

# Jennifer A Kelly

## List of Publications by Year in descending order

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119  
papers

10,127  
citations

28736

57  
h-index

40945

97  
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121  
all docs

121  
docs citations

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times ranked

11144  
citing authors

#	ARTICLE	IF	CITATIONS
1	Variants on the <i>UBE2L3</i> / <i>YDJC</i> Autoimmune Disease Risk Haplotype Increase <i>UBE2L3</i> Expression by Modulating <i>CCCTC</i> Binding Factor and YY1 Binding. <i>Arthritis and Rheumatology</i> , 2022, 74, 163-173.	2.9	4
2	Relationship Between a Vitamin D Genetic Risk Score and Autoantibodies Among First-Degree Relatives of Probands With Rheumatoid Arthritis and Systemic Lupus Erythematosus. <i>Frontiers in Immunology</i> , 2022, 13, .	2.2	5
3	1508...Single-cell epigenetic profiling highlights genetic impact on chromatin accessibility in SLE. , 2021, , .		0
4	Nucleic Acid-Sensing and Interferon-Inducible Pathways Show Differential Methylation in MZ Twins Discordant for Lupus and Overexpression in Independent Lupus Samples: Implications for Pathogenic Mechanism and Drug Targeting. <i>Genes</i> , 2021, 12, 1898.	1.0	6
5	Novel genetic associations with interferon in systemic lupus erythematosus identified by replication and fine-mapping of trait-stratified genome-wide screen. <i>Cytokine</i> , 2020, 132, 154631.	1.4	19
6	American Indians Have a Higher Risk of Sjögren's Syndrome and More Disease Activity Than European Americans and African Americans. <i>Arthritis Care and Research</i> , 2020, 72, 1049-1056.	1.5	14
7	Role of Systemic Lupus Erythematosus Risk Variants With Opposing Functional Effects as a Driver of Hypomorphic Expression of TNIP 1 and Other Genes Within a Three-Dimensional Chromatin Network. <i>Arthritis and Rheumatology</i> , 2020, 72, 780-790.	2.9	9
8	Deep sequencing reveals a DAP1 regulatory haplotype that potentiates autoimmunity in systemic lupus erythematosus. <i>Genome Biology</i> , 2020, 21, 281.	3.8	8
9	Genetic variants at the 16p13 locus confer risk for eosinophilic esophagitis. <i>Genes and Immunity</i> , 2019, 20, 281-292.	2.2	30
10	Association of Epstein-Barr virus serological reactivation with transitioning to systemic lupus erythematosus in at-risk individuals. <i>Annals of the Rheumatic Diseases</i> , 2019, 78, 1235-1241.	0.5	64
11	Latent autoimmunity across disease-specific boundaries in at-risk first-degree relatives of SLE and RA patients. <i>EBioMedicine</i> , 2019, 42, 76-85.	2.7	18
12	A plausibly causal functional lupus-associated risk variant in the <i>STAT1</i> STAT4 locus. <i>Human Molecular Genetics</i> , 2018, 27, 2392-2404.	1.4	34
13	Enhancer histone-QTLs are enriched on autoimmune risk haplotypes and influence gene expression within chromatin networks. <i>Nature Communications</i> , 2018, 9, 2905.	5.8	56
14	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
15	A missense variant in <i>NCF1</i> is associated with susceptibility to multiple autoimmune diseases. <i>Nature Genetics</i> , 2017, 49, 433-437.	9.4	143
16	Combined role of vitamin D status and <i>CYP24A1</i> in the transition to systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2017, 76, 153-158.	0.5	40
17	Brief Report: Association of Natural Killer Cell Ligand Polymorphism HLA-C Asn80Lys With the Development of Anti-SSA/Ro-Associated Congenital Heart Block. <i>Arthritis and Rheumatology</i> , 2017, 69, 2170-2174.	2.9	11
18	Transancestral mapping and genetic load in systemic lupus erythematosus. <i>Nature Communications</i> , 2017, 8, 16021.	5.8	314

