

# Betty P Tsao

## List of Publications by Year in descending order

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140  
papers

10,699  
citations

32410

55  
h-index

37326

100  
g-index

163  
all docs

163  
docs citations

163  
times ranked

11944  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide association scan in women with systemic lupus erythematosus identifies susceptibility variants in ITGAM, PTK, KIAA1542 and other loci. <i>Nature Genetics</i> , 2008, 40, 204-210.	9.4	1,192
2	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. <i>American Journal of Human Genetics</i> , 2007, 80, 1037-1054.	2.6	411
3	Association of increased interferon-inducible gene expression with disease activity and lupus nephritis in patients with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2006, 54, 2951-2962.	6.7	404
4	Sex-specific association of X-linked Toll-like receptor 7 (TLR7) with male systemic lupus erythematosus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 15838-15843.	3.3	324
5	Genetic susceptibility to systemic lupus erythematosus in the genomic era. <i>Nature Reviews Rheumatology</i> , 2010, 6, 683-692.	3.5	319
6	Transancestral mapping and genetic load in systemic lupus erythematosus. <i>Nature Communications</i> , 2017, 8, 16021.	5.8	314
7	Altered Immune Responses in Interleukin 10 Transgenic Mice. <i>Journal of Experimental Medicine</i> , 1997, 185, 2101-2110.	4.2	261
8	Association of a functional variant downstream of TNFAIP3 with systemic lupus erythematosus. <i>Nature Genetics</i> , 2011, 43, 253-258.	9.4	242
9	End-stage Renal Disease in African Americans With Lupus Nephritis Is Associated With <i>APOL1</i> . <i>Arthritis and Rheumatology</i> , 2014, 66, 390-396.	2.9	242
10	A Functional Variant in MicroRNA-146a Promoter Modulates Its Expression and Confers Disease Risk for Systemic Lupus Erythematosus. <i>PLoS Genetics</i> , 2011, 7, e1002128.	1.5	241
11	Identification and characterization of SmD183-119-reactive T cells that provide T cell help for pathogenic anti-double-stranded DNA antibodies. <i>Arthritis and Rheumatism</i> , 2003, 48, 475-485.	6.7	216
12	Differential Genetic Associations for Systemic Lupus Erythematosus Based on Anti-dsDNA Autoantibody Production. <i>PLoS Genetics</i> , 2011, 7, e1001323.	1.5	206
13	Association of Genetic Variants in Complement Factor H and Factor H-Related Genes with Systemic Lupus Erythematosus Susceptibility. <i>PLoS Genetics</i> , 2011, 7, e1002079.	1.5	181
14	The genetics of human systemic lupus erythematosus. <i>Trends in Immunology</i> , 2003, 24, 595-602.	2.9	165
15	Identification of IRF8, TMEM39A, and IKZF3-ZBP2 as Susceptibility Loci for Systemic Lupus Erythematosus in a Large-Scale Multiracial Replication Study. <i>American Journal of Human Genetics</i> , 2012, 90, 648-660.	2.6	161
16	Risk Alleles for Systemic Lupus Erythematosus in a Large Case-Control Collection and Associations with Clinical Subphenotypes. <i>PLoS Genetics</i> , 2011, 7, e1001311.	1.5	154
17	A Comprehensive Analysis of Shared Loci between Systemic Lupus Erythematosus (SLE) and Sixteen Autoimmune Diseases Reveals Limited Genetic Overlap. <i>PLoS Genetics</i> , 2011, 7, e1002406.	1.5	148
18	Comparison of pathogenic and non-pathogenic murine antibodies to DNA: antigen binding and structural characteristics. <i>International Immunology</i> , 1994, 6, 817-830.	1.8	145

