Betty P Tsao

List of Publications by Year
in descending order

140	10,699	28274	32842
papers	citations	55 h-index	100 g-index

163	163	163	10967
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Human SLE variant NCF1-R90H promotes kidney damage and murine lupus through enhanced Tfh2 responses induced by defective efferocytosis of macrophages. Annals of the Rheumatic Diseases, 2022, 81, 255-267.	0.9	14
2	<scp>Upâ€Regulated</scp> Interleukinâ€10 Induced by <scp>E2F</scp> Transcription Factor 2â€“<scp>MicroRNA</scp>â€17â€5p Circuitry in Extrafollicular Effector B Cells Contributes to Autoantibody Production in Systemic Lupus Erythematosus. Arthritis and Rheumatology, 2022, 74, 496-507.	5.6	12
3	Prediction models of treatment response in lupus nephritis. Kidney International, 2022, 101, 379-389.	5.2	18
4	RNASE2 Mediates Age-Associated B Cell Expansion Through Monocyte Derived IL-10 in Patients With Systemic Lupus Erythematosus. Frontiers in Immunology, 2022, 13, 752189.	4.8	9
5	Complement C4, the Major Histocompatibility Complex, and Autoimmunity. Arthritis and Rheumatology, 2022, 74, 1318-1320.	5.6	4
6	Genes and genetics in human SLE. , 2021, , 85-96.		2
7	IFNL4 Genotype Does Not Associate with CD4 T-Cell Recovery in People Living with Human Immunodeficiency Virus. AIDS Research and Human Retroviruses, 2021, 37, 184-188.	1.1	2
8	Lupus susceptibility genes. , 2021, , 25-33.		0
9	Deep sequencing reveals a DAP1 regulatory haplotype that potentiates autoimmunity in systemic lupus erythematosus. Genome Biology, 2020, 21, 281.	8.8	8
10	Reduced Let-7f in Bone Marrow-Derived Mesenchymal Stem Cells Triggers Treg/Th17 Imbalance in Patients With Systemic Lupus Erythematosus. Frontiers in Immunology, 2020, 11, 233.	4.8	30
11	Rigorous Plasma Microbiome Analysis Method Enables Disease Association Discovery in Clinic. Frontiers in Microbiology, 2020, 11, 613268.	3.5	12
12	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.	3.5	36
13	180â€…Examining the transcriptional impact of liganded estrogen receptor alpha in the inflammatory milieu of systemic lupus erythematosus. , 2019, , .		0
14	Genetics of Human SLE. , 2019, , 54-68.		5
15	A plausibly causal functional lupus-associated risk variant in the STAT1â€“STAT4 locus. Human Molecular Genetics, 2018, 27, 2392-2404.	2.9	34
16	GG-07â€…Regulatory polymorphisms in EMSY gene are associated with autoantibodies in healthy individuals. , 2018, , .		0
17	Genetic contributions to lupus nephritis in a multi-ethnic cohort of systemic lupus erythematous patients. PLoS ONE, 2018, 13, e0199003.	2.5	46
18	Transcription Factor SOX5 Promotes the Migration and Invasion of Fibroblast-Like Synoviocytes in Part by Regulating MMP-9 Expression in Collagen-Induced Arthritis. Frontiers in Immunology, 2018, 9, 749.	4.8	33

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19	Genetic variants in systemic lupus erythematosus susceptibility loci, XKR6 and GLT1D1 are associated with childhood-onset SLE in a Korean cohort. Scientific Reports, 2018, 8, 9962.	3.3	25
20	Genetic fine mapping of systemic lupus erythematosus MHC associations in Europeans and African Americans. Human Molecular Genetics, 2018, 27, 3813-3824.	2.9	43
21	A missense variant in NCF1 is associated with susceptibility to multiple autoimmune diseases. Nature Genetics, 2017, 49, 433-437.	21.4	143
22	Updates in Lupus Genetics. Current Rheumatology Reports, 2017, 19, 68.	4.7	99
23	Transancestral mapping and genetic load in systemic lupus erythematosus. Nature Communications, 2017, 8, 16021.	12.8	314
24	X Chromosome Dose and Sex Bias in Autoimmune Diseases: Increased Prevalence of 47,XXX in Systemic Lupus Erythematosus and SjÃ¶gren's Syndrome. Arthritis and Rheumatology, 2016, 68, 1290-1300.	5.6	114
