Edward K Wakeland

List of Publications by Year
in descending order

142 papers	14,747 citations	20797 60 h-index	19169 118 g-index
143 all docs	143 docs citations	143 times ranked	16394 citing authors

#	Article	IF	Citations
1	Association between Antibiotic Exposure and Systemic Immune Parameters in Cancer Patients Receiving Checkpoint Inhibitor Therapy. Cancers, 2022, 14, 1327.	1.7	9
2	A conserved long noncoding RNA, GAPLINC, modulates the immune response during endotoxic shock. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	21
3	Serum IgG Profiling of Toddlers Reveals a Subgroup with Elevated Seropositive Antibodies to Viruses Correlating with Increased Vaccine and Autoantigen Responses. Journal of Clinical Immunology, 2021, 41, 1031-1047.	2.0	3
4	IRF1 governs the differential interferon-stimulated gene responses in human monocytes and macrophages by regulating chromatin accessibility. Cell Reports, 2021, 34, 108891.	2.9	46
5	Association between body mass index, dosing strategy, and efficacy of immune checkpoint inhibitors. , 2021, 9, e002349.		16
6	Outcome and Immune Correlates of a Phase II Trial of High-Dose Interleukin-2 and Stereotactic Ablative Radiotherapy for Metastatic Renal Cell Carcinoma. Clinical Cancer Research, 2021, 27, 6716-6725.	3.2	12
7	Humoral and cellular correlates of a novel immune-related adverse event and its treatment. , 2021, 9, e003585.		10
8	Deep sequencing reveals a DAP1 regulatory haplotype that potentiates autoimmunity in systemic lupus erythematosus. Genome Biology, 2020, 21, 281.	3.8	8
9	sncRNA-1 Is a Small Noncoding RNA Produced by Mycobacterium tuberculosis in Infected Cells That Positively Regulates Genes Coupled to Oleic Acid Biosynthesis. Frontiers in Microbiology, 2020, 11, 1631.	1.5	3
10	Transcriptional profiling identifies caspase-1 as a T cellâ€“intrinsic regulator of Th17 differentiation. Journal of Experimental Medicine, 2020, 217, .	4.2	15
11	Finding a unifying SLE expression signature in a sea of heterogeneity. Nature Reviews Rheumatology, 2020, 16, 357-358.	3.5	1
12	Statin Intolerance, Anti-HMGCR Antibodies, and Immune Checkpoint Inhibitor-Associated Myositis: A â€œTwo-Hitâ€•Autoimmune Toxicity or Clinical Predisposition?. Oncologist, 2020, 25, e1242-e1245.	1.9	10
13	Late-Onset Immunotherapy Toxicity and Delayed Autoantibody Changes: Checkpoint Inhibitorâ€“Induced Raynaud's-Like Phenomenon. Oncologist, 2020, 25, e753-e757.	1.9	17
14	Spatial Network Mapping of Pulmonary Multidrug-Resistant Tuberculosis Cavities Using RNA Sequencing. American Journal of Respiratory and Critical Care Medicine, 2019, 200, 370-380.	2.5	27
15	Amino acid signatures of HLA Class-I and II molecules are strongly associated with SLE susceptibility and autoantibody production in Eastern Asians. PLoS Genetics, 2019, 15, e1008092.	1.5	36
16	Immune dysregulation in cancer patients developing immune-related adverse events. British Journal of Cancer, 2019, 120, 63-68.	2.9	126
17	A plausibly causal functional lupus-associated risk variant in the STAT1â€“STAT4 locus. Human Molecular Genetics, 2018, 27, 2392-2404.	1.4	34
18	Tollâ€Like Receptor 9 Deficiency Breaks Tolerance to RNAâ€Associated Antigens and Upâ€Regulates Tollâ€Like Receptor 7 Protein in <i>Sle1</i> Mice. Arthritis and Rheumatology, 2018, 70, 1597-1609.	2.9	12

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19	GG-04â€¦Pathogenic role of SAT1 variants in monogenic lupus. , 2018, , .		0
20	Co-circulation dynamics and persistence of newly introduced clades of 2012 outbreak associated West Nile Virus in Texas, 2012â€"2015. Infection, Genetics and Evolution, 2018, 66, 13-17.	1.0	0