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19	A stop codon polymorphism of Toll-like receptor 5 is associated with resistance to systemic lupus erythematosus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 10593-10597.	3.3	144
20	A missense variant in NCF1 is associated with susceptibility to multiple autoimmune diseases. <i>Nature Genetics</i> , 2017, 49, 433-437.	9.4	143
21	Update on human systemic lupus erythematosus genetics. <i>Current Opinion in Rheumatology</i> , 2004, 16, 513-521.	2.0	139
22	Genome-Wide Association Study in an Amerindian Ancestry Population Reveals Novel Systemic Lupus Erythematosus Risk Loci and the Role of European Admixture. <i>Arthritis and Rheumatology</i> , 2016, 68, 932-943.	2.9	138
23	Association of a functional IRF7 variant with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2011, 63, 749-754.	6.7	118
24	Recent insights into the genetic basis of systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2013, 72, ii56-ii61.	0.5	117
25	Lupus Nephritis Susceptibility Loci in Women with Systemic Lupus Erythematosus. <i>Journal of the American Society of Nephrology: JASN</i> , 2014, 25, 2859-2870.	3.0	117
26	X Chromosome Dose and Sex Bias in Autoimmune Diseases: Increased Prevalence of 47,XXX in Systemic Lupus Erythematosus and Sjögren's Syndrome. <i>Arthritis and Rheumatology</i> , 2016, 68, 1290-1300.	2.9	114
27	Regulatory polymorphisms modulate the expression of HLA class II molecules and promote autoimmunity. <i>ELife</i> , 2016, 5, .	2.8	113
28	Phenotypic associations of genetic susceptibility loci in systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 1752-1757.	0.5	110
29	Identification of novel genetic susceptibility loci in African American lupus patients in a candidate gene association study. <i>Arthritis and Rheumatism</i> , 2011, 63, 3493-3501.	6.7	109
30	Admixture Mapping in Lupus Identifies Multiple Functional Variants within IFIH1 Associated with Apoptosis, Inflammation, and Autoantibody Production. <i>PLoS Genetics</i> , 2013, 9, e1003222.	1.5	107
31	MicroRNA-3148 Modulates Allelic Expression of Toll-Like Receptor 7 Variant Associated with Systemic Lupus Erythematosus. <i>PLoS Genetics</i> , 2013, 9, e1003336.	1.5	107
32	A loss-of-function variant of PTPN22 is associated with reduced risk of systemic lupus erythematosus. <i>Human Molecular Genetics</i> , 2008, 18, 569-579.	1.4	106
33	Advances in lupus genetics and epigenetics. <i>Current Opinion in Rheumatology</i> , 2014, 26, 482-492.	2.0	104
34	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. <i>Arthritis and Rheumatism</i> , 2001, 44, 432-441.	6.7	103
35	Updates in Lupus Genetics. <i>Current Rheumatology Reports</i> , 2017, 19, 68.	2.1	99
36	PARP alleles within the linked chromosomal region are associated with systemic lupus erythematosus. <i>Journal of Clinical Investigation</i> , 1999, 103, 1135-1140.	3.9	99

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37	Fine mapping of Xq28: both <i>MECP2</i> and <i>IRAK1</i> contribute to risk for systemic lupus erythematosus in multiple ancestral groups. <i>Annals of the Rheumatic Diseases</i> , 2013, 72, 437-444.	0.5	97
38	Abnormal distribution of Fc $\gamma$ receptor type IIa polymorphisms in Korean patients with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 1998, 41, 421-426.	6.7	92
39	Inhibition of Aberrant Circulating Tfh Cell Proportions by Corticosteroids in Patients with Systemic Lupus Erythematosus. <i>PLoS ONE</i> , 2012, 7, e51982.	1.1	91
40	Identification of a Systemic Lupus Erythematosus Risk Locus Spanning <i>ATG16L2</i> , <i>FCHSD2</i> , and <i>P2RY2</i> in Koreans. <i>Arthritis and Rheumatology</i> , 2016, 68, 1197-1209.	2.9	89
41	Analysis of autosomal genes reveals gene $\times$ sex interactions and higher total genetic risk in men with systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 694-699.	0.5	87
42	Association analysis of the R620W polymorphism of protein tyrosine phosphatase <i>PTPN22</i> in systemic lupus erythematosus families: Increased t allele frequency in systemic lupus erythematosus patients with autoimmune thyroid disease. <i>Arthritis and Rheumatism</i> , 2005, 52, 2396-2402.	6.7	80
43	The <i>IRF5</i> $\times$ <i>TNPO3</i> association with systemic lupus erythematosus has two components that other autoimmune disorders variably share. <i>Human Molecular Genetics</i> , 2015, 24, 582-596.	1.4	74
44	Identification of a Systemic Lupus Erythematosus Susceptibility Locus at 11p13 between <i>PDHX</i> and <i>CD44</i> in a Multiethnic Study. <i>American Journal of Human Genetics</i> , 2011, 88, 83-91.	2.6	72
45	A peptide derived from an autoantibody can stimulate t cells in the (nzb $\times$ nzw)F1 mouse model of systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 1993, 36, 355-364.	6.7	70
46	Impact of genetic ancestry and sociodemographic status on the clinical expression of systemic lupus erythematosus in American Indian $\times$ European populations. <i>Arthritis and Rheumatism</i> , 2012, 64, 3687-3694.	6.7	70
47	<i>ABIN1</i> Dysfunction as a Genetic Basis for Lupus Nephritis. <i>Journal of the American Society of Nephrology: JASN</i> , 2013, 24, 1743-1754.	3.0	70
48	Association of two independent functional risk haplotypes in <i>TNIP1</i> with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2012, 64, 3695-3705.	6.7	69
49	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. <i>Arthritis and Rheumatism</i> , 2004, 50, 3203-3210.	6.7	66
50	In vivofunctional analysis of in vitro protein binding sites in the immunoglobulin heavy chain enhancer. <i>Nucleic Acids Research</i> , 1988, 16, 3239-3253.	6.5	64
51	Association of a common complement receptor 2 haplotype with increased risk of systemic lupus erythematosus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 3961-3966.	3.3	62
52	<i>ApoE</i> $\times$ <i>Fas</i> C57BL/6 mice: a novel murine model simultaneously exhibits lupus nephritis, atherosclerosis, and osteopenia. <i>Journal of Lipid Research</i> , 2007, 48, 794-805.	2.0	62
53	Novel identification of the <i>IRF7</i> region as an anticentromere autoantibody propensity locus in systemic sclerosis. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 114-119.	0.5	62
54	Current topics in human SLE genetics. <i>Seminars in Immunopathology</i> , 2006, 28, 97-107.	4.0	61

