List of Publications by Year in descending order

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DONC OINC YE

#	Article	IF	CITATIONS
1	Genome-wide association study in a Chinese Han population identifies nine new susceptibility loci for systemic lupus erythematosus. Nature Genetics, 2009, 41, 1234-1237.	9.4	868
2	Genome-Wide Association Study in Asian Populations Identifies Variants in ETS1 and WDFY4 Associated with Systemic Lupus Erythematosus. PLoS Genetics, 2010, 6, e1000841.	1.5	378
3	Newâ€onset autoimmune phenomena postâ€COVIDâ€19 vaccination. Immunology, 2022, 165, 386-401.	2.0	288
4	Emerging role of long noncoding RNAs in autoimmune diseases. Autoimmunity Reviews, 2015, 14, 798-805.	2.5	226
5	NLRP3: A promising therapeutic target for autoimmune diseases. Autoimmunity Reviews, 2018, 17, 694-702.	2.5	188
6	Emerging role of air pollution in autoimmune diseases. Autoimmunity Reviews, 2019, 18, 607-614.	2.5	188
7	Long noncoding RNAs: Novel insights into gastric cancer. Cancer Letters, 2015, 356, 357-366.	3.2	179
8	Comparative effectiveness and tolerance of treatments for <i>Helicobacter pylori</i> : systematic review and network meta-analysis. BMJ, The, 2015, 351, h4052.	3.0	137
9	Subclinical atherosclerosis in patients with systemic lupus erythematosus: A systemic review and meta-analysis. Autoimmunity Reviews, 2016, 15, 22-37.	2.5	120
10	Meta-analysis of 208370 East Asians identifies 113 susceptibility loci for systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2021, 80, 632-640.	0.5	103
11	Translation of noncoding RNAs: Focus on IncRNAs, pri-miRNAs, and circRNAs. Experimental Cell Research, 2017, 361, 1-8.	1.2	97
12	Causal Effects of Gut Microbiome on Systemic Lupus Erythematosus: A Two-Sample Mendelian Randomization Study. Frontiers in Immunology, 2021, 12, 667097.	2.2	94
13	Identification of long non-coding RNAs GAS5, linc0597 and Inc-DC in plasma as novel biomarkers for systemic lupus erythematosus. Oncotarget, 2017, 8, 23650-23663.	0.8	92
14	Circular <scp>RNA</scp> expression profile and potential function of hsa_circ_0045272 in systemic lupus erythematosus. Immunology, 2018, 155, 137-149.	2.0	74
15	Potential link between m 6 A modification and systemic lupus erythematosus. Molecular Immunology, 2018, 93, 55-63.	1.0	68
16	Competitive endogenous RNA network: potential implication for systemic lupus erythematosus. Expert Opinion on Therapeutic Targets, 2017, 21, 639-648.	1.5	67
17	Influence of social support on health-related quality of life in patients with systemic lupus erythematosus. Clinical Rheumatology, 2009, 28, 265-269.	1.0	54
18	Prevalence of Suicide Attempts among College Students in China: A Meta-Analysis. PLoS ONE, 2015, 10, e0116303.	1.1	50

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19	microRNAs function in CD8+T cell biology. Journal of Leukocyte Biology, 2015, 97, 487-497.	1.5	49
20	Subclinical Atherosclerosis in Patients With Inflammatory Bowel Diseases: A Systematic Review and Meta-Analysis. Angiology, 2017, 68, 447-461.	0.8	47
21	Comprehensive long non-coding RNA expression profiling reveals their potential roles in systemic lupus erythematosus. Cellular Immunology, 2017, 319, 17-27.	1.4	47
22	Interleukin-13: A promising therapeutic target for autoimmune disease. Cytokine and Growth Factor Reviews, 2019, 45, 9-23.	3.2	45
23	Differential Plasma Expression Profiles of Long Non-Coding RNAs Reveal Potential Biomarkers for Systemic Lupus Erythematosus. Biomolecules, 2019, 9, 206.	1.8	44
24	IL-33 in rheumatoid arthritis: Potential role in pathogenesis and therapy. Human Immunology, 2013, 74, 1057-1060.	1.2	41
25	Interleukin-35: a Potential Therapeutic Agent for Autoimmune Diseases. Inflammation, 2017, 40, 303-310.	1.7	41