21	Association of Novel ALX4 Gene Polymorphisms with Antidepressant Treatment Response: Findings from the CO-MED Trial. Molecular Neuropsychiatry, 2018, 4, 7-19.	3.0	12
22	The impact of Staphylococcus aureus genomic variation on clinical phenotype of children with acute hematogenous osteomyelitis. Heliyon, 2018, 4, e00674.	1.4	10
23	T cell-intrinsic IL-1R signaling licenses effector cytokine production by memory CD4 T cells. Nature Communications, 2018, 9, 3185.	5.8	94
24	Drug-Penetration Gradients Associated with Acquired Drug Resistance in Patients with Tuberculosis. American Journal of Respiratory and Critical Care Medicine, 2018, 198, 1208-1219.	2.5	130
25	A missense variant in NCF1 is associated with susceptibility to multiple autoimmune diseases. Nature Genetics, 2017, 49, 433-437.	9.4	143
26	Dysregulated Lymphoid Cell Populations in Mouse Models of Systemic Lupus Erythematosus. Clinical Reviews in Allergy and Immunology, 2017, 53, 181-197.	2.9	24
27	RAD51 interconnects between DNA replication, DNA repair and immunity. Nucleic Acids Research, 2017, 45, 4590-4605.	6.5	127
28	Transancestral mapping and genetic load in systemic lupus erythematosus. Nature Communications, 2017, 8, 16021.	5.8	314
29	Linezolid Dose That Maximizes Sterilizing Effect While Minimizing Toxicity and Resistance Emergence for Tuberculosis. Antimicrobial Agents and Chemotherapy, 2017, 61, .	1.4	81
30	Distinct patterns of innate immune activation by clinical isolates of respiratory syncytial virus. PLoS ONE, 2017, 12, e0184318.	1.1	24
31	Identification of a Systemic Lupus Erythematosus Risk Locus Spanning <i>ATG16L2, FCHSD2</i>, and <i>P2RY2</i> in Koreans. Arthritis and Rheumatology, 2016, 68, 1197-1209.	2.9	89
32	Regulatory polymorphisms modulate the expression of HLA class II molecules and promote autoimmunity. ELife, 2016, 5, .	2.8	113
33	Personalized Immunomonitoring Uncovers Molecular Networks that Stratify Lupus Patients. Cell, 2016, 165, 551-565.	13.5	524
34	RNA sequencing reveals the consequences of a novel insertion in dedicator of cytokinesis-8. Journal of Allergy and Clinical Immunology, 2016, 138, 289-292.e6.	1.5	6
35	Fatty Acid Amide Hydrolase Regulates Peripheral B Cell Receptor Revision, Polyreactivity, and B1 Cells in Lupus. Journal of Immunology, 2016, 196, 1507-1516.	0.4	10
36	DNA polymerase-Î± regulates the activation of type I interferons through cytosolic RNA:DNA synthesis. Nature Immunology, 2016, 17, 495-504.	7.0	123

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37	Autoimmune Diseases in the Bioinformatics Paradigm. Genomics, Proteomics and Bioinformatics, 2015, 13, 205-207.	3.0	0
38	B Cellâ€"Intrinsic CD84 and Ly108 Maintain Germinal Center B Cell Tolerance. Journal of Immunology, 2015, 194, 4130-4143.	0.4	53
39	Differential outcome of TRIF-mediated signaling in TLR4 and TLR3 induced DC maturation. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 13994-13999.	3.3	55
40	Cutting Edge: Inhibiting TBK1 by Compound II Ameliorates Autoimmune Disease in Mice. Journal of Immunology, 2015, 195, 4573-4577.	0.4	61
41	RNA sensing by conventional dendritic cells is central to the development of lupus nephritis. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, E6195-204.	3.3	49
42	Genomic Heterogeneity of Methicillin Resistant Staphylococcus aureus Associated with Variation in Severity of Illness among Children with Acute Hematogenous Osteomyelitis. PLoS ONE, 2015, 10, e0130415.	1.1	18
43	T/Bâ€cell interactions are more transient in response to weak stimuli in SLEâ€prone mice. European Journal of Immunology, 2014, 44, 3522-3531.	1.6	18
