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List of articles citing

Identification of 38 novel loci for systemic lupus erythematosus and genetic heterogeneity between ancestral groups

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#	Paper	IF	Citations
75	snpXplorer: a web application to explore human SNP-associations and annotate SNP-sets. <i>Nucleic Acids Research</i> , 2021 , 49, W603-W612	20.1	3
74	Mergeomics 2.0: a web server for multi-omics data integration to elucidate disease networks and predict therapeutics. <i>Nucleic Acids Research</i> , 2021 , 49, W375-W387	20.1	4
73	Clinical and Immunological Biomarkers for Systemic Lupus Erythematosus. <i>Biomolecules</i> , 2021 , 11,	5.9	5
72	CRISPRa screen on a genetic risk locus shared by multiple autoimmune diseases identifies a dysfunctional enhancer that affects IRF8 expression through cooperative lncRNA and DNA methylation machinery.		
71	Shared genetic study gives insights into the shared and distinct pathogenic immunity components of IgA nephropathy and SLE. <i>Molecular Genetics and Genomics</i> , 2021 , 296, 1017-1026	3.1	0
70	The immunological significance of tumor necrosis factor receptor-associated factors (TRAFs). <i>International Immunology</i> , 2021 ,	4.9	6
69	Polygenic risk scores and rheumatic diseases. <i>Chinese Medical Journal</i> , 2021 , 134, 2521-2524	2.9	1
68	Lupus susceptibility region containing CTLA4 rs17268364 functionally reduces CTLA4 expression by binding EWSR1 and correlates IFN- β signature. <i>Arthritis Research and Therapy</i> , 2021 , 23, 279	5.7	0
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64	Deconvoluting the heterogeneity of SLE: The contribution of ancestry. <i>Journal of Allergy and Clinical Immunology</i> , 2021 ,	11.5	1
63	Systemic lupus erythematosus and immunodeficiency. <i>Rheumatology and Immunology Research</i> , 2021 , 2, 131-138	0.2	
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60	Immunogenetic Relationship of 14 bp Insertion/Deletion Polymorphism and Toll-Like Receptor 9 with Systemic Lupus Erythematosus in Egyptian Patients: A Case-Control Study.. <i>International Journal of General Medicine</i> , 2022 , 15, 661-674	2.3	0
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58	Genetic variants related to systemic lupus erythematosus revealed using bioinformatics. <i>European Journal of Inflammation</i> , 2022 , 20, 205873922110704	0.3	
57	A High Prevalence of Anti-EBNA1 Heteroantibodies in Systemic Lupus Erythematosus (SLE) Supports Anti-EBNA1 as an Origin for SLE Autoantibodies.. <i>Frontiers in Immunology</i> , 2022 , 13, 830993	8.4	○
56	Single-cell eQTL mapping identifies cell type-specific genetic control of autoimmune disease.. <i>Science</i> , 2022 , 376, eabf3041	33.3	8
55	Lupus enhancer risk variant causes dysregulation of IRF8 through cooperative lncRNA and DNA methylation machinery.. <i>Nature Communications</i> , 2022 , 13, 1855	17.4	○
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52	Systematic identification of genomic elements that regulate FCGR2A expression and harbor variants linked with autoimmune disease.. <i>Human Molecular Genetics</i> , 2021 ,	5.6	○
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- 19 The Identification of Significant Genes Related to Systemic Lupus Erythematosus through the Integration of the Results of a Transcriptome-Wide Association Study and an mRNA Expression Profile Analysis. ○
- 18 Interferon signature in systemic autoimmune diseases: what does it mean?. **2022**, 8, e002687 ○
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- 14 The potential causal association of systemic lupus erythematosus with congestive heart failure in the East Asian population: A two-sample mendelian randomization study. 096120332211498 ○
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- 11 Multi-ancestry and multi-trait genome-wide association meta-analyses inform clinical risk prediction for systemic lupus erythematosus. **2023**, 14, ○
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- 7 Extracellular vesicles and their cells of origin: Open issues in autoimmune diseases. 14, ○
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- 2 Trans-ethnic Mendelian randomization study of systemic lupus erythematosus and common female hormone-dependent malignancies. Publish Ahead of Print,
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