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55	Variation in the <i>ICAM1</i> – <i>ICAM4</i> – <i>ICAM5</i> locus is associated with systemic lupus erythematosus susceptibility in multiple ancestries. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 1809-1814.	0.5	60
56	Two Functional Lupus-Associated BLK Promoter Variants Control Cell-Type- and Developmental-Stage-Specific Transcription. <i>American Journal of Human Genetics</i> , 2014, 94, 586-598.	2.6	59
57	PTPN22 Association in Systemic Lupus Erythematosus (SLE) with Respect to Individual Ancestry and Clinical Sub-Phenotypes. <i>PLoS ONE</i> , 2013, 8, e69404.	1.1	57
58	Linkage and interaction of loci on 1q23 and 16q12 may contribute to susceptibility to systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2002, 46, 2928-2936.	6.7	55
59	Association of IRF5 polymorphisms with activation of the interferon $\lambda$ pathway. <i>Annals of the Rheumatic Diseases</i> , 2010, 69, 611-617.	0.5	54
60	Evidence for gene–gene epistatic interactions among susceptibility loci for systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2012, 64, 485-492.	6.7	53
61	Familiality and co-occurrence of clinical features of systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2002, 46, 2678-2685.	6.7	51
62	Evaluation of <i>TRAF6</i> in a large multi-ancestral lupus cohort. <i>Arthritis and Rheumatism</i> , 2012, 64, 1960-1969.	6.7	51
63	Trans-Ancestral Studies Fine Map the SLE-Susceptibility Locus TNFSF4. <i>PLoS Genetics</i> , 2013, 9, e1003554.	1.5	50
64	Association of tumor necrosis factor $\lambda$ polymorphism, but not the shared epitope, with increased radiographic progression in a seropositive rheumatoid arthritis inception cohort. <i>Arthritis and Rheumatism</i> , 2006, 54, 1105-1116.	6.7	49
65	B cells are anergic in transgenic mice that express IgM anti-DNA antibodies. <i>European Journal of Immunology</i> , 1993, 23, 2332-2339.	1.6	48
66	Treatment with apolipoprotein A-1 mimetic peptide reduces lupus-like manifestations in a murine lupus model of accelerated atherosclerosis. <i>Arthritis Research and Therapy</i> , 2010, 12, R93.	1.6	47
67	Genetics of systemic lupus erythematosus: immune responses and end organ resistance to damage. <i>Current Opinion in Immunology</i> , 2014, 31, 87-96.	2.4	47
68	An update on genetic studies of systemic lupus erythematosus. <i>Current Rheumatology Reports</i> , 2002, 4, 359-367.	2.1	46
69	Genetic contributions to lupus nephritis in a multi-ethnic cohort of systemic lupus erythematosus patients. <i>PLoS ONE</i> , 2018, 13, e0199003.	1.1	46
70	Genetic analyses of interferon pathway-related genes reveal multiple new loci associated with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2011, 63, 2049-2057.	6.7	45
71	CD72 polymorphisms associated with alternative splicing modify susceptibility to human systemic lupus erythematosus through epistatic interaction with FCGR2B. <i>Human Molecular Genetics</i> , 2004, 13, 2907-2917.	1.4	43
72	Identification of interferon-inducible genes as diagnostic biomarker for systemic lupus erythematosus. <i>Clinical Rheumatology</i> , 2015, 34, 71-79.	1.0	43