25	Identification of a Systemic Lupus Erythematosus Risk Locus Spanning ATG16L2, FCHSD2, and P2RY2 in Koreans. Arthritis and Rheumatology, 2016, 68, 1197-1209.	5.6	89
26	Regulatory polymorphisms modulate the expression of HLA class II molecules and promote autoimmunity. ELife, 2016, 5, .	6.0	113
27	Genes and Genetics in Human Systemic Lupus Erythematosus. , 2016, , 69-76.		1
28	Genomeâ€Wide Association Study in an Amerindian Ancestry Population Reveals Novel Systemic Lupus Erythematosus Risk Loci and the Role of European Admixture. Arthritis and Rheumatology, 2016, 68, 932-943.	5.6	138
29	Modulation of IL-6 induced RANKL expression in arthritic synovium by a transcription factor SOX5. Scientific Reports, 2016, 6, 32001.	3.3	41
30	CD3Zhypermethylation is associated with severe clinical manifestations in systemic lupus erythematosus and reduces CD3Î¶-chain expression in T cells. Rheumatology, 2016, 56, kew405.	1.9	12
31	Decreased SMG7 expression associates with lupus-risk variants and elevated antinuclear antibody production. Annals of the Rheumatic Diseases, 2016, 75, 2007-2013.	0.9	16
32	Preferential association of a functional variant in complement receptor 2 with antibodies to double-stranded DNA. Annals of the Rheumatic Diseases, 2016, 75, 242-252.	0.9	10
33	Focused transcription from the human CR2/CD21 core promoter is regulated by synergistic activity of TATA and Initiator elements in mature B cells. Cellular and Molecular Immunology, 2016, 13, 119-131.	10.5	3
34	Lupus risk variants in the PXK locus alter B-cell receptor internalization. Frontiers in Genetics, 2015, 5, 450.	2.3	25
35	Lupus Risk Variant Increases pSTAT1 Binding and Decreases ETS1 Expression. American Journal of Human Genetics, 2015, 96, 731-739.	6.2	36
36	Genetic associations of leptin-related polymorphisms with systemic lupus erythematosus. Clinical Immunology, 2015, 161, 157-162.	3.2	10

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37	The IRF5â€“TNPO3 association with systemic lupus erythematosus has two components that other autoimmune disorders variably share. Human Molecular Genetics, 2015, 24, 582-596.	2.9	74
38	Identification of interferon-inducible genes as diagnostic biomarker for systemic lupus erythematosus. Clinical Rheumatology, 2015, 34, 71-79.	2.2	43
39	Restored Immunosuppressive Effect of Mesenchymal Stem Cells on B Cells After Olfactory 1/Early B Cell Factorâ€“Associated Zincâ€Finger Protein Downâ€Regulation in Patients With Systemic Lupus Erythematosus. Arthritis and Rheumatology, 2014, 66, 3413-3423.	5.6	35
40	Genetics of systemic lupus erythematosus: immune responses and end organ resistance to damage. Current Opinion in Immunology, 2014, 31, 87-96.	5.5	47
41	Advances in lupus genetics and epigenetics. Current Opinion in Rheumatology, 2014, 26, 482-492.	4.3	104
42	Endâ€Stage Renal Disease in African Americans With Lupus Nephritis Is Associated With APOL1. Arthritis and Rheumatology, 2014, 66, 390-396.	5.6	242
43	Two Functional Lupus-Associated BLK Promoter Variants Control Cell-Type- and Developmental-Stage-Specific Transcription. American Journal of Human Genetics, 2014, 94, 586-598.	6.2	59
44	Lupus Nephritis Susceptibility Loci in Women with Systemic Lupus Erythematosus. Journal of the American Society of Nephrology: JASN, 2014, 25, 2859-2870.	6.1	117
45	Transcription factor Ikaros Represses Protein Phosphatase 2A (PP2A) Expression through an Intronic Binding Site. Journal of Biological Chemistry, 2014, 289, 13751-13757.	3.4	20
46	Systemic Lupus Erythematosus, Genetics. , 2014, , 1171-1178.		0
47	Plasma levels of osteopontin identify patients at risk for organ damage in systemic lupus erythematosus. Arthritis Research and Therapy, 2013, 15, R18.	3.5	32
48	Brief Report: Singleâ€nucleotide polymorphisms in VKORC1 are risk factors for systemic lupus erythematosus in Asians. Arthritis and Rheumatism, 2013, 65, 211-215.	6.7	10
