 2020, 133, .	1.2	7
440	Structural basis of T cell receptor specificity and cross-reactivity of two HLA-DQ2.5-restricted gluten epitopes in celiac disease. Journal of Biological Chemistry, 2022, 298, 101619.	1.6	7
441	Collision-Induced Affinity Selection Mass Spectrometry for Identification of Ligands. ACS Bio & Med Chem Au, 2022, 2, 450-455.	1.7	7
442	Structural Studies on the $\hat{I}\pm\hat{I}^2$ T-cell Receptor. IUBMB Life, 2005, 57, 575-582.	1.5	6
443	Expression, purification, crystallization and preliminary X-ray characterization of a putative glycosyltransferase of the GT-A fold found in mycobacteria. Acta Crystallographica Section F: Structural Biology Communications, 2008, 64, 428-431.	0.7	6
444	Beta-testing NKT cell self-reactivity. Nature Immunology, 2011, 12, 1135-1137.	7.0	6
445	Cloning, expression, purification and preliminary X-ray diffraction studies of a novel AB5toxin. Acta Crystallographica Section F: Structural Biology Communications, 2013, 69, 912-915.	0.7	6
446	Structural bases of T cell antigen receptor recognition in celiac disease. Current Opinion in Structural Biology, 2022, 74, 102349.	2.6	6
447	HLA-B*27:05 alters immunodominance hierarchy of universal influenza-specific CD8+ T cells. PLoS Pathogens, 2020, 16, e1008714.	2.1	5
448	Rational design of a hydrolysis-resistant mycobacterial phosphoglycolipid antigen presented by CD1c to T cells. Journal of Biological Chemistry, 2021, 297, 101197.	1.6	5
449	Crystal structures of pertussis toxin with NAD+ and analogs provide structural insights into the mechanism of its cytosolic ADP-ribosylation activity. Journal of Biological Chemistry, 2022, 298, 101892.	1.6	5
450	Preliminary X-ray crystallographic studies of a newly defined human theta-class glutathione transferase. Acta Crystallographica Section D: Biological Crystallography, 1998, 54, 148-150.	2.5	4

#	Article	IF	CITATIONS
451	Dichloromethane mediatedin vivoselection and functional characterization of rat glutathioneS-transferase theta 1-1 variants. FEBS Journal, 2001, 268, 4001-4010.	0.2	4
452	The production and purification of the human T-cell receptors, the CD3â^ŠÎ³ and CD3â^ŠÎ′ heterodimers: complex formation and crystallization with OKT3, a therapeutic monoclonal antibody. Acta Crystallographica Section D: Biological Crystallography, 2004, 60, 1425-1428.	2.5	4
453	Expression, purification, crystallization and preliminary X-ray diffraction analysis of an essential lipoprotein implicated in cell-wall biosynthesis inMycobacteria. Acta Crystallographica Section F: Structural Biology Communications, 2005, 61, 1081-1083.	0.7	4
454	The oligomeric assembly of galectin-11 is critical for anti-parasitic activity in sheep (Ovis aries). Communications Biology, 2020, 3, 464.	2.0	4
455	Comparative three-dimensional structure of cholesterol-dependent cytolysins. , 2006, , 659-670.		4
456	Crystallization and preliminary X-ray analysis of a bacterial glutathione transferase. Acta Crystallographica Section D: Biological Crystallography, 1996, 52, 189-191.	2.5	3
457	Valine 10 May Act as a Driver for Product Release from the Active Site of Human Glutathione Transferase P1-1â€,‡. Biochemistry, 2000, 39, 15961-15970.	1.2	3
458	Managing and mining protein crystallization data. Proteins: Structure, Function and Bioinformatics, 2005, 62, 4-7.	1.5	3
459	Human Leukocyte Antigen Class I-Restricted Activation of CD8+ T Cells Provides the Immunogenetic Basis of a Systemic Drug Hypersensitivity. Immunity, 2008, 29, 165.	6.6	3
460	Insight into the basis of autonomous immunoreceptor activation. Trends in Immunology, 2011, 32, 165-170.	2.9	3
461	NKT cells: the smoking gun in fungal-induced asthma?. Nature Medicine, 2013, 19, 1210-1211.	15.2	3
462	A structural characterization of the isoniazidMycobacterium tuberculosisdrug target, Rv2971, in its unliganded form. Acta Crystallographica Section F, Structural Biology Communications, 2014, 70, 572-577.	0.4	3
463	A structural and functional investigation of a novel protein fromMycobacterium smegmatisimplicated in mycobacterial macrophage survivability. Acta Crystallographica Section D: Biological Crystallography, 2014, 70, 2264-2276.	2.5	3
464	Tetramer Immunization and Selection Followed by CELLISA Screening to Generate Monoclonal Antibodies against the Mouse Cytomegalovirus m12 Immunoevasin. Journal of Immunology, 2020, 205, 1709-1717.	0.4	3
465	Elucidating the Motif for CpG Oligonucleotide Binding to the Dendritic Cell Receptor DEC-205 Leads to Improved Adjuvants for Liver-Resident Memory. Journal of Immunology, 2021, 207, 1836-1847.	0.4	3
466	Evaluation of a fit-for-purpose assay to monitor antigen-specific functional CD4+ T-cell subpopulations in rheumatoid arthritis using flow cytometry–based peptide-MHC class-II tetramer staining. Clinical and Experimental Immunology, 2022, 207, 72-83.	1.1	3
467	Overcoming the LAG3 phase problem. Nature Immunology, 2022, 23, 993-995.	7.0	3
468	Dual TCR-α Expression on Mucosal-Associated Invariant T Cells as a Potential Confounder of TCR Interpretation. Journal of Immunology, 2022, 208, 1389-1395.	0.4	2

