

Nan Shen

List of Publications by Year in descending order

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Version: 2024-02-01

126
papers

9,965
citations

36303

51
h-index

38395

95
g-index

135
all docs

135
docs citations

135
times ranked

12878
citing authors

#	ARTICLE	IF	CITATIONS
1	The CD6/ALCAM pathway promotes lupus nephritis via T cell-mediated responses. <i>Journal of Clinical Investigation</i> , 2022, 132, .	8.2	25
2	P2RY8 variants in lupus patients uncover a role for the receptor in immunological tolerance. <i>Journal of Experimental Medicine</i> , 2022, 219, .	8.5	26
3	AKT2 reduces IFN γ production to modulate antiviral responses and systemic lupus erythematosus. <i>EMBO Journal</i> , 2022, 41, e108016.	7.8	5
4	Urinary galectin-3 binding protein (G3BP) as a biomarker for disease activity and renal pathology characteristics in lupus nephritis. <i>Arthritis Research and Therapy</i> , 2022, 24, 77.	3.5	4
5	Expanding Roles of Noncoding RNAs in the Pathogenesis of Systemic Lupus Erythematosus. <i>Current Rheumatology Reports</i> , 2022, 24, 64-75.	4.7	2
6	Lupus enhancer risk variant causes dysregulation of IRF8 through cooperative lncRNA and DNA methylation machinery. <i>Nature Communications</i> , 2022, 13, 1855.	12.8	16
7	TLR7 gain-of-function genetic variation causes human lupus. <i>Nature</i> , 2022, 605, 349-356.	27.8	208
8	Two Distinct Immune Cell Signatures Predict the Clinical Outcomes in Patients With Amyopathic Dermatomyositis With Interstitial Lung Disease. <i>Arthritis and Rheumatology</i> , 2022, 74, 1822-1832.	5.6	10
9	Biological insights into systemic lupus erythematosus through an immune cell-specific transcriptome-wide association study. <i>Annals of the Rheumatic Diseases</i> , 2022, 81, 1273-1280.	0.9	9
10	The NCF1 variant p.R90H aggravates autoimmunity by facilitating the activation of plasmacytoid dendritic cells. <i>Journal of Clinical Investigation</i> , 2022, 132, .	8.2	11
11	MicroRNA in systemic lupus erythematosus. , 2021, , 259-265.		0
12	Long non-coding RNA expression profiles in neutrophils revealed potential biomarker for prediction of renal involvement in SLE patients. <i>Rheumatology</i> , 2021, 60, 1734-1746.	1.9	16
13	Meta-analysis of 208370 East Asians identifies 113 susceptibility loci for systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2021, 80, 632-640.	0.9	103
14	SLE non-coding genetic risk variant determines the epigenetic dysfunction of an immune cell specific enhancer that controls disease-critical microRNA expression. <i>Nature Communications</i> , 2021, 12, 135.	12.8	48
15	Supranutritional selenium suppresses ROS-induced generation of RANKL-expressing osteoclastogenic CD4 ⁺ T cells and ameliorates rheumatoid arthritis. <i>Clinical and Translational Immunology</i> , 2021, 10, e1338.	3.8	7
16	Sustained low-dose interleukin-2 therapy alleviates pathogenic humoral immunity via elevating the Tfr/Tfh ratio in lupus. <i>Clinical and Translational Immunology</i> , 2021, 10, e1293.	3.8	16
17	Identification of 38 novel loci for systemic lupus erythematosus and genetic heterogeneity between ancestral groups. <i>Nature Communications</i> , 2021, 12, 772.	12.8	128
18	The metabolic hormone leptin promotes the function of TFH cells and supports vaccine responses. <i>Nature Communications</i> , 2021, 12, 3073.	12.8	27