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19	Brief Report: Rare X Chromosome Abnormalities in Systemic Lupus Erythematosus and Sjögren's Syndrome. <i>Arthritis and Rheumatology</i> , 2017, 69, 2187-2192.	2.9	35
20	Association of IFIH1 and pro-inflammatory mediators: Potential new clues in SLE-associated pathogenesis. <i>PLoS ONE</i> , 2017, 12, e0171193.	1.1	11
21	Identification of a Sjögren's syndrome susceptibility locus at OAS1 that influences isoform switching, protein expression, and responsiveness to type I interferons. <i>PLoS Genetics</i> , 2017, 13, e1006820.	1.5	60
22	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
23	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
24	Single-cell analysis of glandular T cell receptors in Sjögren's syndrome. <i>JCI Insight</i> , 2016, 1, .	2.3	54
25	Regulatory polymorphisms modulate the expression of HLA class II molecules and promote autoimmunity. <i>ELife</i> , 2016, 5, .	2.8	113
26	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
27	The anti-inflammatory <i>CASPASE-12</i> gene does not influence SLE phenotype in African-Americans. <i>Immunology Letters</i> , 2016, 173, 21-25.	1.1	1
28	Klinefelter's syndrome (47,XXY) is in excess among men with Sjögren's syndrome. <i>Clinical Immunology</i> , 2016, 168, 25-29.	1.4	68
29	Identification of a New Susceptibility Locus for Systemic Lupus Erythematosus on Chromosome 12 in Individuals of European Ancestry. <i>Arthritis and Rheumatology</i> , 2016, 68, 174-183.	2.9	30
30	Development and validation of a simple lupus severity index using ACR criteria for classification of SLE. <i>Lupus Science and Medicine</i> , 2016, 3, e000136.	1.1	25
31	Decreased <i>SMC7</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
32	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
33	Disease Mechanisms in Rheumatology—Tools and Pathways: Defining Functional Genetic Variants in Autoimmune Diseases. <i>Arthritis and Rheumatology</i> , 2015, 67, 1-10.	2.9	22
34	Lupus risk variants in the <i>PXK</i> locus alter B-cell receptor internalization. <i>Frontiers in Genetics</i> , 2015, 5, 450.	1.1	25
35	Lupus Risk Variant Increases pSTAT1 Binding and Decreases ETS1 Expression. <i>American Journal of Human Genetics</i> , 2015, 96, 731-739.	2.6	36
36	Genetic associations of leptin-related polymorphisms with systemic lupus erythematosus. <i>Clinical Immunology</i> , 2015, 161, 157-162.	1.4	10

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37	The IRF5-TNPO3 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
38	Genetics of lupus. , 2015, , 1045-1051.		4
39	Use of next-generation DNA sequencing to analyze genetic variants in rheumatic disease. <i>Arthritis Research and Therapy</i> , 2014, 16, 490.	1.6	13
40	Allelic heterogeneity in NCF2 associated with systemic lupus erythematosus (SLE) susceptibility across four ethnic populations. <i>Human Molecular Genetics</i> , 2014, 23, 1656-1668.	1.4	67
41	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
42	Comparison of the American-European Consensus Group Sjögren's syndrome classification criteria to newly proposed American College of Rheumatology criteria in a large, carefully characterised sicca cohort. <i>Annals of the Rheumatic Diseases</i> , 2014, 73, 31-38.	0.5	161
43	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
44	Variants at multiple loci implicated in both innate and adaptive immune responses are associated with Sjögren's syndrome. <i>Nature Genetics</i> , 2013, 45, 1284-1292.	9.4	427
45	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
46	Preferential Binding to Elk-1 by SLE-Associated IL10 Risk Allele Upregulates IL10 Expression. <i>PLoS Genetics</i> , 2013, 9, e1003870.	1.5	36
47	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
48	Trans-Ancestral Studies Fine Map the SLE-Susceptibility Locus TNFSF4. <i>PLoS Genetics</i> , 2013, 9, e1003554.	1.5	50
49	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
50	Fine mapping of Xq28: both MECP2 and IRAK1 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
51	ABIN1 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
52	Familial Aggregation of High Tumor Necrosis Factor Alpha Levels in Systemic Lupus Erythematosus. <i>Clinical and Developmental Immunology</i> , 2013, 2013, 1-6.	3.3	12
53	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
54	Multiple Autoantibodies Display Association with Lymphopenia, Proteinuria, and Cellular Casts in a Large, Ethnically Diverse SLE Patient Cohort. <i>Autoimmune Diseases</i> , 2012, 2012, 1-11.	2.7	21