49	Variable Association of Reactive Intermediate Genes with Systemic Lupus Erythematosus in Populations with Different African Ancestry. Journal of Rheumatology, 2013, 40, 842-849.	2.0	15
50	Preferential Binding to Elk-1 by SLE-Associated IL10 Risk Allele Upregulates IL10 Expression. PLoS Genetics, 2013, 9, e1003870.	3.5	36
51	Admixture Mapping in Lupus Identifies Multiple Functional Variants within IFIH1 Associated with Apoptosis, Inflammation, and Autoantibody Production. PLoS Genetics, 2013, 9, e1003222.	3.5	107
52	Trans-Ancestral Studies Fine Map the SLE-Susceptibility Locus TNFSF4. PLoS Genetics, 2013, 9, e1003554.	3.5	50
53	MicroRNA-3148 Modulates Allelic Expression of Toll-Like Receptor 7 Variant Associated with Systemic Lupus Erythematosus. PLoS Genetics, 2013, 9, e1003336.	3.5	107
54	Fine mapping of Xq28: bothMECP2 and IRAK1contribute to risk for systemic lupus erythematosus in multiple ancestral groups. Annals of the Rheumatic Diseases, 2013, 72, 437-444.	0.9	97

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55	Recent insights into the genetic basis of systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2013, 72, ii56-ii61.	0.9	117
56	ABIN1 Dysfunction as a Genetic Basis for Lupus Nephritis. Journal of the American Society of Nephrology: JASN, 2013, 24, 1743-1754.	6.1	70
57	Genetics of Human SLE. , 2013, , 35-45.		4
58	PTPN22 Association in Systemic Lupus Erythematosus (SLE) with Respect to Individual Ancestry and Clinical Sub-Phenotypes. PLoS ONE, 2013, 8, e69404.	2.5	57
59	Novel identification of the IRF7 region as an anticentromere autoantibody propensity locus in systemic sclerosis. Annals of the Rheumatic Diseases, 2012, 71, 114-119.	0.9	62
60	Analysis of autosomal genes reveals geneâ€“sex interactions and higher total genetic risk in men with systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2012, 71, 694-699.	0.9	87
61	Impact of genetic ancestry and sociodemographic status on the clinical expression of systemic lupus erythematosus in American Indianâ€“European populations. Arthritis and Rheumatism, 2012, 64, 3687-3694.	6.7	70
62	Association of two independent functional risk haplotypes in TNIP1 with systemic lupus erythematosus. Arthritis and Rheumatism, 2012, 64, 3695-3705.	6.7	69
63	Variation in the ICAM1â€“ICAM4â€“ICAM5 locus is associated with systemic lupus erythematosus susceptibility in multiple ancestries. Annals of the Rheumatic Diseases, 2012, 71, 1809-1814.	0.9	60
64	Evaluation of TRAF6 in a large multiancestral lupus cohort. Arthritis and Rheumatism, 2012, 64, 1960-1969.	6.7	51
65	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.	6.2	161
66	Transcriptional effects of a lupus-associated polymorphism in the 5â€² untranslated region (UTR) of human complement receptor 2 (CR2/CD21). Molecular Immunology, 2012, 52, 165-173.	2.2	12
67	Evidence for geneâ€“gene epistatic interactions among susceptibility loci for systemic lupus erythematosus. Arthritis and Rheumatism, 2012, 64, 485-492.	6.7	53
68	Inhibition of Aberrant Circulating Tfh Cell Proportions by Corticosteroids in Patients with Systemic Lupus Erythematosus. PLoS ONE, 2012, 7, e51982.	2.5	91
69	Differential Genetic Associations for Systemic Lupus Erythematosus Based on Antiâ€“dsDNA Autoantibody Production. PLoS Genetics, 2011, 7, e1001323.	3.5	206
70	Association of a functional variant downstream of TNFAIP3 with systemic lupus erythematosus. Nature Genetics, 2011, 43, 253-258.	21.4	242
71	Identification of a Systemic Lupus Erythematosus Susceptibility Locus at 11p13 between PDHX and CD44 in a Multiethnic Study. American Journal of Human Genetics, 2011, 88, 83-91.	6.2	72
72	Association of a functional IRF7 variant with systemic lupus erythematosus. Arthritis and Rheumatism, 2011, 63, 749-754.	6.7	118