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19	Bach2 attenuates IL-2R signaling to control Treg homeostasis and Tfr development. <i>Cell Reports</i> , 2021, 35, 109096.	6.4	14
20	Selenium-GPX4 axis protects follicular helper T cells from ferroptosis. <i>Nature Immunology</i> , 2021, 22, 1127-1139.	14.5	158
21	Glutamine metabolism is essential for the production of IL-17A in Th17 T cells and skin inflammation. <i>Tissue and Cell</i> , 2021, 71, 101569.	2.2	12
22	Downregulation of Renal Hsa-miR-127-3p Contributes to the Overactivation of Type I Interferon Signaling Pathway in the Kidney of Lupus Nephritis. <i>Frontiers in Immunology</i> , 2021, 12, 747616.	4.8	6
23	SARS-CoV-2-Encoded MiRNAs Inhibit Host Type I Interferon Pathway and Mediate Allelic Differential Expression of Susceptible Gene. <i>Frontiers in Immunology</i> , 2021, 12, 767726.	4.8	17
24	Efficacy and safety of low-dose IL-2 in the treatment of systemic lupus erythematosus: a randomised, double-blind, placebo-controlled trial. <i>Annals of the Rheumatic Diseases</i> , 2020, 79, 141-149.	0.9	223
25	Urine Metabolic Fingerprints Encode Subtypes of Kidney Diseases. <i>Angewandte Chemie - International Edition</i> , 2020, 59, 1703-1710.	13.8	99
26	Zirconia Hybrid Nanoshells for Nutrient and Toxin Detection. <i>Small</i> , 2020, 16, e2003902.	10.0	37
27	MicroRNAs in Systemic Lupus Erythematosus: a Perspective on the Path from Biological Discoveries to Clinical Practice. <i>Current Rheumatology Reports</i> , 2020, 22, 17.	4.7	20
28	Inhibition of Glycolysis in Pathogenic TH17 Cells through Targeting a miR-21-Pel1-c-Rel Pathway Prevents Autoimmunity. <i>Journal of Immunology</i> , 2020, 204, 3160-3170.	0.8	17
29	Urinary activated leukocyte cell adhesion molecule as a novel biomarker of lupus nephritis histology. <i>Arthritis Research and Therapy</i> , 2020, 22, 122.	3.5	23
30	MicroRNA-125a-Loaded Polymeric Nanoparticles Alleviate Systemic Lupus Erythematosus by Restoring Effector/Regulatory T Cells Balance. <i>ACS Nano</i> , 2020, 14, 4414-4429.	14.6	53
31	Taurine Metabolism Aggravates the Progression of Lupus by Promoting the Function of Plasmacytoid Dendritic Cells. <i>Arthritis and Rheumatology</i> , 2020, 72, 2106-2117.	5.6	13
32	Spermidine Suppresses Inflammatory DC Function by Activating the FOXO3 Pathway and Counteracts Autoimmunity. <i>IScience</i> , 2020, 23, 100807.	4.1	49
33	PBX1 expression in uterine natural killer cells drives fetal growth. <i>Science Translational Medicine</i> , 2020, 12, .	12.4	54
34	The MicroRNA miR-22 Represses Th17 Cell Pathogenicity by Targeting PTEN-Regulated Pathways. <i>ImmunoHorizons</i> , 2020, 4, 308-318.	1.8	6
35	MicroRNA-148a facilitates inflammatory dendritic cell differentiation and autoimmunity by targeting MAFB. <i>JCI Insight</i> , 2020, 5, .	5.0	14
36	Evaluation of 12 different assays for detecting ANCA in Chinese patients with GPA and MPA: a multicenter study in China. <i>Clinical Rheumatology</i> , 2019, 38, 3477-3483.	2.2	5

