

# Paul J Lehner

## List of Publications by Year in descending order

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153  
papers

14,943  
citations

13099

68  
h-index

20961

115  
g-index

183  
all docs

183  
docs citations

183  
times ranked

19047  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome surveillance by HUSH-mediated silencing of intronless mobile elements. <i>Nature</i> , 2022, 601, 440-445.	27.8	64
2	Topical TMPRSS2 inhibition prevents SARS-CoV-2 infection in differentiated human airway cultures. <i>Life Science Alliance</i> , 2022, 5, e202101116.	2.8	10
3	Unbiased cell surface proteomics identifies SEMA4A as an effective immunotherapy target for myeloma. <i>Blood</i> , 2022, 139, 2471-2482.	1.4	12
4	SARS-CoV-2 host-shutoff impacts innate NK cell functions, but antibody-dependent NK activity is strongly activated through non-spike antibodies. <i>ELife</i> , 2022, 11, .	6.0	34
5	Dissecting Herpes Simplex Virus 1-Induced Host Shutoff at the RNA Level. <i>Journal of Virology</i> , 2021, 95, .	3.4	25
6	Single-dose BNT162b2 vaccine protects against asymptomatic SARS-CoV-2 infection. <i>ELife</i> , 2021, 10, .	6.0	57
7	Neural stem cells traffic functional mitochondria via extracellular vesicles. <i>PLoS Biology</i> , 2021, 19, e3001166.	5.6	95
8	The SMC5/6 complex compacts and silences unintegrated HIV-1 DNA and is antagonized by Vpr. <i>Cell Host and Microbe</i> , 2021, 29, 792-805.e6.	11.0	49
9	Longitudinal analysis reveals that delayed bystander CD8+ T cell activation and early immune pathology distinguish severe COVID-19 from mild disease. <i>Immunity</i> , 2021, 54, 1257-1275.e8.	14.3	230
10	Human embryonic stem cell-derived cardiomyocyte platform screens inhibitors of SARS-CoV-2 infection. <i>Communications Biology</i> , 2021, 4, 926.	4.4	11
11	CD97 stabilises the immunological synapse between dendritic cells and T cells and is targeted for degradation by the Salmonella effector SteD. <i>PLoS Pathogens</i> , 2021, 17, e1009771.	4.7	17
12	A prenylated dsRNA sensor protects against severe COVID-19. <i>Science</i> , 2021, 374, eabj3624.	12.6	124
13	No evidence for basigin/CD147 as a direct SARS-CoV-2 spike binding receptor. <i>Scientific Reports</i> , 2021, 11, 413.	3.3	156
14	Periphilin self-association underpins epigenetic silencing by the HUSH complex. <i>Nucleic Acids Research</i> , 2020, 48, 10313-10328.	14.5	15
15	Quantitative Proteomics Analysis of Lytic KSHV Infection in Human Endothelial Cells Reveals Targets of Viral Immune Modulation. <i>Cell Reports</i> , 2020, 33, 108249.	6.4	27
16	Point of Care Nucleic Acid Testing for SARS-CoV-2 in Hospitalized Patients: A Clinical Validation Trial and Implementation Study. <i>Cell Reports Medicine</i> , 2020, 1, 100062.	6.5	47
17	TASOR is a pseudo-PARP that directs HUSH complex assembly and epigenetic transposon control. <i>Nature Communications</i> , 2020, 11, 4940.	12.8	59
18	How does SARS-CoV-2 cause COVID-19?. <i>Science</i> , 2020, 369, 510-511.	12.6	159