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73	Genetic analyses of interferon pathway-related genes reveal multiple new loci associated with systemic lupus erythematosus. Arthritis and Rheumatism, 2011, 63, 2049-2057.	6.7	45
74	Association of PPP2CA polymorphisms with systemic lupus erythematosus susceptibility in multiple ethnic groups. Arthritis and Rheumatism, 2011, 63, 2755-2763.	6.7	36
75	Identification of novel genetic susceptibility loci in African American lupus patients in a candidate gene association study. Arthritis and Rheumatism, 2011, 63, 3493-3501.	6.7	109
76	A Functional Variant in MicroRNA-146a Promoter Modulates Its Expression and Confers Disease Risk for Systemic Lupus Erythematosus. PLoS Genetics, 2011, 7, e1002128.	3.5	241
77	Risk Alleles for Systemic Lupus Erythematosus in a Large Case-Control Collection and Associations with Clinical Subphenotypes. PLoS Genetics, 2011, 7, e1001311.	3.5	154
78	A Comprehensive Analysis of Shared Loci between Systemic Lupus Erythematosus (SLE) and Sixteen Autoimmune Diseases Reveals Limited Genetic Overlap. PLoS Genetics, 2011, 7, e1002406.	3.5	148
79	Phenotypic associations of genetic susceptibility loci in systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2011, 70, 1752-1757.	0.9	110
80	Constitutive Genes and Lupus. , 2011, , 47-61.		4
81	Association of Genetic Variants in Complement Factor H and Factor H-Related Genes with Systemic Lupus Erythematosus Susceptibility. PLoS Genetics, 2011, 7, e1002079.	3.5	181
82	A functional RANKL polymorphism associated with younger age at onset of rheumatoid arthritis. Arthritis and Rheumatism, 2010, 62, 2864-2875.	6.7	35
83	Association of IRF5 polymorphisms with activation of the interferon Î± pathway. Annals of the Rheumatic Diseases, 2010, 69, 611-617.	0.9	54
84	Sex-specific association of X-linked Toll-like receptor 7 (TLR7) with male systemic lupus erythematosus. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 15838-15843.	7.1	324
85	Male-only Systemic Lupus. Journal of Rheumatology, 2010, 37, 1480-1487.	2.0	13
86	Treatment with apolipoprotein A-1 mimetic peptide reduces lupus-like manifestations in a murine lupus model of accelerated atherosclerosis. Arthritis Research and Therapy, 2010, 12, R93.	3.5	47
87	Genetic susceptibility to systemic lupus erythematosus in the genomic era. Nature Reviews Rheumatology, 2010, 6, 683-692.	8.0	319
88	Olf1/EBF associated zinc finger protein interfered with antinuclear antibody production in patients with systemic lupus erythematosus. Arthritis Research and Therapy, 2010, 12, R59.	3.5	8
89	European population substructure is associated with mucocutaneous manifestations and autoantibody production in systemic lupus erythematosus. Arthritis and Rheumatism, 2009, 60, 2448-2456.	6.7	27
90	Plasmin immunization preferentially induces potentially prothrombotic IgG anticardiolipin antibodies in MRL/MpJ mice. Arthritis and Rheumatism, 2009, 60, 3108-3117.	6.7	2

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91	Pathogenesis of Systemic Lupus Erythematosus. , 2009, , 1233-1262.		4
92	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.	21.4	1,192
93	A loss-of-function variant of PTPN22 is associated with reduced risk of systemic lupus erythematosus. Human Molecular Genetics, 2008, 18, 569-579.	2.9	106
94	A genome wide association study of systemic lupus erythematosus (SLE) by SLEGEN, the International SLE Genetics Consortium.. FASEB Journal, 2008, 22, 850.1.	0.5	0
95	Association of a common complement receptor 2 haplotype with increased risk of systemic lupus erythematosus. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 3961-3966.	7.1	62
96	ApoEâ^'/â^'Fasâ^'/â^' C57BL/6 mice: a novel murine model simultaneously exhibits lupus nephritis, atherosclerosis, and osteopenia. Journal of Lipid Research, 2007, 48, 794-805.	4.2	62