44	Allelic heterogeneity in NCF2 associated with systemic lupus erythematosus (SLE) susceptibility across four ethnic populations. Human Molecular Genetics, 2014, 23, 1656-1668.	1.4	67
45	Two Functional Lupus-Associated BLK Promoter Variants Control Cell-Type- and Developmental-Stage-Specific Transcription. American Journal of Human Genetics, 2014, 94, 586-598.	2.6	59
46	Whole transcriptome RNA-seq analysis: tumorigenesis and metastasis of melanoma. Gene, 2014, 548, 234-243.	1.0	25
47	Hunting Autoimmune Disease Genes in NOD: Early Steps on a Long Road to Somewhere Important (Hopefully). Journal of Immunology, 2014, 193, 3-6.	0.4	3
48	Dynamic transcriptomes of human myeloid leukemia cells. Genomics, 2013, 102, 250-256.	1.3	32
49	Transcriptome dynamics during human erythroid differentiation and development. Genomics, 2013, 102, 431-441.	1.3	22
50	Trex1 regulates lysosomal biogenesis and interferon-independent activation of antiviral genes. Nature Immunology, 2013, 14, 61-71.	7.0	122
51	Brief Report: Singleâ€nucleotide polymorphisms in <i>VKORC1</i> are risk factors for systemic lupus erythematosus in Asians. Arthritis and Rheumatism, 2013, 65, 211-215.	6.7	10
52	Dense genotyping of immune-related disease regions identifies 14 new susceptibility loci for juvenile idiopathic arthritis. Nature Genetics, 2013, 45, 664-669.	9.4	337
53	DGKE Variants Cause a Glomerular Microangiopathy That Mimics Membranoproliferative GN. Journal of the American Society of Nephrology: JASN, 2013, 24, 377-384.	3.0	130
54	A Role for Ly108 in the Induction of Promyelocytic Zinc Finger Transcription Factor in Developing Thymocytes. Journal of Immunology, 2013, 190, 2121-2128.	0.4	53

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55	SLE Peripheral Blood B Cell, T Cell and Myeloid Cell Transcriptomes Display Unique Profiles and Each Subset Contributes to the Interferon Signature. PLoS ONE, 2013, 8, e67003.	1.1	165
56	Complete Genome Analysis of Three Acinetobacter baumannii Clinical Isolates in China for Insight into the Diversification of Drug Resistance Elements. PLoS ONE, 2013, 8, e66584.	1.1	107
57	Few Single Nucleotide Variations in Exomes of Human Cord Blood Induced Pluripotent Stem Cells. PLoS ONE, 2013, 8, e59908.	1.1	31
58	B Cell TLR7 Expression Drives Anti-RNA Autoantibody Production and Exacerbates Disease in Systemic Lupus Erythematosusâ€"Prone Mice. Journal of Immunology, 2012, 189, 5786-5796.	0.4	111
59	Association of two independent functional risk haplotypes in <i>TNIP1</i> with systemic lupus erythematosus. Arthritis and Rheumatism, 2012, 64, 3695-3705.	6.7	69
60	Influence of Carbon Monoxide on Growth and Apoptosis of Human Umbilical Artery Smooth Muscle Cells and Vein Endothelial Cells. International Journal of Biological Sciences, 2012, 8, 1431-1446.	2.6	16
61	Identification of IRF8, TMEM39A, and IKZF3-ZPBP2 as Susceptibility Loci for Systemic Lupus Erythematosus in a Large-Scale Multiracial Replication Study. American Journal of Human Genetics, 2012, 90, 648-660.	2.6	161
62	Genetic predisposition to autoimmunity â€" What have we learned?. Seminars in Immunology, 2011, 23, 67-83.	2.7	68
63	The Antibacterial Lectin RegIIIÎ³ Promotes the Spatial Segregation of Microbiota and Host in the Intestine. Science, 2011, 334, 255-258.	6.0	1,163
64	Internal standard-based analysis of microarray data2â€"Analysis of functional associations between HVE-genes. Nucleic Acids Research, 2011, 39, 7881-7899.	6.5	14
65	Dysregulated expression of CXCR4/CXCL12 in subsets of patients with systemic lupus erythematosus. Arthritis and Rheumatism, 2010, 62, 3436-3446.	6.7	79