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37	Genomic test ends a long diagnostic odyssey in a patient with resistance to thyroid hormones. <i>Thyroid Research</i> , 2019, 12, 7.	1.5	1
38	Interferon- β exacerbates neuropsychiatric phenotypes in lupus-prone mice. <i>Arthritis Research and Therapy</i> , 2019, 21, 205.	3.5	14
39	Excessive CD11c ⁺ Tbet ⁺ B cells promote aberrant T _{FH} differentiation and affinity-based germinal center selection in lupus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 18550-18560.	7.1	68
40	Lupus-associated atypical memory B cells are mTORC1-hyperactivated and functionally dysregulated. <i>Annals of the Rheumatic Diseases</i> , 2019, 78, 1090-1100.	0.9	61
41	Identification of Renal Long Non-coding RNA RP11-2B6.2 as a Positive Regulator of Type I Interferon Signaling Pathway in Lupus Nephritis. <i>Frontiers in Immunology</i> , 2019, 10, 975.	4.8	52
42	Functional rare and low frequency variants in BLK and BANK1 contribute to human lupus. <i>Nature Communications</i> , 2019, 10, 2201.	12.8	73
43	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.	3.5	36
44	Systemic lupus erythematosus: A new autoimmune disorder in Kabuki syndrome. <i>European Journal of Medical Genetics</i> , 2019, 62, 103538.	1.3	10
45	Epigenetics of Lupus. , 2019, , 69-85.		0
46	MicroRNA-663 induces immune dysregulation by inhibiting TGF- β 1 production in bone marrow-derived mesenchymal stem cells in patients with systemic lupus erythematosus. <i>Cellular and Molecular Immunology</i> , 2019, 16, 260-274.	10.5	50
47	Type I interferons promote the survival and proinflammatory properties of transitional B cells in systemic lupus erythematosus patients. <i>Cellular and Molecular Immunology</i> , 2019, 16, 367-379.	10.5	40
48	A plausibly causal functional lupus-associated risk variant in the STAT1-STAT4 locus. <i>Human Molecular Genetics</i> , 2018, 27, 2392-2404.	2.9	34
49	MicroRNA networks associated with active systemic juvenile idiopathic arthritis regulate CD163 expression and anti-inflammatory functions in macrophages through two distinct mechanisms. <i>Journal of Leukocyte Biology</i> , 2018, 103, 71-85.	3.3	19
50	Peli1 negatively regulates noncanonical NF- κ B signaling to restrain systemic lupus erythematosus. <i>Nature Communications</i> , 2018, 9, 1136.	12.8	55
51	Exome-wide association study identifies four novel loci for systemic lupus erythematosus in Han Chinese population. <i>Annals of the Rheumatic Diseases</i> , 2018, 77, 417-417.	0.9	50
52	Identification of LncRNA Linc00513 Containing Lupus-Associated Genetic Variants as a Novel Regulator of Interferon Signaling Pathway. <i>Frontiers in Immunology</i> , 2018, 9, 2967.	4.8	56
53	Association of Abnormal Elevations in IFIT3 With Overactive Cyclic GMP-AMP Synthase/Stimulator of Interferon Genes Signaling in Human Systemic Lupus Erythematosus Monocytes. <i>Arthritis and Rheumatology</i> , 2018, 70, 2036-2045.	5.6	57
54	Iron Drives T Helper Cell Pathogenicity by Promoting RNA-Binding Protein PCBP1-Mediated Proinflammatory Cytokine Production. <i>Immunity</i> , 2018, 49, 80-92.e7.	14.3	107

