

# Yuan-Yuan Qi

## List of Publications by Year in Descending Order

**Source:** <https://exaly.com/author-pdf/5206473/yuan-yuan-qi-publications-by-year.pdf>  
**Version:** 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.  
The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

22 papers	349 citations	10 h-index	18 g-index
22 ext. papers	476 ext. citations	6.1 avg, IF	2.94 L-index

#	Paper	IF	Citations
22	The comprehensive analysis of clinical trials registration for IgA nephropathy therapy on ClinicalTrials.gov.. <i>Renal Failure</i> , <b>2022</b> , 44, 461-472	2.9	
21	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> , <b>2021</b> , 11, 806264	5.3	1
20	Lupus susceptibility region containing CTLA4 rs17268364 functionally reduces CTLA4 expression by binding EWSR1 and correlates IFN- $\beta$ signature. <i>Arthritis Research and Therapy</i> , <b>2021</b> , 23, 279	5.7	0
19	The ZNF76 rs10947540 polymorphism associated with systemic lupus erythematosus risk in Chinese populations. <i>Scientific Reports</i> , <b>2021</b> , 11, 5186	4.9	1
18	Association of the Variant rs6984094, Which Lengthens Telomeres, with Systemic Lupus Erythematosus Susceptibility in Chinese Populations. <i>Journal of Immunology Research</i> , <b>2021</b> , 2021, 7079359	4.5	0
17	Albumin-to-globulin ratio (AGR) as a potential marker of predicting lupus nephritis in Chinese patients with systemic lupus erythematosus. <i>Lupus</i> , <b>2021</b> , 30, 412-420	2.6	3
16	Single Nucleotide Polymorphisms in Associated with Systemic Lupus Erythematosus in Chinese Populations. <i>Journal of Immunology Research</i> , <b>2020</b> , 2020, 7285747	4.5	1
15	Is Likely a Susceptibility Gene for Systemic Lupus Erythematosus for Chinese Populations. <i>Journal of Immunology Research</i> , <b>2020</b> , 2020, 8897936	4.5	2
14	A Three Protein-Coding Gene Prognostic Model Predicts Overall Survival in Bladder Cancer Patients. <i>BioMed Research International</i> , <b>2020</b> , 2020, 7272960	3	3
13	Autophagy and immunological aberrations in systemic lupus erythematosus. <i>European Journal of Immunology</i> , <b>2019</b> , 49, 523-533	6.1	22
12	A Rare Variant (rs933717) at FBXO31-MAP1LC3B in Chinese Is Associated With Systemic Lupus Erythematosus. <i>Arthritis and Rheumatology</i> , <b>2018</b> , 70, 287-297	9.5	14
11	Increased autophagy is cytoprotective against podocyte injury induced by antibody and interferon- $\gamma$ in lupus nephritis. <i>Annals of the Rheumatic Diseases</i> , <b>2018</b> , 77, 1799-1809	2.4	35
10	Autophagy-related gene LRRK2 is likely a susceptibility gene for systemic lupus erythematosus in northern Han Chinese. <i>Oncotarget</i> , <b>2017</b> , 8, 13754-13761	3.3	12
9	Novel identified associations of RGS1 and RASGRP1 variants in IgA Nephropathy. <i>Scientific Reports</i> , <b>2016</b> , 6, 35781	4.9	9
8	High-density genotyping of immune-related loci identifies new SLE risk variants in individuals with Asian ancestry. <i>Nature Genetics</i> , <b>2016</b> , 48, 323-30	36.3	156
7	Elevated Plasma $\beta$ -Defensins (HNP1-3) Levels Correlated with IgA1 Glycosylation and Susceptibility to IgA Nephropathy. <i>Disease Markers</i> , <b>2016</b> , 2016, 8123138	3.2	5
6	Polymorphism rs3828903 within MICB Is Associated with Susceptibility to Systemic Lupus Erythematosus in a Northern Han Chinese Population. <i>Journal of Immunology Research</i> , <b>2016</b> , 2016, 1343780	4.5	4

5	Association of variants in CCR6 with susceptibility to lupus nephritis in Chinese. <i>Arthritis and Rheumatology</i> , <b>2015</b> , 67, 3091-3	9.5	10
4	Rare Variants of ATG5 Are Likely to Be Associated With Chinese Patients With Systemic Lupus Erythematosus. <i>Medicine (United States)</i> , <b>2015</b> , 94, e939	1.8	14
3	Detecting Genetic Associations between ATG5 and Lupus Nephritis by trans-eQTL. <i>Journal of Immunology Research</i> , <b>2015</b> , 2015, 153132	4.5	11
2	Cumulative effects of variants identified by genome-wide association studies in IgA nephropathy. <i>Scientific Reports</i> , <b>2014</b> , 4, 4904	4.9	13
1	Brief Report: identification of MTMR3 as a novel susceptibility gene for lupus nephritis in northern Han Chinese by shared-gene analysis with IgA nephropathy. <i>Arthritis and Rheumatology</i> , <b>2014</b> , 66, 2842-8	9.5	33