97	Gene Copy-Number Variation and Associated Polymorphisms of Complement Component C4 in Human Systemic Lupus Erythematosus (SLE): Low Copy Number Is a Risk Factor for and High Copy Number Is a Protective Factor against SLE Susceptibility in European Americans. American Journal of Human Genetics, 2007, 80, 1037-1054.	6.2	411
98	Current topics in human SLE genetics. Seminars in Immunopathology, 2006, 28, 97-107.	4.0	61
99	Association of tumor necrosis factor Î± polymorphism, but not the shared epitope, with increased radiographic progression in a seropositive rheumatoid arthritis inception cohort. Arthritis and Rheumatism, 2006, 54, 1105-1116.	6.7	49
100	Association of increased interferon-inducible gene expression with disease activity and lupus nephritis in patients with systemic lupus erythematosus. Arthritis and Rheumatism, 2006, 54, 2951-2962.	6.7	404
101	Association analysis of the R620W polymorphism of protein tyrosine phosphatase PTPN22 in systemic lupus erythematosus families: Increased t allele frequency in systemic lupus erythematosus patients with autoimmune thyroid disease. Arthritis and Rheumatism, 2005, 52, 2396-2402.	6.7	80
102	Maternal HLA class II compatibility in men with systemic lupus erythematosus. Arthritis and Rheumatism, 2005, 52, 2768-2773.	6.7	36
103	A stop codon polymorphism of Toll-like receptor 5 is associated with resistance to systemic lupus erythematosus. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 10593-10597.	7.1	144
104	CD72 polymorphisms associated with alternative splicing modify susceptibility to human systemic lupus erythematosus through epistatic interaction with FCGR2B. Human Molecular Genetics, 2004, 13, 2907-2917.	2.9	43
105	Current advances in the human lupus genetics. Current Rheumatology Reports, 2004, 6, 391-398.	4.7	35
106	Association of Fc? receptor IIA, but not IIB and IIIA, polymorphisms with systemic lupus erythematosus: A family-based association study in Caucasians. Arthritis and Rheumatism, 2004, 50, 671-673.	6.7	34
107	Systemic lupus erythematosus genome scan: Support for linkage at 1q23, 2q33, 16q12-13, and 17q21-23 and novel evidence at 3p24, 10q23-24, 13q32, and 18q22-23. Arthritis and Rheumatism, 2004, 50, 3203-3210.	6.7	66
108	Interaction betweenRANKL andHLA-DRB1 genotypes may contribute to younger age at onset of seropositive rheumatoid arthritis in an inception cohort. Arthritis and Rheumatism, 2004, 50, 3093-3103.	6.7	42

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109	Update on human systemic lupus erythematosus genetics. Current Opinion in Rheumatology, 2004, 16, 513-521.	4.3	139
110	Identification and characterization of SmD183-119-reactive T cells that provide T cell help for pathogenic anti-double-stranded DNA antibodies. Arthritis and Rheumatism, 2003, 48, 475-485.	6.7	216
111	Identification and characterization of a peptide mimetic that may detect a species of disease-associated anticardiolipin antibodies in patients with the antiphospholipid syndrome. Arthritis and Rheumatism, 2003, 48, 737-745.	6.7	14
112	The genetics of human systemic lupus erythematosus. Trends in Immunology, 2003, 24, 595-602.	6.8	165
113	Familiality and co-occurrence of clinical features of systemic lupus erythematosus. Arthritis and Rheumatism, 2002, 46, 2678-2685.	6.7	51
114	Linkage and interaction of loci on 1q23 and 16q12 may contribute to susceptibility to systemic lupus erythematosus. Arthritis and Rheumatism, 2002, 46, 2928-2936.	6.7	55
115	An update on genetic studies of systemic lupus erythematosus. Current Rheumatology Reports, 2002, 4, 359-367.	4.7	46
116	FcÎ³ receptor IIIA polymorphism in Korean patients with systemic lupus erythematosus. Rheumatology International, 2002, 21, 222-226.	3.0	17
117	Genetics and systemic lupus erythematosus. Current Rheumatology Reports, 2001, 3, 183-190.	4.7	22