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19	Treatment of COVID-19 with remdesivir in the absence of humoral immunity: a case report. <i>Nature Communications</i> , 2020, 11, 6385.	12.8	103
20	A trimeric Rab7 GEF controls NPC1-dependent lysosomal cholesterol export. <i>Nature Communications</i> , 2020, 11, 5559.	12.8	52
21	Ubiquitin-mediated regulation of sterol homeostasis. <i>Current Opinion in Cell Biology</i> , 2020, 65, 103-111.	5.4	29
22	Stromal cell protein kinase C- $\beta$ inhibition enhances chemosensitivity in B cell malignancies and overcomes drug resistance. <i>Science Translational Medicine</i> , 2020, 12, .	12.4	18
23	Integrative functional genomics decodes herpes simplex virus 1. <i>Nature Communications</i> , 2020, 11, 2038.	12.8	61
24	A genome-wide CRISPR screen identifies regulation factors of the TLR3 signalling pathway. <i>Innate Immunity</i> , 2020, 26, 459-472.	2.4	6
25	Antagonism of PP2A is an independent and conserved function of HIV-1 Vif and causes cell cycle arrest. <i>ELife</i> , 2020, 9, .	6.0	15
26	Screening of healthcare workers for SARS-CoV-2 highlights the role of asymptomatic carriage in COVID-19 transmission. <i>ELife</i> , 2020, 9, .	6.0	423
27	Effective control of SARS-CoV-2 transmission between healthcare workers during a period of diminished community prevalence of COVID-19. <i>ELife</i> , 2020, 9, .	6.0	40
28	An Evolutionarily Conserved Function of Polycomb Silences the MHC Class I Antigen Presentation Pathway and Enables Immune Evasion in Cancer. <i>Cancer Cell</i> , 2019, 36, 385-401.e8.	16.8	359
29	Monocytes Latently Infected with Human Cytomegalovirus Evade Neutrophil Killing. <i>IScience</i> , 2019, 12, 13-26.	4.1	29
30	Temporal Proteomic Analysis of BK Polyomavirus Infection Reveals Virus-Induced G <sub>2</sub> Arrest and Highly Effective Evasion of Innate Immune Sensing. <i>Journal of Virology</i> , 2019, 93, .	3.4	28
31	Promiscuous Targeting of Cellular Proteins by Vpr Drives Systems-Level Proteomic Remodeling in HIV-1 Infection. <i>Cell Reports</i> , 2019, 27, 1579-1596.e7.	6.4	75
32	Interferon-Responsive Genes Are Targeted during the Establishment of Human Cytomegalovirus Latency. <i>MBio</i> , 2019, 10, .	4.1	33
33	An Interferon-Driven Oxysterol-Based Defense against Tumor-Derived Extracellular Vesicles. <i>Cancer Cell</i> , 2019, 35, 33-45.e6.	16.8	125
34	Differential viral accessibility (DIVA) identifies alterations in chromatin architecture through large-scale mapping of lentiviral integration sites. <i>Nature Protocols</i> , 2019, 14, 153-170.	12.0	7
35	Functional proteomic atlas of HIV infection in primary human CD4+ T cells. <i>ELife</i> , 2019, 8, .	6.0	34
36	Suppression of costimulation by human cytomegalovirus promotes evasion of cellular immune defenses. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 4998-5003.	7.1	61