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73	Genetic fine mapping of systemic lupus erythematosus MHC associations in Europeans and African Americans. <i>Human Molecular Genetics</i> , 2018, 27, 3813-3824.	1.4	43
74	Interaction between RANKL and HLA-DRB1 genotypes may contribute to younger age at onset of seropositive rheumatoid arthritis in an inception cohort. <i>Arthritis and Rheumatism</i> , 2004, 50, 3093-3103.	6.7	42
75	Modulation of IL-6 induced RANKL expression in arthritic synovium by a transcription factor SOX5. <i>Scientific Reports</i> , 2016, 6, 32001.	1.6	41
76	Maternal HLA class II compatibility in men with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2005, 52, 2768-2773.	6.7	36
77	Association of <i>PPP2CA</i> polymorphisms with systemic lupus erythematosus susceptibility in multiple ethnic groups. <i>Arthritis and Rheumatism</i> , 2011, 63, 2755-2763.	6.7	36
78	Preferential Binding to Elk-1 by SLE-Associated IL10 Risk Allele Upregulates IL10 Expression. <i>PLoS Genetics</i> , 2013, 9, e1003870.	1.5	36
79	Lupus Risk Variant Increases pSTAT1 Binding and Decreases ETS1 Expression. <i>American Journal of Human Genetics</i> , 2015, 96, 731-739.	2.6	36
80	Amino acid signatures of HLA Class-I and II molecules are strongly associated with SLE susceptibility and autoantibody production in Eastern Asians. <i>PLoS Genetics</i> , 2019, 15, e1008092.	1.5	36
81	Current advances in the human lupus genetics. <i>Current Rheumatology Reports</i> , 2004, 6, 391-398.	2.1	35
82	A functional <i>RANKL</i> polymorphism associated with younger age at onset of rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 2010, 62, 2864-2875.	6.7	35
83	Restored Immunosuppressive Effect of Mesenchymal Stem Cells on B Cells After Olfactory 1/Early B Cell Factor-Associated Zinc Finger Protein Down-Regulation in Patients With Systemic Lupus Erythematosus. <i>Arthritis and Rheumatology</i> , 2014, 66, 3413-3423.	2.9	35
84	Association of Fc $\gamma$ receptor IIA, but not IIB and IIIA, polymorphisms with systemic lupus erythematosus: A family-based association study in Caucasians. <i>Arthritis and Rheumatism</i> , 2004, 50, 671-673.	6.7	34
85	A plausibly causal functional lupus-associated risk variant in the STAT1-STAT4 locus. <i>Human Molecular Genetics</i> , 2018, 27, 2392-2404.	1.4	34
86	Transcription Factor SOX5 Promotes the Migration and Invasion of Fibroblast-Like Synoviocytes in Part by Regulating MMP-9 Expression in Collagen-Induced Arthritis. <i>Frontiers in Immunology</i> , 2018, 9, 749.	2.2	33
87	Plasma levels of osteopontin identify patients at risk for organ damage in systemic lupus erythematosus. <i>Arthritis Research and Therapy</i> , 2013, 15, R18.	1.6	32
88	Reduced Let-7f in Bone Marrow-Derived Mesenchymal Stem Cells Triggers Treg/Th17 Imbalance in Patients With Systemic Lupus Erythematosus. <i>Frontiers in Immunology</i> , 2020, 11, 233.	2.2	30
89	Single-nucleotide polymorphisms of T cell receptor $\gamma$ chain in patients with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 1999, 42, 2601-2605.	6.7	29
90	European population substructure is associated with mucocutaneous manifestations and autoantibody production in systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2009, 60, 2448-2456.	6.7	27