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55	Effects of IRF5 Lupus Risk Haplotype on Pathways Predicted to Influence B Cell Functions. <i>Journal of Biomedicine and Biotechnology</i> , 2012, 2012, 1-12.	3.0	8
56	Genetic and physical interaction of the B-cell systemic lupus erythematosus-associated genes <i>BANK1</i> and <i>BLK</i> . <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 136-142.	0.5	67
57	Analysis of autosomal genes reveals gene-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
58	IRF5 haplotypes demonstrate diverse serological associations which predict serum interferon alpha activity and explain the majority of the genetic association with systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 463-469.	0.5	127
59	Impact of genetic ancestry and sociodemographic status on the clinical expression of systemic lupus erythematosus in American Indian-European populations. <i>Arthritis and Rheumatism</i> , 2012, 64, 3687-3694.	6.7	70
60	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
61	Comparison of autoantibody specificities between traditional and bead-based assays in a large, diverse collection of patients with systemic lupus erythematosus and family members. <i>Arthritis and Rheumatism</i> , 2012, 64, 3677-3686.	6.7	72
62	Variation in the <i>ICAM1-ICAM4-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
63	Genetics of Sjögren's syndrome in the genome-wide association era. <i>Journal of Autoimmunity</i> , 2012, 39, 57-63.	3.0	61
64	Brief Report: Large-scale analysis of tumor necrosis factor $\pm$ levels in systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2012, 64, 2947-2952.	6.7	76
65	Genome-Wide Association Study of African and European Americans Implicates Multiple Shared and Ethnic Specific Loci in Sarcoidosis Susceptibility. <i>PLoS ONE</i> , 2012, 7, e43907.	1.1	105
66	Evaluation of <i>TRAF6</i> in a large multiancestral lupus cohort. <i>Arthritis and Rheumatism</i> , 2012, 64, 1960-1969.	6.7	51
67	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
68	The genomics of autoimmune disease in the era of genome-wide association studies and beyond. <i>Autoimmunity Reviews</i> , 2012, 11, 267-275.	2.5	58
69	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
70	Association of a functional variant downstream of TNFAIP3 with systemic lupus erythematosus. <i>Nature Genetics</i> , 2011, 43, 253-258.	9.4	242
71	Identification of a Systemic Lupus Erythematosus Susceptibility Locus at 11p13 between PDHX and CD44 in a Multiethnic Study. <i>American Journal of Human Genetics</i> , 2011, 88, 83-91.	2.6	72
72	Network analysis of associations between serum interferon $\pm$ activity, autoantibodies, and clinical features in systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2011, 63, 1044-1053.	6.7	222

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73	Fine-mapping and transethnic genotyping establish IL2/IL21 genetic association with lupus and localize this genetic effect to IL21. <i>Arthritis and Rheumatism</i> , 2011, 63, 1689-1697.	6.7	49
74	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
75	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
76	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
77	Early disease onset is predicted by a higher genetic risk for lupus and is associated with a more severe phenotype in lupus patients. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 151-156.	0.5	155
78	The Lupus Family Registry and Repository. <i>Rheumatology</i> , 2011, 50, 47-59.	0.9	82
79	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
80	Phenotypic associations of genetic susceptibility loci in systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 1752-1757.	0.5	110
81	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
82	Analysis of maternal-offspring HLA compatibility, parent-of-origin effects, and noninherited maternal antigen effects for HLA-DRB1 in systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2010, 62, 1712-1717.	6.7	8
83	Genetically determined Amerindian ancestry correlates with increased frequency of risk alleles for systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2010, 62, 3722-3729.	6.7	70
84	Identification of Unique MicroRNA Signature Associated with Lupus Nephritis. <i>PLoS ONE</i> , 2010, 5, e10344.	1.1	187
85	The Role of Genetic Variation Near Interferon-Kappa in Systemic Lupus Erythematosus. <i>Journal of Biomedicine and Biotechnology</i> , 2010, 2010, 1-11.	3.0	44
86	Polymorphisms in the Hsp70 gene locus are genetically associated with systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2010, 69, 1983-1989.	0.5	32
87	Identification of <i>IRAK1</i> as a risk gene with critical role in the pathogenesis of systemic lupus erythematosus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 6256-6261.	3.3	218
88	High-Density SNP Screening of the Major Histocompatibility Complex in Systemic Lupus Erythematosus Demonstrates Strong Evidence for Independent Susceptibility Regions. <i>PLoS Genetics</i> , 2009, 5, e1000696.	1.5	109
89	Variants within <i>MECP2</i> , a key transcription regulator, are associated with increased susceptibility to lupus and differential gene expression in patients with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2009, 60, 1076-1084.	6.7	80
90	High-density genotyping of STAT4 reveals multiple haplotypic associations with systemic lupus erythematosus in different racial groups. <i>Arthritis and Rheumatism</i> , 2009, 60, 1085-1095.	6.7	82