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37	Neuropathic MORC2 mutations perturb GHKL ATPase dimerization dynamics and epigenetic silencing by multiple structural mechanisms. <i>Nature Communications</i> , 2018, 9, 651.	12.8	58
38	The sterol-responsive RNF145 E3 ubiquitin ligase mediates the degradation of HMG-CoA reductase together with gp78 and Hrd1. <i>ELife</i> , 2018, 7, .	6.0	85
39	Notch2 controls non-autonomous Wnt-signalling in chronic lymphocytic leukaemia. <i>Nature Communications</i> , 2018, 9, 3839.	12.8	51
40	Regulation of Human $\gamma\delta$ T Cells by BTN3A1 Protein Stability and ATP-Binding Cassette Transporters. <i>Frontiers in Immunology</i> , 2018, 9, 662.	4.8	18
41	The HUSH complex cooperates with TRIM28 to repress young retrotransposons and new genes. <i>Genome Research</i> , 2018, 28, 836-845.	5.5	141
42	MARCH6 and TRC8 facilitate the quality control of cytosolic and tail-anchored proteins. <i>EMBO Reports</i> , 2018, 19, .	4.5	65
43	Hyperactivation of HUSH complex function by Charcot-Marie-Tooth disease mutation in MORC2. <i>Nature Genetics</i> , 2017, 49, 1035-1044.	21.4	105
44	MARCH9-mediated ubiquitination regulates MHC I export from the TGN. <i>Immunology and Cell Biology</i> , 2017, 95, 753-764.	2.3	31
45	Multiple E2 ubiquitin-conjugating enzymes regulate human cytomegalovirus US2-mediated immunoreceptor downregulation. <i>Journal of Cell Science</i> , 2017, 130, 2883-2892.	2.0	18
46	CMTM6 maintains the expression of PD-L1 and regulates anti-tumour immunity. <i>Nature</i> , 2017, 549, 101-105.	27.8	624
47	A genome-wide CRISPR screen reconciles the role of N-linked glycosylation in galectin-3 transport to the cell surface. <i>Journal of Cell Science</i> , 2017, 130, 3234-3247.	2.0	38
48	Control of immune ligands by members of a cytomegalovirus gene expansion suppresses natural killer cell activation. <i>ELife</i> , 2017, 6, .	6.0	67
49	Position-effect variegation revisited: HUSHing up heterochromatin in human cells. <i>BioEssays</i> , 2016, 38, 333-343.	2.5	36
50	Human Cytomegalovirus Infection Upregulates the Mitochondrial Transcription and Translation Machineries. <i>MBio</i> , 2016, 7, e00029.	4.1	55
51	Manipulation of immunometabolism by HIV accessories to the crime?. <i>Current Opinion in Virology</i> , 2016, 19, 65-70.	5.4	13
52	A Genetic Screen Identifies a Critical Role for the WDR81-WDR91 Complex in the Trafficking and Degradation of Tetherin. <i>Traffic</i> , 2016, 17, 940-958.	2.7	21
53	NOTCH1 mediates a switch between two distinct secretomes during senescence. <i>Nature Cell Biology</i> , 2016, 18, 979-992.	10.3	365
54	Genetic dissection of mammalian ERAD through comparative haploid and CRISPR forward genetic screens. <i>Nature Communications</i> , 2016, 7, 11786.	12.8	64

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55	ATF7IP-Mediated Stabilization of the Histone Methyltransferase SETDB1 Is Essential for Heterochromatin Formation by the HUSH Complex. <i>Cell Reports</i> , 2016, 17, 653-659.	6.4	94
56	Mitochondrial Protein Lipoylation and the 2-Oxoglutarate Dehydrogenase Complex Controls HIF1 $\alpha$ Stability in Aerobic Conditions. <i>Cell Metabolism</i> , 2016, 24, 740-752.	16.2	112
57	Temporal proteomic analysis of HIV infection reveals remodelling of the host phosphoproteome by lentiviral Vif variants. <i>ELife</i> , 2016, 5, .	6.0	76
58	A non-canonical ESCRT pathway, including histidine domain phosphotyrosine phosphatase (HD-PTP), is used for down-regulation of virally ubiquitinated MHC class I. <i>Biochemical Journal</i> , 2015, 471, 79-88.	3.7	35
59	A non-proteolytic role for ubiquitin in deadenylation of MHC-I mRNA by the RNA-binding E3-ligase MEX-3C. <i>Nature Communications</i> , 2015, 6, 8670.	12.8	41
60	Epigenetic silencing by the HUSH complex mediates position-effect variegation in human cells. <i>Science</i> , 2015, 348, 1481-1485.	12.6	250
61	UBE2L3 Polymorphism Amplifies NF- $\kappa$ B Activation and Promotes Plasma Cell Development, Linking Linear Ubiquitination to Multiple Autoimmune Diseases. <i>American Journal of Human Genetics</i> , 2015, 96, 221-234.	6.2	84
62	<scp>TRIM</scp> 5 $\alpha$ requires Ube2W to anchor Lys63 $\alpha$ -linked ubiquitin chains and restrict reverse transcription. <i>EMBO Journal</i> , 2015, 34, 2078-2095.	7.8	89
63	Plasma Membrane Profiling Defines an Expanded Class of Cell Surface Proteins Selectively Targeted for Degradation by HCMV US2 in Cooperation with UL141. <i>PLoS Pathogens</i> , 2015, 11, e1004811.	4.7	73
64	Human cytomegalovirus: taking the strain. <i>Medical Microbiology and Immunology</i> , 2015, 204, 273-284.	4.8	119
65	Cell Surface Proteomic Map of HIV Infection Reveals Antagonism of Amino Acid Metabolism by Vpu and Nef. <i>Cell Host and Microbe</i> , 2015, 18, 409-423.	11.0	158
66	Identifying the ERAD ubiquitin E3 ligases for viral and cellular targeting of MHC class I. <i>Molecular Immunology</i> , 2015, 68, 106-111.	2.2	38
67	Antibody-Free Magnetic Cell Sorting of Genetically Modified Primary Human CD4+ T Cells by One-Step Streptavidin Affinity Purification. <i>PLoS ONE</i> , 2014, 9, e111437.	2.5	20
68	Two Novel Human Cytomegalovirus NK Cell Evasion Functions Target MICA for Lysosomal Degradation. <i>PLoS Pathogens</i> , 2014, 10, e1004058.	4.7	123
69	TMEM129 is a Derlin-1 associated ERAD E3 ligase essential for virus-induced degradation of MHC-I. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 11425-11430.	7.1	92
70	HCMV pUL135 Remodels the Actin Cytoskeleton to Impair Immune Recognition of Infected Cells. <i>Cell Host and Microbe</i> , 2014, 16, 201-214.	11.0	67
71	Cleavage by signal peptide peptidase is required for the degradation of selected tail-anchored proteins. <i>Journal of Cell Biology</i> , 2014, 205, 847-862.	5.2	73
72	Quantitative Temporal Viromics: An Approach to Investigate Host-Pathogen Interaction. <i>Cell</i> , 2014, 157, 1460-1472.	28.9	409