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91	Lupus risk variants in the PXX locus alter B-cell receptor internalization. <i>Frontiers in Genetics</i> , 2015, 5, 450.	1.1	25
92	Genetic variants in systemic lupus erythematosus susceptibility loci, XKR6 and GLT1D1 are associated with childhood-onset SLE in a Korean cohort. <i>Scientific Reports</i> , 2018, 8, 9962.	1.6	25
93	Genetics and systemic lupus erythematosus. <i>Current Rheumatology Reports</i> , 2001, 3, 183-190.	2.1	22
94	Transcription factor Ikaros Represses Protein Phosphatase 2A (PP2A) Expression through an Intronic Binding Site. <i>Journal of Biological Chemistry</i> , 2014, 289, 13751-13757.	1.6	20
95	Prediction models of treatment response in lupus nephritis. <i>Kidney International</i> , 2022, 101, 379-389.	2.6	18
96	Fc $\gamma$ 3 receptor IIIA polymorphism in Korean patients with systemic lupus erythematosus. <i>Rheumatology International</i> , 2002, 21, 222-226.	1.5	17
97	Decreased <i>SMG7</i> expression associates with lupus-risk variants and elevated antinuclear antibody production. <i>Annals of the Rheumatic Diseases</i> , 2016, 75, 2007-2013.	0.5	16
98	Variable Association of Reactive Intermediate Genes with Systemic Lupus Erythematosus in Populations with Different African Ancestry. <i>Journal of Rheumatology</i> , 2013, 40, 842-849.	1.0	15
99	Genetics and systemic lupus erythematosus. <i>Current Rheumatology Reports</i> , 2000, 2, 13-18.	2.1	14
100	Identification and characterization of a peptide mimetic that may detect a species of disease-associated anticardiolipin antibodies in patients with the antiphospholipid syndrome. <i>Arthritis and Rheumatism</i> , 2003, 48, 737-745.	6.7	14
101	Human SLE variant <i>NCF1</i> -R90H promotes kidney damage and murine lupus through enhanced Tfh2 responses induced by defective efferocytosis of macrophages. <i>Annals of the Rheumatic Diseases</i> , 2022, 81, 255-267.	0.5	14
102	Male-only Systemic Lupus. <i>Journal of Rheumatology</i> , 2010, 37, 1480-1487.	1.0	13
103	Transcriptional effects of a lupus-associated polymorphism in the 5' untranslated region (UTR) of human complement receptor 2 (CR2/CD21). <i>Molecular Immunology</i> , 2012, 52, 165-173.	1.0	12
104	CD3Z hypermethylation is associated with severe clinical manifestations in systemic lupus erythematosus and reduces CD3Z-chain expression in T cells. <i>Rheumatology</i> , 2016, 56, kew405.	0.9	12
105	Rigorous Plasma Microbiome Analysis Method Enables Disease Association Discovery in Clinic. <i>Frontiers in Microbiology</i> , 2020, 11, 613268.	1.5	12
106	Upregulated Interleukin-10 Induced by E2F Transcription Factor 2 MicroRNA Circuitry in Extrafollicular Effector B Cells Contributes to Autoantibody Production in Systemic Lupus Erythematosus. <i>Arthritis and Rheumatology</i> , 2022, 74, 496-507.	2.9	12
107	Commentary: Genetics of systemic lupus erythematosus. <i>Current Opinion in Rheumatology</i> , 1997, 9, 377-379.	2.0	10
108	Poly(ADP-ribose) polymerase and susceptibility to systemic lupus erythematosus and primary antiphospholipid syndrome: Comment on the article by Delrieu et al. <i>Arthritis and Rheumatism</i> , 2000, 43, 1421-1422.	6.7	10