26	Serum resistin levels in patients with rheumatoid arthritis and systemic lupus erythematosus: a meta-analysis. Clinical Rheumatology, 2015, 34, 1713-1720.	1.0	40
27	Differentially expressed circular RNAs in systemic lupus erythematosus and their clinical significance. Biomedicine and Pharmacotherapy, 2018, 107, 1720-1727.	2.5	36
28	Meta-analysis of GWAS on two Chinese populations followed by replication identifies novel genetic variants on the X chromosome associated with systemic lupus erythematosus. Human Molecular Genetics, 2015, 24, 274-284.	1.4	35
29	Circular RNAs and systemic lupus erythematosus. Experimental Cell Research, 2016, 346, 248-254.	1.2	35
30	Intratumoral and peritumoral expression of CD68 and CD206 in hepatocellular carcinoma and their prognostic value. Oncology Reports, 2017, 38, 886-898.	1.2	35
31	The correlation between monocyte chemoattractant protein-1 and the arthritis of systemic lupus erythematosus among Chinese. Archives of Dermatological Research, 2005, 296, 366-371.	1.1	34
32	Identification of <i>ST3AGL4</i> , <i>MFHAS1, CSNK2A2</i> and <i>CD226</i> as loci associated with systemic lupus erythematosus (SLE) and evaluation of SLE genetics in drug repositioning. Annals of the Rheumatic Diseases, 2018, 77, 1078-1084.	0.5	34
33	Hypoxia-inducible factor-1α: a promising therapeutic target for autoimmune diseases. Expert Opinion on Therapeutic Targets, 2017, 21, 715-723.	1.5	33
34	Association of long noncoding RNAs expression levels and their gene polymorphisms with systemic lupus erythematosus. Scientific Reports, 2017, 7, 15119.	1.6	33
35	Associated Variables of Myositis in Systemic Lupus Erythematosus: A Cross-Sectional Study. Medical Science Monitor, 2017, 23, 2543-2549.	0.5	30
36	Three SNPs in chromosome 11q23.3 are independently associated with systemic lupus erythematosus in Asians. Human Molecular Genetics, 2014, 23, 524-533.	1.4	29

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37	Polymorphisms in the promoter region of RANTES in Han Chinese and their relationship with systemic lupus erythematosus. Archives of Dermatological Research, 2005, 297, 108-113.	1.1	28
38	Association between tumor necrosis factor-α (TNF-α) promoter â^'308 G/A and response to TNF-α blockers in rheumatoid arthritis: a meta-analysis. Modern Rheumatology, 2013, 23, 489-495.	0.9	28
39	A Meta-Analysis of Cardiovascular Events in Systemic Lupus Erythematosus. Immunological Investigations, 2019, 48, 505-520.	1.0	28
40	Association between VDR polymorphisms and multiple sclerosis: systematic review and updated meta-analysis of case-control studies. Neurological Sciences, 2018, 39, 225-234.	0.9	27
41	The impact of SLE on health-related quality of life assessed with SF-36: a systemic review and meta-analysis. Lupus, 2019, 28, 371-382.	0.8	27
42	Identification of new susceptibility loci associated with rheumatoid arthritis. Annals of the Rheumatic Diseases, 2020, 79, 1565-1571.	0.5	27
43	Circadian clock genes as promising therapeutic targets for autoimmune diseases. Autoimmunity Reviews, 2021, 20, 102866.	2.5	27
44	Evidence for genetic association of TBX21 and IFNG with systemic lupus erythematosus in a Chinese Han population. Scientific Reports, 2016, 6, 22081.	1.6	26
45	Emerging role of adipokines in systemic lupus erythematosus. Immunologic Research, 2016, 64, 820-830.	1.3	26
46	The prevalence and risk factors for serositis in patients with systemic lupus erythematosus: a cross-sectional study. Rheumatology International, 2017, 37, 305-311.	1.5	26
47	Therapeutic potential of IL-15 in rheumatoid arthritis. Human Immunology, 2015, 76, 812-818.	1.2	25
48	Associations Between PADI4 Gene Polymorphisms and Rheumatoid Arthritis: An Updated Meta-analysis. Archives of Medical Research, 2015, 46, 317-325.	1.5	24