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55	A missense variant in NCF1 is associated with susceptibility to multiple autoimmune diseases. <i>Nature Genetics</i> , 2017, 49, 433-437.	21.4	143
56	Confirmation of five novel susceptibility loci for Systemic Lupus Erythematosus (SLE) and integrated network analysis of 82 SLE susceptibility loci. <i>Human Molecular Genetics</i> , 2017, 26, ddx026.	2.9	47
57	Novel insights of microRNAs in the development of systemic lupus erythematosus. <i>Current Opinion in Rheumatology</i> , 2017, 29, 450-457.	4.3	20
58	Antigen-specific CD8+ T cell feedback activates NLRP3 inflammasome in antigen-presenting cells through perforin. <i>Nature Communications</i> , 2017, 8, 15402.	12.8	61
59	The role of long non-coding RNAs in rheumatic diseases. <i>Nature Reviews Rheumatology</i> , 2017, 13, 657-669.	8.0	65
60	FGF2 cooperates with IL-17 to promote autoimmune inflammation. <i>Scientific Reports</i> , 2017, 7, 7024.	3.3	22
61	T-bet+CD11c+ B cells are critical for antichromatin immunoglobulin G production in the development of lupus. <i>Arthritis Research and Therapy</i> , 2017, 19, 225.	3.5	58
62	MiR-125a Is a critical modulator for neutrophil development. <i>PLoS Genetics</i> , 2017, 13, e1007027.	3.5	19
63	Identification of Cyclin-Dependent Kinase 1 as a Novel Regulator of Type I Interferon Signaling in Systemic Lupus Erythematosus. <i>Arthritis and Rheumatology</i> , 2016, 68, 1222-1232.	5.6	35
64	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.	5.6	89
65	MicroRNA in Systemic Lupus Erythematosus. , 2016, , 231-236.		2
66	Dendritic Cells in Systemic Lupus Erythematosus: From Pathogenic Players to Therapeutic Tools. <i>Mediators of Inflammation</i> , 2016, 2016, 1-12.	3.0	43
67	In Vivo Therapeutic Success of MicroRNA-155 Antagomir in a Mouse Model of Lupus Alveolar Hemorrhage. <i>Arthritis and Rheumatology</i> , 2016, 68, 953-964.	5.6	57
68	Monocyte MicroRNA Expression in Active Systemic Juvenile Idiopathic Arthritis Implicates MicroRNA-125a in Polarized Monocyte Phenotypes. <i>Arthritis and Rheumatology</i> , 2016, 68, 2300-2313.	5.6	62
69	MicroRNA-130b Ameliorates Murine Lupus Nephritis Through Targeting the Type I Interferon Pathway on Renal Mesangial Cells. <i>Arthritis and Rheumatology</i> , 2016, 68, 2232-2243.	5.6	59
70	Identification of the long noncoding RNA NEAT1 as a novel inflammatory regulator acting through MAPK pathway in human lupus. <i>Journal of Autoimmunity</i> , 2016, 75, 96-104.	6.5	233
71	Low-dose interleukin-2 treatment selectively modulates CD4+ T cell subsets in patients with systemic lupus erythematosus. <i>Nature Medicine</i> , 2016, 22, 991-993.	30.7	457
72	High-density genotyping of immune-related loci identifies new SLE risk variants in individuals with Asian ancestry. <i>Nature Genetics</i> , 2016, 48, 323-330.	21.4	219