66	The role of SLAM/CD2 polymorphisms in systemic autoimmunity. Current Opinion in Immunology, 2010, 22, 706-714.	2.4	44
67	Optimal Germinal Center Responses RequireÂa Multistage T Cell:B Cell Adhesion Process Involving Integrins, SLAM-Associated Protein, and CD84. Immunity, 2010, 32, 253-265.	6.6	341
68	<i>Slam</i> Haplotypes Modulate the Response to Lipopolysaccharide In Vivo through Control of NKT Cell Number and Function. Journal of Immunology, 2010, 185, 144-156.	0.4	14
69	Type I Interferons Produced by Resident Renal Cells May Promote End-Organ Disease in Autoantibody-Mediated Glomerulonephritis. Journal of Immunology, 2009, 183, 6831-6838.	0.4	82
70	CXCR4/CXCL12 Hyperexpression Plays a Pivotal Role in the Pathogenesis of Lupus. Journal of Immunology, 2009, 182, 4448-4458.	0.4	109
71	Kallikrein genes are associated with lupus and glomerular basement membraneâ€"specific antibodyâ€"induced nephritis in mice and humans. Journal of Clinical Investigation, 2009, 119, 911-923.	3.9	114
72	Systemic IFNâ€Î± drives kidney nephritis in B6.<i>Sle123 </i>mice. European Journal of Immunology, 2008, 38, 1948-1960.	1.6	89

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73	<i>Yaa</i> autoimmune phenotypes are conferred by overexpression of TLR7. European Journal of Immunology, 2008, 38, 1971-1978.	1.6	150
74	A nonsynonymous functional variant in integrin-Î±M (encoded by ITGAM) is associated with systemic lupus erythematosus. Nature Genetics, 2008, 40, 152-154.	9.4	277
75	Genome-wide association scan in women with systemic lupus erythematosus identifies susceptibility variants in ITGAM, PXK, KIAA1542 and other loci. Nature Genetics, 2008, 40, 204-210.	9.4	1,192
76	Antigen-specific responses and ANA production in B6.Sle1b mice: A role for SAP. Journal of Autoimmunity, 2008, 31, 345-353.	3.0	17
77	Interferon regulatory factor 5 participates in Tollâ€like receptor 7 signaling. FASEB Journal, 2008, 22, 434-434.	0.2	0
78	Tissue kallkreins protect mice against antiâ€GBM induced nephritis and are potential Sle3 candidate genes. FASEB Journal, 2008, 22, 466-466.	0.2	4
79	Death-effector domain-containing protein DEDD is an inhibitor of mitotic Cdk1/cyclin B1. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 2289-2294.	3.3	27
80	PI3K/AKT/mTOR hypersignaling in autoimmune lymphoproliferative disease engendered by the epistatic interplay of Sle1b and FASlpr. International Immunology, 2007, 19, 509-522.	1.8	34
81	Autoantibody profiling to identify individuals at risk for systemic lupus erythematosus. Journal of Autoimmunity, 2006, 27, 153-160.	3.0	162
82	The role of SAP and the SLAM family in autoimmunity. Current Opinion in Immunology, 2006, 18, 656-664.	2.4	29
83	Regulation of B Cell Tolerance by the Lupus Susceptibility Gene Ly108. Science, 2006, 312, 1665-1669.	6.0	233
84	Systemic Lupus Erythematosus: Multiple Immunological Phenotypes in a Complex Genetic Disease. Advances in Immunology, 2006, 92, 1-69.	1.1	165
85	Immune dysregulation accelerates atherosclerosis and modulates plaque composition in systemic lupus erythematosus. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 7018-7023.	3.3	71
86	A Tlr7 translocation accelerates systemic autoimmunity in murine lupus. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 9970-9975.	3.3	567
87	Epistatic Suppression of Systemic Lupus Erythematosus: Fine Mapping of<i>Sles1</i>to Less Than 1 Mb. Journal of Immunology, 2005, 175, 1062-1072.	0.4	39
88	Genetic Dissection of the Murine Lupus Susceptibility Locus<i>Sle2</i>: Contributions to Increased Peritoneal B-1a Cells and Lupus Nephritis Map to Different Loci. Journal of Immunology, 2005, 175, 936-943.	0.4	55