49	Who benefited from the New Rural Cooperative Medical System in China? A case study on Anhui Province. BMC Health Services Research, 2016, 16, 195.	0.9	23
50	Association between IL-33 Gene Polymorphisms (rs1929992, rs7044343) and Systemic Lupus Erythematosus in a Chinese Han Population. Immunological Investigations, 2016, 45, 575-583.	1.0	23
51	Prevalence and risk factors of chronic obstructive pulmonary disease in Anhui Province, China: a population-based survey. BMC Pulmonary Medicine, 2019, 19, 102.	0.8	23
52	Long Non-coding RNAs Genes Polymorphisms and Their Expression Levels in Patients With Rheumatoid Arthritis. Frontiers in Immunology, 2019, 10, 2529.	2.2	23
53	A meta-analysis of the association of <i>STAT4</i> polymorphism with systemic lupus erythematosus. Modern Rheumatology, 2010, 20, 257-262.	0.9	22
54	Emerging role of semaphorin-3A in autoimmune diseases. Inflammopharmacology, 2018, 26, 655-665.	1.9	22

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55	Association of CTLA-4 variants with susceptibility to inflammatory bowel disease: A meta-analysis. Human Immunology, 2014, 75, 227-233.	1.2	21
56	Single nucleotide polymorphisms of HSP90AA1 gene influence response of SLE patients to glucocorticoids treatment. SpringerPlus, 2016, 5, 222.	1.2	21
5 7	Association between the serum level of vitamin D and systemic sclerosis in a Chinese population: a case control study. International Journal of Rheumatic Diseases, 2017, 20, 1002-1008.	0.9	21
58	Coagulation cascade and complement system in systemic lupus erythematosus. Oncotarget, 2018, 9, 14862-14881.	0.8	21
59	Relationship between the IL12B (rs3212227) gene polymorphism and susceptibility to multiple autoimmune diseases: A meta-analysis. Modern Rheumatology, 2016, 26, 749-756.	0.9	20
60	An Empirical Analysis of Rural-Urban Differences in Out-Of-Pocket Health Expenditures in a Low-Income Society of China. PLoS ONE, 2016, 11, e0154563.	1.1	20
61	<scp>HIV</scp> / <scp>AIDS</scp> stigma among older <scp>PLWHA</scp> in south rural <scp>C</scp> hina. International Journal of Nursing Practice, 2015, 21, 221-228.	0.8	19
62	MicroRNA-210 and its theranostic potential. Expert Opinion on Therapeutic Targets, 2016, 20, 1325-1338.	1.5	19
63	Meta-analysis of GWASÂonÂboth Chinese and European populations identifies GPR173 as a novel X chromosome susceptibility gene for SLE. Arthritis Research and Therapy, 2018, 20, 92.	1.6	19
64	Expression of several long noncoding RNAs in peripheral blood mononuclear cells of patients with systemic lupus erythematosus. Advances in Medical Sciences, 2019, 64, 430-436.	0.9	19
65	Two follicle-stimulating hormone receptor polymorphisms and polycystic ovary syndrome risk: a meta-analysis. European Journal of Obstetrics, Gynecology and Reproductive Biology, 2014, 182, 27-32.	0.5	18
66	Increased circulating CXCL13 levels in systemic lupus erythematosus and rheumatoid arthritis: a meta-analysis. Clinical Rheumatology, 2020, 39, 281-290.	1.0	18
67	Association of KIR genotype with susceptibility to HLA-B27-positive ankylosing spondylitis. Modern Rheumatology, 2013, 23, 538-541.	0.9	17
68	HIV, other sexually transmitted infections, and risk behaviors among female sex workers in Liuzhou, China. International Journal of Gynecology and Obstetrics, 2015, 128, 18-22.	1.0	17
69	Plasma levels of adipokines in systemic lupus erythematosus patients. Cytokine, 2016, 86, 15-20.	1.4	17
70	Association between serum/plasma adiponectin levels and immune-mediated diseases: a meta-analysis. Archives of Dermatological Research, 2017, 309, 625-635.	1.1	17
71	Association between HLA-DQB1 polymorphisms and pemphigus vulgaris: A meta-analysis. Immunological Investigations, 2018, 47, 101-112.	1.0	17
72	Association Study and Fine-Mapping MajorÂHistocompatibility Complex AnalysisÂof Pemphigus Vulgaris in aÂHanÂChinese Population. Journal of Investigative Dermatology, 2018, 138, 2307-2314.	0.3	17