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73	MicroRNA-499 Rs3746444 polymorphism and biliary atresia. <i>Digestive and Liver Disease</i> , 2016, 48, 423-428.	0.9	9
74	miR-744 enhances type I interferon signaling pathway by targeting PTP1B in primary human renal mesangial cells. <i>Scientific Reports</i> , 2015, 5, 12987.	3.3	23
75	MiR-125a-5p Decreases the Sensitivity of Treg cells Toward IL-6-Mediated Conversion by Inhibiting IL-6R and STAT3 Expression. <i>Scientific Reports</i> , 2015, 5, 14615.	3.3	33
76	Association of large intergenic noncoding RNA expression with disease activity and organ damage in systemic lupus erythematosus. <i>Arthritis Research and Therapy</i> , 2015, 17, 131.	3.5	92
77	Type I Interferon Inhibition of MicroRNA-146a Maturation Through Up-Regulation of Monocyte Chemoattractant Protein-1 in Systemic Lupus Erythematosus. <i>Arthritis and Rheumatology</i> , 2015, 67, 3209-3218.	5.6	51
78	Predominant Role of Plasmacytoid Dendritic Cells in Stimulating Systemic Autoimmunity. <i>Frontiers in Immunology</i> , 2015, 6, 526.	4.8	31
79	Effect of TACI Signaling on Humoral Immunity and Autoimmune Diseases. <i>Journal of Immunology Research</i> , 2015, 2015, 1-12.	2.2	57
80	Lupus risk variants in the PXX locus alter B-cell receptor internalization. <i>Frontiers in Genetics</i> , 2015, 5, 450.	2.3	25
81	NF- κ B-induced microRNA-31 promotes epidermal hyperplasia by repressing protein phosphatase 6 in psoriasis. <i>Nature Communications</i> , 2015, 6, 7652.	12.8	191
82	miRNAs in the Pathogenesis of Systemic Lupus Erythematosus. <i>International Journal of Molecular Sciences</i> , 2015, 16, 9557-9572.	4.1	55
83	Lupus Risk Variant Increases pSTAT1 Binding and Decreases ETS1 Expression. <i>American Journal of Human Genetics</i> , 2015, 96, 731-739.	6.2	36
84	MiR-125a targets effector programs to stabilize Treg-mediated immune homeostasis. <i>Nature Communications</i> , 2015, 6, 7096.	12.8	133
85	Posttranscriptional T cell gene regulation to limit Tfh cells and autoimmunity. <i>Current Opinion in Immunology</i> , 2015, 37, 21-27.	5.5	14
86	Growth Factor FGF2 Cooperates with Interleukin-17 to Repair Intestinal Epithelial Damage. <i>Immunity</i> , 2015, 43, 488-501.	14.3	174
87	Brief Report: Identification of <i>MTMR3</i> as a Novel Susceptibility Gene for Lupus Nephritis in Northern Han Chinese by Shared-Gene Analysis With IgA Nephropathy. <i>Arthritis and Rheumatology</i> , 2014, 66, 2842-2848.	5.6	44
88	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.	6.2	59
89	Chest imaging manifestations in lupus nephritis. <i>Clinical Rheumatology</i> , 2014, 33, 817-823.	2.2	1
90	Alterations in the Microbiota Drive Interleukin-17C Production from Intestinal Epithelial Cells to Promote Tumorigenesis. <i>Immunity</i> , 2014, 40, 140-152.	14.3	153

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91	Elevated microRNA-155 promotes foam cell formation by targeting HBP1 in atherogenesis. <i>Cardiovascular Research</i> , 2014, 103, 100-110.	3.8	131
92	microRNA-mediated regulation of innate immune response in rheumatic diseases. <i>Arthritis Research and Therapy</i> , 2013, 15, 210.	3.5	34
93	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
94	Type I interferons in Sjögren's syndrome. <i>Autoimmunity Reviews</i> , 2013, 12, 558-566.	5.8	97
95	Meta-analysis Followed by Replication Identifies Loci in or near <i>CDKN1B</i> , <i>TET3</i> , <i>CD80</i> , <i>DRAM1</i> , and <i>ARID5B</i> as Associated with Systemic Lupus Erythematosus in Asians. <i>American Journal of Human Genetics</i> , 2013, 92, 41-51.	6.2	184
96	Epigenetics of Lupus. , 2013, , 46-56.		1
97	Trans-Ancestral Studies Fine Map the SLE-Susceptibility Locus <i>TNFSF4</i> . <i>PLoS Genetics</i> , 2013, 9, e1003554.	3.5	50
98	MicroRNA involvement in lupus. <i>Current Opinion in Rheumatology</i> , 2012, 24, 489-498.	4.3	30
99	miR-132 regulates the differentiation of dopamine neurons by directly targeting <i>Nurr1</i> expression. <i>Journal of Cell Science</i> , 2012, 125, 1673-82.	2.0	132
100	Distinct roles of myeloid and plasmacytoid dendritic cells in systemic lupus erythematosus. <i>Autoimmunity Reviews</i> , 2012, 11, 890-897.	5.8	77
101	Identification of microRNA-31 as a novel regulator contributing to impaired interleukin-2 production in T cells from patients with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2012, 64, 3715-3725.	6.7	97
102	T-614 alters the production of matrix metalloproteinases (MMP-1 and MMP-3) and inhibits the migratory expansion of rheumatoid synovial fibroblasts, in vitro. <i>International Immunopharmacology</i> , 2012, 13, 54-60.	3.8	29
103	MicroRNAs as novel regulators of systemic lupus erythematosus pathogenesis. <i>Nature Reviews Rheumatology</i> , 2012, 8, 701-709.	8.0	143
104	MicroRNA-125b/Lin28 Pathway Contributes to the Mesendodermal Fate Decision of Embryonic Stem Cells. <i>Stem Cells and Development</i> , 2012, 21, 1524-1537.	2.1	25
105	A Novel Vector-Based Method for Exclusive Overexpression of Star-Form MicroRNAs. <i>PLoS ONE</i> , 2012, 7, e41504.	2.5	9
106	The microRNA miR-23b suppresses IL-17-associated autoimmune inflammation by targeting <i>TAB2</i> , <i>TAB3</i> and <i>IKK-β</i> . <i>Nature Medicine</i> , 2012, 18, 1077-1086.	30.7	397
107	Gene-gene interaction of <i>BLK</i> , <i>TNFSF4</i> , <i>TRAF1</i> , <i>TNFAIP3</i> , and <i>REL</i> in systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2012, 64, 222-231.	6.7	73
108	Single-nucleotide polymorphisms and haplotype of <i>CYP2E1</i> gene associated with systemic lupus erythematosus in Chinese population. <i>Arthritis Research and Therapy</i> , 2011, 13, R11.	3.5	11