89	Sle1ab Mediates the Aberrant Activation of STAT3 and Ras-ERK Signaling Pathways in B Lymphocytes. Journal of Immunology, 2005, 174, 1630-1637.	0.4	40
90	Enhanced Egg-Induced Immunopathology Correlates With High IFN-Î³ in Murine Schistosomiasis: Identification of Two Epistatic Genetic Intervals. Journal of Immunology, 2005, 174, 435-440.	0.4	23

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91	Selective Expression of the 21-Kilodalton Tyrosine-Phosphorylated Form of TCR Î¶ Promotes the Emergence of T Cells with Autoreactive Potential. Journal of Immunology, 2005, 174, 6071-6079.	0.4	15
92	T cell hyperactivity in lupus as a consequence of hyperstimulatory antigen-presenting cells. Journal of Clinical Investigation, 2005, 115, 1869-1878.	3.9	108
93	Association of Extensive Polymorphisms in the SLAM/CD2 Gene Cluster with Murine Lupus. Immunity, 2004, 21, 769-780.	6.6	253
94	The mapping of quantitative trait loci underlying strain differences in locomotor activity between 129S6 and C57BL/6J mice. Mammalian Genome, 2003, 14, 692-702.	1.0	36
95	A Murine Locus on Chromosome 18 Controls NKT Cell Homeostasis and Th Cell Differentiation. Journal of Immunology, 2003, 171, 4613-4620.	0.4	25
96	Genetic Modulation of Tau Phosphorylation in the Mouse. Journal of Neuroscience, 2003, 23, 187-192.	1.7	80
97	Genetic Modifiers of Systemic Lupus Erythematosus in FcÎ³RIIBâˆ’/âˆ’ Mice. Journal of Experimental Medicine, 2002, 195, 1167-1174.	4.2	238
98	Genetic Dissection of Systemic Lupus Erythematosus Pathogenesis: Evidence for Functional Expression of<i>Sle3/5</i>by Non-T Cells. Journal of Immunology, 2002, 169, 4025-4032.	0.4	50
99	The Major Murine Systemic Lupus Erythematosus Susceptibility Locus<i>Sle1</i>Results in Abnormal Functions of Both B and T Cells. Journal of Immunology, 2002, 169, 2694-2700.	0.4	85
100	Delineating the Genetic Basis of Systemic Lupus Erythematosus. Immunity, 2001, 15, 397-408.	6.6	529
101	Cr2, a Candidate Gene in the Murine Sle1c Lupus Susceptibility Locus, Encodes a Dysfunctional Protein. Immunity, 2001, 15, 775-785.	6.6	214
102	The genetics of lupus. Current Opinion in Nephrology and Hypertension, 2001, 10, 437-443.	1.0	7
103	The genetics of complex autoimmune diseases: non-MHC susceptibility genes. Nature Immunology, 2001, 2, 802-809.	7.0	338
104	Genetic dissection of systemic lupus erythematosus. Current Opinion in Immunology, 1999, 11, 701-707.	2.4	148
105	Multiplex inheritance of component phenotypes in a murine model of lupus. Mammalian Genome, 1999, 10, 176-181.	1.0	91
106	Epistatic Modifiers of Autoimmunity in a Murine Model of Lupus Nephritis. Immunity, 1999, 11, 131-139.	6.6	177
107	Genetic dissection of lupus pathogenesis: a recipe for nephrophilic autoantibodies. Journal of Clinical Investigation, 1999, 103, 1685-1695.	3.9	162
108	Mouse chromosome 3. Mammalian Genome, 1998, 8, S50-S67.	1.0	1

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109	Susceptibility to lupus nephritis in the NZB/W model system. Current Opinion in Immunology, 1998, 10, 718-725.	2.4	71
110	Genetic dissection of lupus nephritis in murine models of SLE. Journal of Clinical Immunology, 1997, 17, 272-281.	2.0	61
111	Mouse chromosome 3. Mammalian Genome, 1997, 7, S45-S59.	1.0	4
112	Ancestral Polymorphism of <i>Mhc</i> Class II Genes in Mice: Implications for Balancing Selection and the Mammalian Molecular Clock. Genetics, 1997, 146, 655-668.	1.2	61
113	Evolutionary origins of retroposon lineages of Mhc class II Ab alleles. Immunogenetics, 1996, 43, 115-24.	1.2	3