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73	Latency-Associated Degradation of the MRP1 Drug Transporter During Latent Human Cytomegalovirus Infection. <i>Science</i> , 2013, 340, 199-202.	12.6	129
74	Role for the obesity-related <i>FTO</i> gene in the cellular sensing of amino acids. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 2557-2562.	7.1	150
75	A novel post-transcriptional role for ubiquitin in the differential regulation of MHC class I allotypes. <i>Molecular Immunology</i> , 2013, 55, 135-138.	2.2	14
76	Studying Ubiquitination of MHC Class I Molecules. <i>Methods in Molecular Biology</i> , 2013, 960, 109-125.	0.9	6
77	Haploid Genetic Screens Identify an Essential Role for PLP2 in the Downregulation of Novel Plasma Membrane Targets by Viral E3 Ubiquitin Ligases. <i>PLoS Pathogens</i> , 2013, 9, e1003772.	4.7	42
78	Sterol metabolism regulates neuroserpin polymer degradation in the absence of the unfolded protein response in the dementia FENIB. <i>Human Molecular Genetics</i> , 2013, 22, 4616-4626.	2.9	21
79	Tapasin-related protein TAPBPR is an additional component of the MHC class I presentation pathway. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 3465-3470.	7.1	107
80	MHC class I molecules are preferentially ubiquitinated on endoplasmic reticulum luminal residues during HRD1 ubiquitin E3 ligase-mediated dislocation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 14290-14295.	7.1	58
81	The RNA-binding E3 ubiquitin ligase MEX-3C links ubiquitination with MHC-I mRNA degradation. <i>EMBO Journal</i> , 2012, 31, 3596-3606.	7.8	74
82	Proteomic Plasma Membrane Profiling Reveals an Essential Role for gp96 in the Cell Surface Expression of LDLR Family Members, Including the LDL Receptor and LRP6. <i>Journal of Proteome Research</i> , 2012, 11, 1475-1484.	3.7	68
83	Fluorescence-Based Phenotypic Selection Allows Forward Genetic Screens in Haploid Human Cells. <i>PLoS ONE</i> , 2012, 7, e39651.	2.5	24
84	Endosomal transport via ubiquitination. <i>Trends in Cell Biology</i> , 2011, 21, 647-655.	7.9	88
85	The dominantly expressed class I molecule of the chicken MHC is explained by coevolution with the polymorphic peptide transporter (TAP) genes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 8396-8401.	7.1	88
86	HRD1 and UBE2J1 target misfolded MHC class I heavy chains for endoplasmic reticulum-associated degradation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 2034-2039.	7.1	116
87	What Has the Study of the K3 and K5 Viral Ubiquitin E3 Ligases Taught Us about Ubiquitin-Mediated Receptor Regulation?. <i>Viruses</i> , 2011, 3, 118-131.	3.3	45
88	RNA-binding E3 ubiquitin ligases: novel players in nucleic acid regulation. <i>Biochemical Society Transactions</i> , 2010, 38, 1621-1626.	3.4	44
89	Efficient Internalization of MHC I Requires Lysine <sup>11</sup> and Lysine <sup>63</sup> Mixed Linkage Polyubiquitin Chains. <i>Traffic</i> , 2010, 11, 210-220.	2.7	111
90	Stabilization of an E3 Ligase-Ubiquitin Complex Increases Cell Surface MHC Class I Expression. <i>Journal of Immunology</i> , 2010, 184, 6978-6985.	0.8	17