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109	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.	3.5	241
110	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.	3.5	181
111	Multidimensional Single Cell Based STAT Phosphorylation Profiling Identifies a Novel Biosignature for Evaluation of Systemic Lupus Erythematosus Activity. <i>PLoS ONE</i> , 2011, 6, e21671.	2.5	41
112	miR-155 and its star-form partner miR-155* cooperatively regulate type I interferon production by human plasmacytoid dendritic cells. <i>Blood</i> , 2010, 116, 5885-5894.	1.4	233
113	MicroRNA-125a contributes to elevated inflammatory chemokine RANTES levels via targeting KLF13 in systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2010, 62, 3425-3435.	6.7	212
114	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.	7.1	324
115	Genome-Wide Association Study in Asian Populations Identifies Variants in ETS1 and WDFY4 Associated with Systemic Lupus Erythematosus. <i>PLoS Genetics</i> , 2010, 6, e1000841.	3.5	378
116	Non-synonymous variant (Gly307Ser) in CD226 is associated with susceptibility to multiple autoimmune diseases. <i>Rheumatology</i> , 2010, 49, 1239-1244.	1.9	64
117	MicroRNA-21 and MicroRNA-148a Contribute to DNA Hypomethylation in Lupus CD4+ T Cells by Directly and Indirectly Targeting DNA Methyltransferase 1. <i>Journal of Immunology</i> , 2010, 184, 6773-6781.	0.8	499
118	MicroRNA-146a contributes to abnormal activation of the type I interferon pathway in human lupus by targeting the key signaling proteins. <i>Arthritis and Rheumatism</i> , 2009, 60, 1065-1075.	6.7	679
119	T-614, a novel immunomodulator, attenuates joint inflammation and articular damage in collagen-induced arthritis. <i>Arthritis Research and Therapy</i> , 2008, 10, R136.	3.5	72
120	Association of elevated transcript levels of interferon-inducible chemokines with disease activity and organ damage in systemic lupus erythematosus patients. <i>Arthritis Research and Therapy</i> , 2008, 10, R112.	3.5	81
121	Interferon-induced protein IFIT4 is associated with systemic lupus erythematosus and promotes differentiation of monocytes into dendritic cell-like cells. <i>Arthritis Research and Therapy</i> , 2008, 10, R91.	3.5	25
122	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.	7.1	62
123	Effects of FTY720 on BXSB lupus-prone mice: a preliminary study. <i>APLAR Journal of Rheumatology</i> , 2007, 10, 214-220.	0.2	0
124	Current advances in the human lupus genetics. <i>Current Rheumatology Reports</i> , 2004, 6, 391-398.	4.7	35
125	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
126	PARP alleles within the linked chromosomal region are associated with systemic lupus erythematosus. <i>Journal of Clinical Investigation</i> , 1999, 103, 1135-1140.	8.2	99