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73	Association of interleukin-1 family cytokines single nucleotide polymorphisms with susceptibility to systemic sclerosis: an independent case–control study and a meta-analysis. Immunologic Research, 2016, 64, 1041-1052.	1.3	16
74	Association of leptin and leptin receptor gene polymorphisms with systemic lupus erythematosus in a Chinese population. Journal of Cellular and Molecular Medicine, 2017, 21, 1732-1741.	1.6	16
75	RNAi Silencing of HIF-11̂± Ameliorates Lupus Development in MRL/lpr Mice. Inflammation, 2018, 41, 1717-1730.	1.7	16
76	Risk of gestational diabetes mellitus in systemic lupus erythematosus pregnancy: a systematic review and meta-analysis. BMC Pregnancy and Childbirth, 2019, 19, 179.	0.9	16
77	Causes and Factors Associated with Frequent Hospitalization in Chinese Patients with Systemic Lupus Erythematosus: An Ambispective Cohort Study. Medical Science Monitor, 2019, 25, 8061-8068.	0.5	16
78	Ambient air pollutants increase the risk of immunoglobulin E–mediated allergic diseases: a systematic review and meta-analysis. Environmental Science and Pollution Research, 2022, 29, 49534-49552.	2.7	16
79	Integrated analysis of IncRNA, miRNA and mRNA expression profiling in patients with systemic lupus erythematosus. Archives of Medical Science, 2019, 15, 872-879.	0.4	15
80	TREX1 As a Potential Therapeutic Target for Autoimmune and Inflammatory Diseases. Current Pharmaceutical Design, 2019, 25, 3239-3247.	0.9	15
81	Geneâ€Based Metaâ€Analysis of Genomeâ€Wide Association Study Data Identifies Independent Singleâ€Nucleotide Polymorphisms in <i>ANXA6</i> as Being Associated With Systemic Lupus Erythematosus in Asian Populations. Arthritis and Rheumatology, 2015, 67, 2966-2977.	2.9	14
82	UBASH3A gene polymorphisms and expression profile in rheumatoid arthritis. Autoimmunity, 2019, 52, 21-26.	1.2	14
83	Review on the Alteration of Gut Microbiota: The Role of HIV Infection and Old Age. AIDS Research and Human Retroviruses, 2020, 36, 556-565.	0.5	14
84	Strong policies control the spread of COVIDâ€19 in China. Journal of Medical Virology, 2020, 92, 1980-1987.	2.5	14
85	Association of TNFSF4 polymorphisms with systemic lupus erythematosus: a meta-analysis. Modern Rheumatology, 2013, 23, 686-693.	0.9	13
86	An Updated Meta-Analysis: Risk Conferred by Glutathione S-Transferases (<i>GSTM1</i> and <i>GSTT1</i>) Polymorphisms to Age-Related Cataract. Journal of Ophthalmology, 2015, 2015, 1-10.	0.6	13
87	Association study of TRAP1 gene polymorphisms with susceptibility and glucocorticoids efficacy of systemic lupus erythematosus. Gene, 2018, 671, 117-126.	1.0	13
88	Associations of Vitamin D Receptor Single Nucleotide Polymorphisms with Susceptibility to Systemic Sclerosis. Archives of Medical Research, 2019, 50, 368-376.	1.5	13
89	Elevated plasma midkine and pleiotrophin levels in patients with systemic lupus erythematosus. Oncotarget, 2017, 8, 40181-40189.	0.8	13
90	Altered microRNAs expression in T cells of patients with SLE involved in the lack of vitamin D. Oncotarget, 2017, 8, 62099-62110.	0.8	13

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91	Involvement of N6-methyladenosine modifications of long noncoding RNAs in systemic lupus erythematosus. Molecular Immunology, 2022, 143, 77-84.	1.0	13
92	TCR-CD3ζ gene polymorphisms and expression profile in rheumatoid arthritis. Autoimmunity, 2016, 49, 466-471.	1.2	12
93	A meta-analysis of the relationship between MYO9B gene polymorphisms and susceptibility to Crohn's disease and ulcerative colitis. Human Immunology, 2016, 77, 990-996.	1.2	12
94	Decreased flow-mediated dilatation in patients with rheumatoid arthritis: a meta-analysis. Postgraduate Medical Journal, 2017, 93, 260-265.	0.9	12