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91	First Report of <i>Salmonella enterica</i> Serotype Paratyphi A Azithromycin Resistance Leading to Treatment Failure. <i>Journal of Clinical Microbiology</i> , 2010, 48, 4655-4657.	3.9	62
92	Kaposi's Sarcoma-Associated Herpesvirus K3 and K5 Proteins Block Distinct Steps in Transendothelial Migration of Effector Memory CD4+ T Cells by Targeting Different Endothelial Proteins. <i>Journal of Immunology</i> , 2010, 184, 5186-5192.	0.8	33
93	Ubiquitination of lysine-331 by Kaposi's sarcoma-associated herpesvirus protein K5 targets HFE for lysosomal degradation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 16240-16245.	7.1	16
94	Identification of a Lysosomal Pathway Regulating Degradation of the Bone Morphogenetic Protein Receptor Type II. <i>Journal of Biological Chemistry</i> , 2010, 285, 37641-37649.	3.4	59
95	Comparative analysis of techniques to purify plasma membrane proteins. <i>Journal of Biomolecular Techniques</i> , 2010, 21, 108-15.	1.5	62
96	Analysis of the human E2 ubiquitin conjugating enzyme protein interaction network. <i>Genome Research</i> , 2009, 19, 1905-1911.	5.5	134
97	The TRC8 E3 ligase ubiquitinates MHC class I molecules before dislocation from the ER. <i>Journal of Cell Biology</i> , 2009, 186, 685-692.	5.2	132
98	Stable Isotope Labeling by Amino Acids in Cell Culture and Differential Plasma Membrane Proteome Quantitation Identify New Substrates for the MARCH9 Transmembrane E3 Ligase. <i>Molecular and Cellular Proteomics</i> , 2009, 8, 1959-1971.	3.8	49
99	The trafficking and regulation of membrane receptors by the RING-CH ubiquitin E3 ligases. <i>Experimental Cell Research</i> , 2009, 315, 1593-1600.	2.6	96
100	Viral avoidance and exploitation of the ubiquitin system. <i>Nature Cell Biology</i> , 2009, 11, 527-534.	10.3	204
101	Jenner's Irony: Cowpox Taps into T Cell Evasion. <i>Cell Host and Microbe</i> , 2009, 6, 395-397.	11.0	4
102	ESCRT proteins and the regulation of endocytic delivery to lysosomes. <i>Biochemical Society Transactions</i> , 2009, 37, 178-180.	3.4	15
103	The TRC8 E3 ligase ubiquitinates MHC class I molecules before dislocation from the ER. <i>Journal of Experimental Medicine</i> , 2009, 206, i22-i22.	8.5	0
104	The Ubiquitin E3 Ligase MARCH7 is Differentially Regulated by the Deubiquitylating Enzymes USP7 and USP9X. <i>Traffic</i> , 2008, 9, 1130-1145.	2.7	72
105	Orthotopic liver transplantation for subacute hepatic failure following partial treatment of isoniazid-resistant tuberculosis. <i>Transplant Infectious Disease</i> , 2008, 10, 272-275.	1.7	8
106	Down-regulation of NKG2D and Nkp80 ligands by Kaposi's sarcoma-associated herpesvirus K5 protects against NK cell cytotoxicity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 1656-1661.	7.1	159
107	The Varicellovirus UL49.5 Protein Blocks the Transporter Associated with Antigen Processing (TAP) by Inhibiting Essential Conformational Transitions in the 6+6 Transmembrane TAP Core Complex. <i>Journal of Immunology</i> , 2008, 181, 4894-4907.	0.8	32
108	Natural killer cell evasion by an E3 ubiquitin ligase from Kaposi's sarcoma-associated herpesvirus. <i>Biochemical Society Transactions</i> , 2008, 36, 459-463.	3.4	31



