

Xianyong Yin

List of Publications by Year in descending order

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49
papers

3,033
citations

279798

23
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51
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all docs

63
docs citations

63
times ranked

5918
citing authors

#	ARTICLE	IF	CITATIONS
1	The power of genetic diversity in genome-wide association studies of lipids. <i>Nature</i> , 2021, 600, 675-679.	27.8	353
2	Genomic Dissection of Population Substructure of Han Chinese and Its Implication in Association Studies. <i>American Journal of Human Genetics</i> , 2009, 85, 762-774.	6.2	338
3	Genome-wide association meta-analysis in Chinese and European individuals identifies ten new loci associated with systemic lupus erythematosus. <i>Nature Genetics</i> , 2016, 48, 940-946.	21.4	283
4	Deep sequencing of the MHC region in the Chinese population contributes to studies of complex disease. <i>Nature Genetics</i> , 2016, 48, 740-746.	21.4	188
5	Meta-analysis Followed by Replication Identifies Loci in or near CDKN1B, TET3, CD80, DRAM1, and ARID5B as Associated with Systemic Lupus Erythematosus in Asians. <i>American Journal of Human Genetics</i> , 2013, 92, 41-51.	6.2	184
6	A large-scale screen for coding variants predisposing to psoriasis. <i>Nature Genetics</i> , 2014, 46, 45-50.	21.4	183
7	Exome sequencing of Finnish isolates enhances rare-variant association power. <i>Nature</i> , 2019, 572, 323-328.	27.8	161
8	Genome-wide meta-analysis identifies multiple novel associations and ethnic heterogeneity of psoriasis susceptibility. <i>Nature Communications</i> , 2015, 6, 6916.	12.8	154
9	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
10	Whole-exome SNP array identifies 15 new susceptibility loci for psoriasis. <i>Nature Communications</i> , 2015, 6, 6793.	12.8	118
11	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
12	Epigenome-Wide Association Analysis Identified Nine Skin DNA Methylation Loci for Psoriasis. <i>Journal of Investigative Dermatology</i> , 2016, 136, 779-787.	0.7	75
13	Genetic Susceptibility to Vitiligo: GWAS Approaches for Identifying Vitiligo Susceptibility Genes and Loci. <i>Frontiers in Genetics</i> , 2016, 7, 3.	2.3	69
14	Genome-wide association analyses in Han Chinese identify two new susceptibility loci for amyotrophic lateral sclerosis. <i>Nature Genetics</i> , 2013, 45, 697-700.	21.4	67
15	Genome-wide association studies of metabolites in Finnish men identify disease-relevant loci. <i>Nature Communications</i> , 2022, 13, 1644.	12.8	63
16	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
17	Exome sequencing identifies SLC17A9 pathogenic gene in two Chinese pedigrees with disseminated superficial actinic prokeratosis. <i>Journal of Medical Genetics</i> , 2014, 51, 699-704.	3.2	32
18	Epigenome-wide association data implicates DNA methylation-mediated genetic risk in psoriasis. <i>Clinical Epigenetics</i> , 2016, 8, 131.	4.1	31