95	The expression levels of long noncoding RNAs Inc0640 and Inc5150 and its gene singleâ€nucleotide polymorphisms in rheumatoid arthritis patients. Journal of Cellular Biochemistry, 2018, 119, 10095-10106.	1.2	12
96	Circulating antioxidant levels in systemic lupus erythematosus patients: a systematic review and meta-analysis. Biomarkers in Medicine, 2019, 13, 1137-1152.	0.6	12
97	Elevated Blood and Urinary ICAM-1 is a Biomarker for Systemic Lupus Erythematosus: A Systematic Review and Meta-Analysis. Immunological Investigations, 2020, 49, 15-31.	1.0	12
98	Decreased H19, GAS5, and linc0597 Expression and Association Analysis of Related Gene Polymorphisms in Rheumatoid Arthritis. Biomolecules, 2020, 10, 55.	1.8	12
99	BMI, disease activity, and health-related quality-of-life in systemic lupus erythematosus. Clinical Rheumatology, 2010, 29, 1413-1417.	1.0	11
100	Association study ofinterleukin-19 rs2243188polymorphism with systemic lupus erythematosus in a Chinese population. Autoimmunity, 2014, 47, 378-382.	1.2	11
101	Association of UBASH3A gene polymorphisms and systemic lupus erythematosus in a Chinese population. Gene, 2015, 565, 116-121.	1.0	11
102	Association of <i>interleukin-10</i> gene single nucleotide polymorphisms with rheumatoid arthritis in a Chinese population. Postgraduate Medical Journal, 2018, 94, 284-288.	0.9	11
103	Diagnostic value of urinary monocyte chemoattractant protein-1 in evaluating the activity of lupus nephritis: a meta-analysis. Lupus, 2020, 29, 599-606.	0.8	11
104	Genetic variant in <i>microRNA-146a</i> gene is associated with risk of rheumatoid arthritis. Annals of Medicine, 2021, 53, 824-829.	1.5	11
105	Willingness to use HIV pre-exposure prophylaxis and associated factors among men who have sex with men in Liuzhou, China. AIDS Research and Therapy, 2021, 18, 46.	0.7	10
106	Association Study of Matrix Metalloproteinases Gene Polymorphisms with Susceptibility to Rheumatoid Arthritis: A Meta-Analysis. Immunological Investigations, 2015, 44, 603-615.	1.0	9
107	Association of adiponectin and adiponectin receptor gene polymorphisms with rheumatoid arthritis in a Chinese population. Postgraduate Medical Journal, 2020, 96, 149-155.	0.9	9
108	Baseline survey for malaria prevalence in Khyber Pakhtunkhwa Province, Pakistan. Eastern Mediterranean Health Journal, 2020, 26, 453-460.	0.3	9

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109	Prevalence and influencing factors of anxiety and depression symptoms among surgical nurses during COVIDâ€19 pandemic: A largeâ€scale crossâ€sectional study. Nursing Open, 2022, 9, 752-764.	1.1	9
110	Short-term association of NO2 with hospital visits for chronic kidney disease and effect modification by temperature in Hefei, China: A time series study. Ecotoxicology and Environmental Safety, 2022, 237, 113505.	2.9	9
111	Biological insights into systemic lupus erythematosus through an immune cell-specific transcriptome-wide association study. Annals of the Rheumatic Diseases, 2022, 81, 1273-1280.	0.5	9
112	Effects of Disease Activity and Inflammatory Response on Hypercoagulability in Patients with Systemic Lupus Erythematosus. Archives of Medical Research, 2016, 47, 573-579.	1.5	8
113	Association of HLA-DQB1 polymorphisms with rheumatoid arthritis: a meta-analysis. Postgraduate Medical Journal, 2017, 93, 618-625.	0.9	8
114	Features associated with pulmonary arterial hypertension in Chinese hospitalized systemic lupus erythematosus patients. Clinical Rheumatology, 2018, 37, 1547-1553.	1.0	8
115	Seasonal variation in systemic lupus erythematosus and rheumatoid arthritis: An ecological study based on internet searches. Autoimmunity Reviews, 2019, 18, 825-827.	2.5	8