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109	A 58-Year-Old Woman with Abdominal Symptoms and Elevated C-Reactive Protein. <i>PLoS Medicine</i> , 2008, 5, e149.	8.4	3
110	HIV-1 Nef-induced Down-Regulation of MHC Class I Requires AP-1 and Clathrin but Not PACS-1 and Is Impeded by AP-2. <i>Molecular Biology of the Cell</i> , 2007, 18, 3351-3365.	2.1	92
111	MARCH-IX mediates ubiquitination and downregulation of ICAM-1. <i>FEBS Letters</i> , 2007, 581, 45-51.	2.8	51
112	Structure and Function: Heat Shock Proteins and Adaptive Immunity. <i>Journal of Immunology</i> , 2007, 179, 2035-2040.	0.8	106
113	Dendritic Cell Stimulation by Mycobacterial Hsp70 Is Mediated Through CCR5. <i>Science</i> , 2006, 314, 454-458.	12.6	162
114	Lysine-63-linked ubiquitination is required for endolysosomal degradation of class I molecules. <i>EMBO Journal</i> , 2006, 25, 1635-1645.	7.8	234
115	Degradation of Endocytosed Epidermal Growth Factor and Virally Ubiquitinated Major Histocompatibility Complex Class I Is Independent of Mammalian ESCRTII. <i>Journal of Biological Chemistry</i> , 2006, 281, 5094-5105.	3.4	160
116	Loss of function of a lupus-associated Fcγ3RIIb polymorphism through exclusion from lipid rafts. <i>Nature Medicine</i> , 2005, 11, 1056-1058.	30.7	301
117	Downregulation of cell surface receptors by the K3 family of viral and cellular ubiquitin E3 ligases. <i>Immunological Reviews</i> , 2005, 207, 112-125.	6.0	117
118	An in vitro model for the regulation of human cytomegalovirus latency and reactivation in dendritic cells by chromatin remodelling. <i>Journal of General Virology</i> , 2005, 86, 2949-2954.	2.9	163
119	Latency, chromatin remodeling, and reactivation of human cytomegalovirus in the dendritic cells of healthy carriers. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 4140-4145.	7.1	322
120	Solution Structure of the Kaposi's Sarcoma-associated Herpesvirus K3 N-terminal Domain Reveals a Novel E2-binding C4HC3-type RING Domain. <i>Journal of Biological Chemistry</i> , 2004, 279, 53840-53847.	3.4	85
121	Molecular studies of anti-HLA-A2 using light-chain shuffling: a structural model for HLA antibody binding. <i>Tissue Antigens</i> , 2004, 63, 345-354.	1.0	4
122	Recent developments in MHC-class-I-mediated antigen presentation. <i>Current Opinion in Immunology</i> , 2004, 16, 82-89.	5.5	76
123	HSP70 Peptide Binding Mutants Separate Antigen Delivery from Dendritic Cell Stimulation. <i>Immunity</i> , 2004, 20, 95-106.	14.3	111
124	Viral Degradation of the MHC Class I Peptide Loading Complex. <i>Immunity</i> , 2004, 20, 305-317.	14.3	99
125	Peptides complexed with the protein HSP70 generate efficient human cytolytic T-lymphocyte responses. <i>Biochemical Society Transactions</i> , 2004, 32, 622-625.	3.4	26
126	The ABC-transporter signature motif is required for peptide translocation but not peptide binding by TAP. <i>European Journal of Immunology</i> , 2003, 33, 422-427.	2.9	18