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109	Brief Report: Single nucleotide polymorphisms in <i>VKORC1</i> are risk factors for systemic lupus erythematosus in Asians. <i>Arthritis and Rheumatism</i> , 2013, 65, 211-215.	6.7	10
110	Genetic associations of leptin-related polymorphisms with systemic lupus erythematosus. <i>Clinical Immunology</i> , 2015, 161, 157-162.	1.4	10
111	Preferential association of a functional variant in complement receptor 2 with antibodies to double-stranded DNA. <i>Annals of the Rheumatic Diseases</i> , 2016, 75, 242-252.	0.5	10
112	RNASE2 Mediates Age-Associated B Cell Expansion Through Monocyte Derived IL-10 in Patients With Systemic Lupus Erythematosus. <i>Frontiers in Immunology</i> , 2022, 13, 752189.	2.2	9
113	T cell up-regulation of B cells via their idiotypes contributing to the development of systemic lupus erythematosus. <i>American Journal of Medicine</i> , 1988, 85, 32-34.	0.6	8
114	Olf1/EBF associated zinc finger protein interfered with antinuclear antibody production in patients with systemic lupus erythematosus. <i>Arthritis Research and Therapy</i> , 2010, 12, R59.	1.6	8
115	Deep sequencing reveals a DAP1 regulatory haplotype that potentiates autoimmunity in systemic lupus erythematosus. <i>Genome Biology</i> , 2020, 21, 281.	3.8	8
116	Membrane association and orientation of rat renal activities capable of degrading glutathione. <i>International Journal of Biochemistry &amp; Cell Biology</i> , 1980, 12, 219-222.	0.8	7
117	The role of cytoplasmic free calcium concentration in B-cell tolerance. <i>Cellular Immunology</i> , 1987, 108, 335-342.	1.4	7
118	Central suppression of monoclonal B cells: DNP-MGG suppresses proliferation and immunoglobulin synthesis in anti-DNP-secreting hybridoma and myeloma. <i>Cellular Immunology</i> , 1984, 88, 96-108.	1.4	6
119	Evidence that the hydrophobic domain of rat renal $\Gamma^3$ -glutamyltransferase spans the brush border membrane. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1982, 690, 199-206.	1.4	5
120	Genetics of Human SLE. , 2019, , 54-68.		5
121	Constitutive Genes and Lupus. , 2011, , 47-61.		4
122	Genetics of Human SLE. , 2013, , 35-45.		4
123	Pathogenesis of Systemic Lupus Erythematosus. , 2009, , 1233-1262.		4
124	Complement <i>C4</i> , the Major Histocompatibility Complex, and Autoimmunity. <i>Arthritis and Rheumatology</i> , 2022, 74, 1318-1320.	2.9	4
125	Contribution of Major Histocompatibility Complex (MHC) to Upregulation of Anti-DNA Antibody in Transgenic Mice. <i>Journal of Autoimmunity</i> , 1993, 6, 1-9.	3.0	3
126	Focused transcription from the human CR2/CD21 core promoter is regulated by synergistic activity of TATA and Initiator elements in mature B cells. <i>Cellular and Molecular Immunology</i> , 2016, 13, 119-131.	4.8	3



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127	Ig-transgenic mice as models for studying the regulation and role of anti-DNA antibodies in murine lupus. <i>ImmunoMethods</i> , 1992, 1, 185-190.	0.8	2
128	Plasmin immunization preferentially induces potentially prothrombotic IgG anticardiolipin antibodies in MRL/MpJ mice. <i>Arthritis and Rheumatism</i> , 2009, 60, 3108-3117.	6.7	2
129	Genes and genetics in human SLE. , 2021, , 85-96.		2
130	IFNL4 Genotype Does Not Associate with CD4 T-Cell Recovery in People Living with Human Immunodeficiency Virus. <i>AIDS Research and Human Retroviruses</i> , 2021, 37, 184-188.	0.5	2
131	Idiotype selection is an immunoregulatory mechanism which contributes to the pathogenesis of systemic lupus erythematosus. <i>Journal of Autoimmunity</i> , 1988, 1, 673-681.	3.0	1
132	Genes and Genetics in Human Systemic Lupus Erythematosus. , 2016, , 69-76.		1
133	Macrophage-derived soluble factors mediate suppression induced by 2,4-dinitrophenyl-conjugated mouse IgG in hybridoma cells. <i>Cellular Immunology</i> , 1985, 91, 362-374.	1.4	0
134	Autoimmunity and Tolerance in Ig-Transgenic Mice: Murine SLE as a Model to Study B Cell Tolerance. <i>International Reviews of Immunology</i> , 1994, 11, 305-320.	1.5	0
135	CG-07â€¦Regulatory polymorphisms in EMSY gene are associated with autoantibodies in healthy individuals. , 2018, , .		0
136	180â€¦Examining the transcriptional impact of liganded estrogen receptor alpha in the inflammatory milieu of systemic lupus erythematosus. , 2019, , .		0
137	Lupus susceptibility genes. , 2021, , 25-33.		0
138	A genome wide association study of systemic lupus erythematosus (SLE) by SLEGEN, the International SLE Genetics Consortium.. <i>FASEB Journal</i> , 2008, 22, 850.1.	0.2	0
139	Systemic Lupus Erythematosus, <i>Genetics</i> . , 2014, , 1171-1178.		0
140	Autoantibodies as a Source of Peptides That Regulate Autoantibody Production. , 1999, , 371-388.		0