114	Communal nesting and communal nursing in house mice, Mus musculus domesticus. Animal Behaviour, 1995, 50, 741-751.	0.8	163
115	The Rab Protein Family: Genetic Mapping of Six Rab Genes in the Mouse. Genomics, 1995, 30, 439-444.	1.3	30
116	Polygenic control of susceptibility to murine systemic lupus erythematosus. Immunity, 1994, 1, 219-229.	6.6	476
117	Evolution of MHC genetic diversity: a tale of incest, pestilence and sexual preference. Trends in Genetics, 1993, 9, 408-412.	2.9	254
118	Communal nesting patterns in mice implicate MHC genes in kin recognition. Nature, 1992, 360, 581-583.	13.7	227
119	Mouse lipocortin I gene structure and chromosomal assignment: Gene duplication and the origins of a gene family. Genomics, 1991, 10, 365-374.	1.3	38
120	Mating patterns in seminatural populations of mice influenced by MHC genotype. Nature, 1991, 352, 619-621.	13.7	640
121	Molecular and Genetic Mechanisms Involved in the Generation of Mhc Diversity. , 1991, , 139-154.		4
122	The Generation and Maintenance of MHC Class II Gene Polymorphism in Rodents. Immunological Reviews, 1990, 113, 207-226.	2.8	48
123	Ancestral polymorphisms of MHC class II genes: Divergent allele advantage. Immunologic Research, 1990, 9, 115-122.	1.3	125
124	Revised nomenclature of mouseH-2 genes. Immunogenetics, 1990, 32, 147-149.	1.2	49
125	Genetic polymorphisms ofQ region genes from wild-derived mice: Implications forQ region evolution. Immunogenetics, 1990, 31, 315-325.	1.2	16
126	The mouse lymph node homing receptor is identical with the lymphocyte cell surface marker Ly-22: Role of the EGF domain in endothelial binding. Cell, 1990, 61, 611-622.	13.5	126

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127	Evolution of diversity at the major histocompatibility complex. Trends in Ecology and Evolution, 1990, 5, 181-187.	4.2	197
128	The origin of MHC class II gene polymorphism within the genus Mus. Nature, 1988, 332, 651-654.	13.7	130
129	Polymorphisms of DQ ? genes in HLA-DR4 haplotypes from healthy and diabetic individuals. Immunogenetics, 1987, 25, 152-160.	1.2	21
130	Production of 35 H-2 homozygous strains from wild mice. Immunogenetics, 1987, 26, 115-119.	1.2	15
131	DNA polymorphism of MHC III genes in inbred and wild mouse strains. Immunogenetics, 1987, 25, 290-298.	1.2	25
132	Extraction of cellular DNA from human cells and tissues fixed in ethanol. Analytical Biochemistry, 1987, 160, 135-138.	1.1	56
133	Serological and biochemical characterization of class II antigens in B10.W lines. Tissue Antigens, 1982, 19, 40-52.	1.0	0
134	Minor structural variants of H-2K-controlled molecules in wild mice. Immunogenetics, 1982, 16, 491-493.	1.2	5
135	An H-2 haplotype possibly derived by crossing-over between the (A Î± A Î²) duplex and the E Î²locus. Immunogenetics, 1981, 14, 273-281.	1.2	9
136	Heterozygosity of Hâ€“2 loci in wild mice. Nature, 1979, 281, 603-605.	13.7	41
137	The histocompatibility-2 system in wild mice. Immunogenetics, 1979, 8, 27-39.	1.2	35
138	Genetic nomenclature for chicken immunoglobulin allotypes: An extensive survey of inbred lines and antisera. Immunogenetics, 1979, 8, 385-404.	1.2	11
139	Structural comparisons of serologically identical IA- and IE-encoded antigens from inbred and wild mice. Immunogenetics, 1979, 9, 535-550.	1.2	15
140	Histocompatibility antigens controlled by theI region of the murineH-2 complex. Immunogenetics, 1977, 5, 445-451.	1.2	13
141	Structural and genetic analysis of four chicken 7S immunoglobulin allotypes. Immunogenetics, 1975, 2, 531-541.	1.2	10
142	The Importance of Epistatic Interactions in the Development of Autoimmunity. Novartis Foundation Symposium, 0, , 76-93.	1.2	5