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127	The Calculus of Immunity. <i>Immunity</i> , 2003, 18, 315-317.	14.3	15
128	Tapasin Is a Facilitator, Not an Editor, of Class I MHC Peptide Binding. <i>Journal of Immunology</i> , 2003, 171, 5287-5295.	0.8	103
129	Powering the peptide pump: TAP crosstalk with energetic nucleotides. <i>Trends in Biochemical Sciences</i> , 2002, 27, 454-461.	7.5	50
130	Ubiquitylation of MHC class I by the K3 viral protein signals internalization and TSG101-dependent degradation. <i>EMBO Journal</i> , 2002, 21, 2418-2429.	7.8	177
131	CD40 Is a Cellular Receptor Mediating Mycobacterial Heat Shock Protein 70 Stimulation of CC-Chemokines. <i>Immunity</i> , 2001, 15, 971-983.	14.3	253
132	The human cytomegalovirus gene product US6 inhibits ATP binding by TAP. <i>EMBO Journal</i> , 2001, 20, 387-396.	7.8	155
133	Cytomegalovirus: from evasion to suppression?. <i>Nature Immunology</i> , 2001, 2, 993-994.	14.5	17
134	Distinct functions and cooperative interaction of the subunits of the transporter associated with antigen processing (TAP). <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 7431-7436.	7.1	69
135	Mobilization of MHC class I molecules from late endosomes to the cell surface following activation of CD34-derived human Langerhans cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 3982-3987.	7.1	78
136	Tartrate-resistant Acid Phosphatase (Acp 5): Identification in Diverse Human Tissues and Dendritic Cells. <i>Journal of Histochemistry and Cytochemistry</i> , 2001, 49, 675-683.	2.5	85
137	Antigen presentation: Peptides and proteins scramble for the exit. <i>Current Biology</i> , 2000, 10, R839-R842.	3.9	3
138	Inhibition of MHC class I-restricted antigen presentation by gamma 2-herpesviruses. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000, 97, 8455-8460.	7.1	201
139	Lemierre Syndrome: Forgotten but Not Extinct—Report of Four Cases. <i>Radiology</i> , 1999, 213, 369-374.	7.3	114
140	Nephropathia Epidemica and Puumala Virus in Austria. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 1999, 18, 467-472.	2.9	32
141	Antigen presentation: TAP dances with ATP. <i>Current Biology</i> , 1999, 9, R820-R824.	3.9	13
142	The N-terminal region of tapasin is required to stabilize the MHC class I loading complex. <i>European Journal of Immunology</i> , 1999, 29, 1858-1870.	2.9	142
143	Antigen presentation: Coming out gracefully. <i>Current Biology</i> , 1998, 8, R605-R608.	3.9	47
144	Soluble Tapasin Restores MHC Class I Expression and Function in the Tapasin-Negative Cell Line .220. <i>Immunity</i> , 1998, 8, 221-231.	14.3	260

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145	Interferon- $\hat{I}^3$ Rapidly Increases Peptide Transporter (TAP) Subunit Expression and Peptide Transport Capacity in Endothelial Cells. <i>Journal of Biological Chemistry</i> , 1997, 272, 16585-16590.	3.4	80
146	The human cytomegalovirus US6 glycoprotein inhibits transporter associated with antigen processing-dependent peptide translocation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1997, 94, 6904-6909.	7.1	262
147	A Critical Role for Tapasin in the Assembly and Function of Multimeric MHC Class I-TAP Complexes. <i>Science</i> , 1997, 277, 1306-1309.	12.6	477
148	Regulation of MHC class I heterodimer stability and interaction with TAP by tapasin. <i>Immunogenetics</i> , 1997, 46, 477-483.	2.4	77
149	Cd8 <sup>high</sup> + (CD57 <sup>+</sup> ) T cells in patients with rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 1997, 40, 237-248.	6.7	44
150	Roles for Calreticulin and a Novel Glycoprotein, Tapasin, in the Interaction of MHC Class I Molecules with TAP. <i>Immunity</i> , 1996, 5, 103-114.	14.3	644
151	Processing and delivery of peptides presented by MHC class I molecules. <i>Current Opinion in Immunology</i> , 1996, 8, 59-67.	5.5	157
152	Human HLA-A0201-restricted cytotoxic T lymphocyte recognition of influenza A is dominated by T cells bearing the V beta 17 gene segment.. <i>Journal of Experimental Medicine</i> , 1995, 181, 79-91.	8.5	274
153	CD8 <sup>high</sup> (CD57 <sup>+</sup> ) T cells in normal, healthy individuals specifically suppress the generation of cytotoxic T lymphocytes to Epstein-Barr virus-transformed B cell lines. <i>European Journal of Immunology</i> , 1994, 24, 2903-2909.	2.9	45