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91	A polymorphism within <i>IL21R</i> confers risk for systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2009, 60, 2402-2407.	6.7	108
92	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
93	Genetic susceptibility to SLE: new insights from fine mapping and genome-wide association studies. <i>Nature Reviews Genetics</i> , 2009, 10, 285-290.	7.7	251
94	Genome-wide linkage screen for stature and body mass index in 3,032 families: evidence for sex- and population-specific genetic effects. <i>European Journal of Human Genetics</i> , 2009, 17, 258-266.	1.4	16
95	Kallikrein genes are associated with lupus and glomerular basement membrane-specific antibody-induced nephritis in mice and humans. <i>Journal of Clinical Investigation</i> , 2009, 119, 911-923.	3.9	114
96	Association of the IRF5 risk haplotype with high serum interferon activity in systemic lupus erythematosus patients. <i>Arthritis and Rheumatism</i> , 2008, 58, 2481-2487.	6.7	246
97	Klinefelter's syndrome (47,XXY) in male systemic lupus erythematosus patients: Support for the notion of a gene-dose effect from the X chromosome. <i>Arthritis and Rheumatism</i> , 2008, 58, 2511-2517.	6.7	308
98	A nonsynonymous functional variant in integrin- $\beta$ M (encoded by ITGAM) is associated with systemic lupus erythematosus. <i>Nature Genetics</i> , 2008, 40, 152-154.	9.4	277
99	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
100	Addressing genetic heterogeneity in complex disease: Finding seizure genes in systemic lupus erythematosus. <i>Epilepsia</i> , 2008, 49, 527-530.	2.6	14
101	Common Variants within MECP2 Confer Risk of Systemic Lupus Erythematosus. <i>PLoS ONE</i> , 2008, 3, e1727.	1.1	125
102	The Genetics of Lupus. , 2007, , 74-86.		4
103	Localization and replication of the systemic lupus erythematosus linkage signal at 4p16: interaction with 2p11, 12q24 and 19q13 in European Americans. <i>Human Genetics</i> , 2006, 120, 623-631.	1.8	20
104	Unraveling the genetics of systemic lupus erythematosus. <i>Seminars in Immunopathology</i> , 2006, 28, 119-130.	4.0	127
105	Evaluation of the genetic association of the PTPN22 R620W polymorphism in familial and sporadic systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2006, 54, 2533-2540.	6.7	43
106	Linkage at 5q14.3-15 in multiplex systemic lupus erythematosus pedigrees stratified by autoimmune thyroid disease. <i>Arthritis and Rheumatism</i> , 2005, 52, 3646-3650.	6.7	34
107	Genetic linkage of systemic lupus erythematosus to 13q32 in African American families with affected male members. <i>Human Genetics</i> , 2005, 118, 309-321.	1.8	15
108	Mapping the Systematic Lupus Erythematosus Susceptibility Genes. , 2004, 102, 011-030.		5

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109	SLEN2 (2q34-35) and SLEN1 (10q22.3) Replication in Systemic Lupus Erythematosus Stratified by Nephritis. <i>American Journal of Human Genetics</i> , 2004, 75, 346-348.	2.6	20
110	Thrombocytopenia identifies a severe familial phenotype of systemic lupus erythematosus and reveals genetic linkages at 1q22 and 11p13. <i>Blood</i> , 2003, 101, 992-997.	0.6	67
111	Evidence for a susceptibility gene (SLEH1) on chromosome 11q14 for systemic lupus erythematosus (SLE) families with hemolytic anemia. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 11766-11771.	3.3	35
112	Genetic linkage and association of Fc $\gamma$ receptor IIIA (CD16A) on chromosome 1q23 with human systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2002, 46, 2132-2140.	6.7	127
113	Stratification of pedigrees multiplex for systemic lupus erythematosus and for self-reported rheumatoid arthritis detects a systemic lupus erythematosus susceptibility gene (SLER1) at 5p15.3. <i>Arthritis and Rheumatism</i> , 2002, 46, 2937-2945.	6.7	36
114	SLEB3 in systemic lupus erythematosus (SLE) is strongly related to SLE families ascertained through neuropsychiatric manifestations. <i>Human Genetics</i> , 2002, 111, 54-58.	1.8	30
115	Genetic basis of systemic lupus erythematosus: a review of the unique genetic contributions in African Americans. <i>Journal of the National Medical Association</i> , 2002, 94, 670-7.	0.6	12
116	Evidence for a Susceptibility Gene, SLEV1, on Chromosome 17p13 in Families with Vitiligo-Related Systemic Lupus Erythematosus. <i>American Journal of Human Genetics</i> , 2001, 69, 1401-1406.	2.6	124
117	Linkage analysis of angiotensin-converting enzyme (ACE) insertion/deletion polymorphism and systemic lupus erythematosus. <i>Molecular and Cellular Endocrinology</i> , 2001, 177, 81-85.	1.6	73
118	Linkage analysis of human systemic lupus erythematosus-related traits: A principal component approach. <i>Arthritis and Rheumatism</i> , 2001, 44, 2807-2818.	6.7	37
119	Confirmation of genetic linkage between human systemic lupus erythematosus and chromosome 1q41. <i>Arthritis and Rheumatism</i> , 1999, 42, 1902-1907.	6.7	64