

# Yuan-Yuan Qi

## List of Publications by Year in descending order

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22  
papers

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citations

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677123

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22  
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times ranked

1151  
citing authors

#	ARTICLE	IF	CITATIONS
1	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
2	Increased autophagy is cytoprotective against podocyte injury induced by antibody and interferon- $\gamma$ in lupus nephritis. <i>Annals of the Rheumatic Diseases</i> , 2018, 77, 1799-1809.	0.9	79
3	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
4	Autophagy and immunological aberrations in systemic lupus erythematosus. <i>European Journal of Immunology</i> , 2019, 49, 523-533.	2.9	37
5	Autophagy-related gene <i>LRK2</i> is likely a susceptibility gene for systemic lupus erythematosus in northern Han Chinese. <i>Oncotarget</i> , 2017, 8, 13754-13761.	1.8	20
6	A Rare Variant (rs933717) at <i>FBXO31</i> MAP1LC3B in Chinese Is Associated With Systemic Lupus Erythematosus. <i>Arthritis and Rheumatology</i> , 2018, 70, 287-297.	5.6	18
7	Novel identified associations of RGS1 and RASGRP1 variants in IgA Nephropathy. <i>Scientific Reports</i> , 2016, 6, 35781.	3.3	17
8	Rare Variants of ATG5 Are Likely to Be Associated With Chinese Patients With Systemic Lupus Erythematosus. <i>Medicine (United States)</i> , 2015, 94, e939.	1.0	16
9	Cumulative Effects of Variants Identified by Genome-wide Association Studies in IgA Nephropathy. <i>Scientific Reports</i> , 2014, 4, 4904.	3.3	16
10	Polymorphism rs3828903 within <i>MICB</i> Is Associated with Susceptibility to Systemic Lupus Erythematosus in a Northern Han Chinese Population. <i>Journal of Immunology Research</i> , 2016, 2016, 1-6.	2.2	13
11	Albumin-to-globulin ratio (AGR) as a potential marker of predicting lupus nephritis in Chinese patients with systemic lupus erythematosus. <i>Lupus</i> , 2021, 30, 412-420.	1.6	13
12	Association of Variants in <i>CCR6</i> With Susceptibility to Lupus Nephritis in Chinese. <i>Arthritis and Rheumatology</i> , 2015, 67, 3091-3093.	5.6	12
13	Detecting Genetic Associations between <i>ATG5</i> and Lupus Nephritis by <i>trans</i> -eQTL. <i>Journal of Immunology Research</i> , 2015, 2015, 1-7.	2.2	12
14	Identification of a Hypoxia-Related Gene Model for Predicting the Prognosis and Formulating the Treatment Strategies in Kidney Renal Clear Cell Carcinoma. <i>Frontiers in Oncology</i> , 2021, 11, 806264.	2.8	8
15	Lupus susceptibility region containing CTLA4 rs17268364 functionally reduces CTLA4 expression by binding EWSR1 and correlates IFN- $\gamma$ signature. <i>Arthritis Research and Therapy</i> , 2021, 23, 279.	3.5	7
16	Elevated Plasma $\alpha$ -Defensins (HNP1-3) Levels Correlated with IgA1 Glycosylation and Susceptibility to IgA Nephropathy. <i>Disease Markers</i> , 2016, 2016, 1-7.	1.3	6
17	A Three Protein-Coding Gene Prognostic Model Predicts Overall Survival in Bladder Cancer Patients. <i>BioMed Research International</i> , 2020, 2020, 1-10.	1.9	6
18	The comprehensive analysis of clinical trials registration for IgA nephropathy therapy on ClinicalTrials.gov. <i>Renal Failure</i> , 2022, 44, 461-472.	2.1	6

#	ARTICLE	IF	CITATIONS
19	SCUBE3 Is Likely a Susceptibility Gene for Systemic Lupus Erythematosus for Chinese Populations. Journal of Immunology Research, 2020, 2020, 1-7.	2.2	3
20	The ZNF76 rs10947540 polymorphism associated with systemic lupus erythematosus risk in Chinese populations. Scientific Reports, 2021, 11, 5186.	3.3	2
21	Association of the PINX1 Variant rs6984094, Which Lengthens Telomeres, with Systemic Lupus Erythematosus Susceptibility in Chinese Populations. Journal of Immunology Research, 2021, 2021, 1-7.	2.2	2
22	Single Nucleotide Polymorphisms in PPAR $\alpha$ Associated with Systemic Lupus Erythematosus in Chinese Populations. Journal of Immunology Research, 2020, 2020, 1-7.	2.2	1