116	Association of omentin-1, adiponectin, and resistin genetic polymorphisms with systemic lupus erythematosus in a Chinese population. International Immunopharmacology, 2020, 83, 106343.	1.7	8
117	Low ambient temperature increases hospital re-admissions for systemic lupus erythematosus in humid subtropical region: a time series study. Environmental Science and Pollution Research, 2021, 28, 530-537.	2.7	8
118	Association of MALAT-1 gene single nucleotide polymorphisms with genetic susceptibility to systemic lupus erythematosus. Lupus, 2021, 30, 1923-1930.	0.8	8
119	Prevalence and associated factors of HIV infection among men who have sex with men in Hefei, China, 2013–2014: a cross-sectional study. International Journal of STD and AIDS, 2016, 27, 305-312.	0.5	7
120	Genetic variant of IL-10RA and susceptibility to rheumatoid arthritis in a Chinese population. Clinical Rheumatology, 2017, 36, 825-830.	1.0	7
121	Metaâ€analysis of associations between <i><scp>XRCC</scp>1</i> gene polymorphisms and susceptibility to systemic lupus erythematosus and rheumatoid arthritis. International Journal of Rheumatic Diseases, 2018, 21, 179-185.	0.9	7
122	Diagnostic accuracy of anti-keratin antibody for rheumatoid arthritis: a meta-analysis. Clinical Rheumatology, 2019, 38, 1841-1849.	1.0	7
123	Serum/plasma homocysteine levels in patients with systemic lupus erythematosus: a systematic review and meta-analysis. Clinical Rheumatology, 2020, 39, 1725-1736.	1.0	7
124	Diagnostic value of antiâ€citrullinated fibrinogen antibody in rheumatoid arthritis: A metaâ€analysis. International Journal of Rheumatic Diseases, 2019, 22, 599-607.	0.9	6
125	Association between ambient air pollution and multiple sclerosis: a systemic review and meta-analysis. Environmental Science and Pollution Research, 2021, 28, 58142-58153.	2.7	6
126	Association of lymphotoxin alpha polymorphism with systemic lupus erythematosus and rheumatoid arthritis: a metaâ€analysis. International Journal of Rheumatic Diseases, 2015, 18, 398-407.	0.9	5

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127	Genetic Polymorphism (rs329498) in the Pellino-1 Gene as Possible Predisposal Factor for Systemic Lupus Erythematosus in a Chinese Population. Immunological Investigations, 2016, 45, 181-190.	1.0	5
128	Association between rheumatoid arthritis and genetic variants of natural resistanceâ€associated macrophage protein 1 gene: A metaâ€analysis. International Journal of Rheumatic Diseases, 2018, 21, 1651-1658.	0.9	5
129	Levels of the macrophage migration inhibitory factor and polymorphisms in systemic lupus erythematosus: a meta-analysis. Archives of Medical Science, 2021, 17, 1232-1240.	0.4	5
130	Association of NCF2, NCF4, and CYBA Gene Polymorphisms with Rheumatoid Arthritis in a Chinese Population. Journal of Immunology Research, 2020, 2020, 1-11.	0.9	5
131	Non-causal effects of smoking and alcohol use on the risk of systemic lupus erythematosus. Autoimmunity Reviews, 2021, 20, 102890.	2.5	5
132	Altered mRNA expression levels of vaspin and adiponectin in peripheral blood mononuclear cells of systemic lupus erythematosus patients. Clinical and Experimental Rheumatology, 2019, 37, 458-464.	0.4	5
133	Proton pump inhibitors induce changes in the gut microbiome composition of systemic lupus erythematosus patients. BMC Microbiology, 2022, 22, 117.	1.3	5
134	Lack of association of Toll-like receptor 9 polymorphisms with susceptibility to systemic lupus erythematosus in an Asian population: a meta-analysis. Modern Rheumatology, 2012, 22, 550-556.	0.9	4
135	Identification of Mutations in Myocilin and Beta-1,4-galactosyltransferase 3 Genes in a Chinese Family with Primary Open-angle Glaucoma. Chinese Medical Journal, 2016, 129, 2810-2815.	0.9	4
136	Safety of measles-containing vaccines in post-marketing surveillance in Anhui, China. PLoS ONE, 2017, 12, e0172108.	1.1	4