#	Article	IF	CITATIONS
469	Alloreactivity. Methods in Molecular Biology, 2013, 1034, 3-39.	0.4	1
470	Cloning, expression, purification and preliminary X-ray diffraction studies of a mycobacterial protein implicated in bacterial survival in macrophages. Acta Crystallographica Section F: Structural Biology Communications, 2013, 69, 566-569.	0.7	1
471	Cutting Edge: Unconventional CD8 <sup>+</sup> T Cell Recognition of a Naturally Occurring HLA-A*02:01–Restricted 20mer Epitope. Journal of Immunology, 2022, , ji2101208.	0.4	1
472	ORGAN SPECIFICITY OF CROSS-REACTIVE ALLOGENEIC RESPONSES BY VIRAL SPECIFIC MEMORY T-CELLS. Transplantation, 2010, 90, 239.	0.5	0
473	How opposites attract. Immunology and Cell Biology, 2011, 89, 163-164.	1.0	Ο
474	Immune self-reactivity triggered by drug-modified human leukocyte antigen-peptide presentation. Molecular Immunology, 2012, 51, 19.	1.0	0
475	A structural basis for selection of the mucosal associated invariant T cell receptor in MR1-restricted antigen recognition. Molecular Immunology, 2012, 51, 25.	1.0	Ο
476	Shedding light on MHC class I antigen loading: a UV-labile peptide ligand approach. Molecular Immunology, 2012, 51, 33.	1.0	0
477	Missense single nucleotide polymorphisms in the human T cell receptor loci control variable gene usage in the T cell repertoire. British Journal of Haematology, 2014, 166, 148-152.	1.2	Ο
478	OR26 BW4 + HLA class I and KIR3DL1 allotypic pairing influences licensing and effector function of KIR3DL1 + NK cells. Human Immunology, 2017, 78, 25.	1.2	0
479	Editorial. Seminars in Cell and Developmental Biology, 2018, 84, 1.	2.3	Ο
480	Reply to Roudier et al.: HLA-DRB1 polymorphism, anti-citrullinated protein antibodies, and rheumatoid arthritis. Journal of Biological Chemistry, 2018, 293, 7039.	1.6	0
481	Reply:. Hepatology, 2021, 73, 1239-1239.	3.6	Ο
482	Katharina Gaus 1972–2021. Nature Immunology, 2021, 22, 535-536.	7.0	0
483	The human pi-class glutathione S-transferase: a target for structure-based drug design. Acta Crystallographica Section A: Foundations and Advances, 1996, 52, C112-C112.	0.3	Ο
484	HLA-B*27:05 alters immunodominance hierarchy of universal influenza-specific CD8+ T cells. , 2020, 16, e1008714.		0
485	HLA-B*27:05 alters immunodominance hierarchy of universal influenza-specific CD8+ T cells. , 2020, 16, e1008714.		0
486	HLA-B*27:05 alters immunodominance hierarchy of universal influenza-specific CD8+ T cells. , 2020, 16,		0

e1008714.

#	Article	IF	CITATIONS
487	HLA-B*27:05 alters immunodominance hierarchy of universal influenza-specific CD8+ T cells. , 2020, 16, e1008714.		0