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19	Identification of seven novel loci associated with amino acid levels using single-variant and gene-based tests in 8545 Finnish men from the METSIM study. <i>Human Molecular Genetics</i> , 2018, 27, 1664-1674.	2.9	30
20	Causal Relationship and Shared Genetic Loci between Psoriasis and Type 2 Diabetes through Trans-Disease Meta-Analysis. <i>Journal of Investigative Dermatology</i> , 2021, 141, 1493-1502.	0.7	29
21	Confirmation of C4 gene copy number variation and the association with systemic lupus erythematosus in Chinese Han population. <i>Rheumatology International</i> , 2012, 32, 3047-3053.	3.0	28
22	Discovery of a novel genetic susceptibility locus on X chromosome for systemic lupus erythematosus. <i>Arthritis Research and Therapy</i> , 2015, 17, 349.	3.5	26
23	Down-regulated expression of IKZF1 mRNA in peripheral blood mononuclear cells from patients with systemic lupus erythematosus. <i>Rheumatology International</i> , 2011, 31, 819-822.	3.0	25
24	A Weighted Polygenic Risk Score Using 14 Known Susceptibility Variants to Estimate Risk and Age Onset of Psoriasis in Han Chinese. <i>PLoS ONE</i> , 2015, 10, e0125369.	2.5	22
25	Association of the Polymorphism rs13259960 in <i>SLEAR</i> With Predisposition to Systemic Lupus Erythematosus. <i>Arthritis and Rheumatology</i> , 2020, 72, 985-996.	5.6	22
26	Meta-analysis of NOD2/CARD15 polymorphisms with psoriasis and psoriatic arthritis. <i>Rheumatology International</i> , 2012, 32, 1893-1900.	3.0	21
27	Association of the Late Cornified Envelope-3 Genes with Psoriasis and Psoriatic Arthritis: A Systematic Review. <i>Journal of Genetics and Genomics</i> , 2015, 42, 49-56.	3.9	19
28	Common susceptibility variants are shared between schizophrenia and psoriasis in the Han Chinese population. <i>Journal of Psychiatry and Neuroscience</i> , 2016, 41, 413-421.	2.4	19
29	Common variants explain a large fraction of the variability in the liability to psoriasis in a Han Chinese population. <i>BMC Genomics</i> , 2014, 15, 87.	2.8	16
30	Platelet-derived Growth Factor Receptor Alpha Gene Mutations in Vitiligo Vulgaris. <i>Acta Dermato-Venereologica</i> , 2010, 90, 131-135.	1.3	15
31	Genome-wide meta-analysis identifies a novel susceptibility signal at <i>CACNA2D3</i> for nicotine dependence. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 557-567.	1.7	15
32	Genotype imputation for Han Chinese population using Haplotype Reference Consortium as reference. <i>Human Genetics</i> , 2018, 137, 431-436.	3.8	15
33	Several Critical Cell Types, Tissues, and Pathways Are Implicated in Genome-Wide Association Studies for Systemic Lupus Erythematosus. <i>G3: Genes, Genomes, Genetics</i> , 2016, 6, 1503-1511.	1.8	12
34	Fine mapping and subphenotyping implicates <i>ADRA1B</i> gene variants in psoriasis susceptibility in a Chinese population. <i>Epigenomics</i> , 2019, 11, 455-467.	2.1	10
35	Four genetic variants interact to confer susceptibility to atopic dermatitis in Chinese Han population. <i>Molecular Genetics and Genomics</i> , 2015, 290, 1493-1498.	2.1	9
36	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

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37	Combined effect of five single nucleotide polymorphisms related to IL23/Th17 pathway in the risk of psoriasis. <i>Immunogenetics</i> , 2014, 66, 215-218.	2.4	8
38	JAK2 and PTPRC mRNA expression in peripheral blood mononuclear cells from patients with systemic lupus erythematosus. <i>Clinical Rheumatology</i> , 2020, 39, 443-448.	2.2	8
39	Five regulatory genes detected by matching signatures of eQTL and GWAS in psoriasis. <i>Journal of Dermatological Science</i> , 2014, 76, 139-142.	1.9	7
40	More heritability probably captured by psoriasis genome-wide association study in Han Chinese. <i>Gene</i> , 2015, 573, 46-49.	2.2	6
41	Whole Exome Sequencing Enhanced Imputation Identifies 85 Metabolite Associations in the Alpine CHRIS Cohort. <i>Metabolites</i> , 2022, 12, 604.	2.9	6
42	Integration of expression quantitative trait loci and pleiotropy identifies a novel psoriasis susceptibility gene, <i>PTPN1</i> . <i>Journal of Gene Medicine</i> , 2017, 19, e2939.	2.8	5
43	A catalog of potential putative functional variants in psoriasis genome-wide association regions. <i>PLoS ONE</i> , 2018, 13, e0196635.	2.5	5
44	Identification of cell types, tissues and pathways affected by risk loci in psoriasis. <i>Molecular Genetics and Genomics</i> , 2016, 291, 1005-1012.	2.1	3
45	Rs4948496 within <i>ARID5B</i> gene is associated with clinical features of systemic lupus erythematosus in the Chinese Han population. <i>Journal of Dermatology</i> , 2015, 42, 608-612.	1.2	2
46	A rare variant in COL11A1 is strongly associated with adult height in Chinese Han population. <i>Journal of Genetics and Genomics</i> , 2016, 43, 549-554.	3.9	2
47	Discovery of Novel Genetic Risk Loci for Acute Central Serous Chorioretinopathy and Genetic Pleiotropic Effect With Age-Related Macular Degeneration. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 696885.	3.7	2
48	The Contribution of Meta-Analysis of Genome-Wide Association Studies in Investigating the Genetic Susceptibility to Lupus. <i>Journal of Investigative Dermatology Symposium Proceedings</i> , 2015, 17, 51-52.	0.8	0
49	Introduction of the Institute of Dermatology, Anhui Medical University. <i>Journal of Investigative Dermatology Symposium Proceedings</i> , 2015, 17, 8.	0.8	0