137	Diagnostic accuracy of miRNAs as potential biomarkers for systemic lupus erythematosus: a meta-analysis. Clinical Rheumatology, 2018, 37, 2999-3007.	1.0	4
138	Serum 14-3-3η is a Marker that Complements Current Biomarkers for the Diagnosis of RA: Evidence from a Meta-analysis. Immunological Investigations, 2020, , 1-17.	1.0	4
139	Association of Midkine and Pleiotrophin Gene Polymorphisms With Systemic Lupus Erythematosus Susceptibility in Chinese Han Population. Frontiers in Immunology, 2020, 11, 110.	2.2	4
140	Antimalarials may reduce cancer risk in patients with systemic lupus erythematosus: a systematic review and meta-analysis of prospective studies. Annals of Medicine, 2021, 53, 1688-1696.	1.5	4
141	Circulating Insulin-like Growth Factor-1 Levels in Patients with Rheumatoid Arthritis: A Meta-analysis. Current Pharmaceutical Design, 2019, 25, 1091-1098.	0.9	4
142	Global Public Interest and Seasonal Variations in Alzheimer's Disease: Evidence From Google Trends. Frontiers in Medicine, 2021, 8, 778930.	1.2	3
143	Association of HLA-DR1, HLA-DR13, and HLA-DR16 Polymorphisms with Systemic Lupus Erythematosus: A Meta-Analysis. Journal of Immunology Research, 2022, 2022, 1-17.	0.9	3
144	ALKBH5 Expression could Affect the Function of T Cells in Systemic Lupus Erythematosus Patients: A Case-control Study. Current Pharmaceutical Design, 2022, 28, 2270-2278.	0.9	3

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145	Association study between X-linked susceptibility genes and clinical features in Chinese female patients with systemic lupus erythematosus. Autoimmunity, 2019, 52, 289-293.	1.2	2
146	The association between reproductive factors and systemic sclerosis in Chinese women: A case ontrol study and metaâ€analysis. International Journal of Rheumatic Diseases, 2019, 22, 1832-1840.	0.9	2
147	X chromosome and female bias in systemic lupus erythematosus: Focus on population-based evidence. Autoimmunity Reviews, 2019, 18, 109-111.	2.5	2
148	Elevated Urinary and Blood Vascular Cell Adhesion Molecule-1 as Potential Biomarkers for Active Systemic Lupus Erythematosus: A Meta-analysis. Current Pharmaceutical Design, 2020, 26, 5998-6006.	0.9	2
149	Association of HLA-B27 genetic polymorphisms with ankylosing spondylitis susceptibility worldwide: a meta-analysis. Modern Rheumatology, 2013, , 1.	0.9	1
150	Comparison of the adhesion of Streptococcus sanguinis to commonly used dental alloys stratified by gold content. Journal of Dental Sciences, 2016, 11, 437-442.	1.2	1
151	Intention to undergo HIV testing and associated factors among women in one high–HIV prevalence city. International Journal of Nursing Practice, 2017, 23, e12533.	0.8	1
152	Association of the rs17250932, rs4794067 and rs2240017 polymorphism in the TBX21 gene with autoimmune diseases. Allergologia Et Immunopathologia, 2021, 49, 83-90.	1.0	1
153	Increased circulating sclerostin levels in rheumatoid arthritis patients: an updated meta-analysis. Zeitschrift Fur Rheumatologie, 2023, 82, 51-58.	0.5	1
154	439: Association of Adverse Childhood Experiences and Health Risks among College Students in China. American Journal of Epidemiology, 2005, 161, S110-S110.	1.6	0
155	Response to the comment on â€~â€~Relationship between the IL12B (rs3212227) gene polymorphism and susceptibility to multiple autoimmune diseases: A meta-analysis''. Modern Rheumatology, 2017, 27, 180-181.	0.9	0
156	The Effect of Rosuvastatin on plasma/serum levels of high sensitivity C-reactive protein, Interleukin-6 and D-dimer in people living with Human Immunodeficiency Virus: a systematic review and meta-analysis AIDS Research and Human Retroviruses, 2021, 37, 821-833.	0.5	0
157	Predicting Malaria Incidence in Northern and Northwestern, Pakistan. Iranian Journal of Public Health, 2018, 47, 1961-1962.	0.3	0