118	Treatment with a consensus peptide based on amino acid sequences in autoantibodies prevents T cell activation by autoantigens and delays disease onset in murine lupus. Arthritis and Rheumatism, 2001, 44, 432-441.	6.7	103
119	Poly(ADP)â€ribose polymerase and susceptibility to systemic lupus erythematosus and primary antiphospholipid syndrome: Comment on the article by Delrieu et al. Arthritis and Rheumatism, 2000, 43, 1421-1422.	6.7	10
120	Genetics and systemic lupus erythematosus. Current Rheumatology Reports, 2000, 2, 13-18.	4.7	14
121	Single-nucleotide polymorphisms of T cell receptor ? chain in patients with systemic lupus erythematosus. Arthritis and Rheumatism, 1999, 42, 2601-2605.	6.7	29
122	PARP alleles within the linked chromosomal region are associated with systemic lupus erythematosus. Journal of Clinical Investigation, 1999, 103, 1135-1140.	8.2	99
123	Autoantibodies as a Source of Peptides That Regulate Autoantibody Production. , 1999, , 371-388.		0
124	Abnormal distribution of Fc? receptor type IIa polymorphisms in Korean patients with systemic lupus erythematosus. Arthritis and Rheumatism, 1998, 41, 421-426.	6.7	92
125	Altered Immune Responses in Interleukin 10 Transgenic Mice. Journal of Experimental Medicine, 1997, 185, 2101-2110.	8.5	261
126	Commentary: Genetics of systemic lupus erythematosus. Current Opinion in Rheumatology, 1997, 9, 377-379.	4.3	10

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127	Autoimmunity and Tolerance in Ig-Transgenic Mice: Murine SLE as a Model to Study B Cell Tolerance. International Reviews of Immunology, 1994, 11, 305-320.	3.3	0
128	Comparison of pathogenic and non-pathogenic murine antibodies to DNA: antigen binding and structural characteristics. International Immunology, 1994, 6, 817-830.	4.0	145
129	A peptide derived from an autoantibody can stimulate t cells in the (nzb Ã— nzw)f1 mouse model of systemic lupus erythematosus. Arthritis and Rheumatism, 1993, 36, 355-364.	6.7	70
130	B cells are anergic in transgenic mice that express IgM anti-DNA antibodies. European Journal of Immunology, 1993, 23, 2332-2339.	2.9	48
131	Contribution of Major Histocompatibility Complex (MHC) to Upregulation of Anti-DNA Antibody in Transgenic Mice. Journal of Autoimmunity, 1993, 6, 1-9.	6.5	3
132	Ig-transgenic mice as models for studying the regulation and role of anti-DNA antibodies in murine lupus. ImmunoMethods, 1992, 1, 185-190.	0.8	2
133	T cell up-regulation of B cells via their idiotypes contributing to the development of systemic lupus erythematosus. American Journal of Medicine, 1988, 85, 32-34.	1.5	8
134	Idiotype selection is an immunoregulatory mechanism which contributes to the pathogenesis of systemic lupus erythematosus. Journal of Autoimmunity, 1988, 1, 673-681.	6.5	1
135	In vivofunctional analysis ofin vitroprotein binding sites in the immunoglobulin heavy chain enhancer. Nucleic Acids Research, 1988, 16, 3239-3253.	14.5	64
136	The role of cytoplasmic free calcium concentration in B-cell tolerance. Cellular Immunology, 1987, 108, 335-342.	3.0	7
137	Macrophage-derived soluble factors mediate suppression induced by 2,4-dinitrophenyl-conjugated mouse IgG in hybridoma cells. Cellular Immunology, 1985, 91, 362-374.	3.0	0
138	Central suppression of monoclonal B cells: DNP-MGG suppresses proliferation and immunoglobulin synthesis in anti-DNP-secreting hybridoma and myeloma. Cellular Immunology, 1984, 88, 96-108.	3.0	6
139	Evidence that the hydrophobic domain of rat renal Î³-glutamyltransferase spans the brush border membrane. Biochimica Et Biophysica Acta - Biomembranes, 1982, 690, 199-206.	2.6	5
140	Membrane association and orientation of rat renal activities capable of degrading glutathione. International Journal of Biochemistry & Cell Biology, 1980, 12, 219-222.	0.5	7