

Kwangwoo Kim

List of Publications by Year in descending order

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Version: 2024-02-01

43
papers

1,732
citations

361296

20
h-index

302012

39
g-index

46
all docs

46
docs citations

46
times ranked

3101
citing authors

#	ARTICLE	IF	CITATIONS
1	Novel susceptibility loci for steroid-associated osteonecrosis of the femoral head in systemic lupus erythematosus. <i>Human Molecular Genetics</i> , 2022, 31, 1082-1095.	1.4	1
2	Recent advances in understanding the genetic basis of systemic lupus erythematosus. <i>Seminars in Immunopathology</i> , 2022, 44, 29-46.	2.8	27
3	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.	0.5	9
4	Allele-specific Quantification of HLA-DRB1 Transcripts Reveals Imbalanced Allelic Expression That Modifies the Amino Acid Effects in HLA-DR β 1. <i>Arthritis and Rheumatology</i> , 2021, 73, 381-391.	2.9	4
5	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.	0.5	103
6	Large-scale meta-analysis across East Asian and European populations updated genetic architecture and variant-driven biology of rheumatoid arthritis, identifying 11 novel susceptibility loci. <i>Annals of the Rheumatic Diseases</i> , 2021, 80, 558-565.	0.5	93
7	Genetic variants shape rheumatoid arthritis-specific transcriptomic features in CD4 ⁺ T cells through differential DNA methylation, explaining a substantial proportion of heritability. <i>Annals of the Rheumatic Diseases</i> , 2021, 80, 876-883.	0.5	12
8	Clinical and Genetic Risk Factors Associated With the Presence of Lupus Nephritis. <i>Journal of Rheumatic Diseases</i> , 2021, 28, 150-158.	0.4	7
9	Identifying damage clusters in patients with systemic lupus erythematosus. <i>International Journal of Rheumatic Diseases</i> , 2020, 23, 84-91.	0.9	6
10	Host Genetic and Gut Microbial Signatures in Familial Inflammatory Bowel Disease. <i>Clinical and Translational Gastroenterology</i> , 2020, 11, e00213.	1.3	9
11	Genome-wide association study in a Korean population identifies six novel susceptibility loci for rheumatoid arthritis. <i>Annals of the Rheumatic Diseases</i> , 2020, 79, 1438-1445.	0.5	26
12	MHC associations of ankylosing spondylitis in East Asians are complex and involve non-HLA-B27 HLA contributions. <i>Arthritis Research and Therapy</i> , 2020, 22, 74.	1.6	13
13	Allelic frequency differences of DAOA variants between Caucasians and Asians and their association with major mood disorders. <i>Signal Transduction and Targeted Therapy</i> , 2019, 4, 39.	7.1	1
14	Deletion at 2q14.3 is associated with worse response to TNF- α blockers in patients with rheumatoid arthritis. <i>Arthritis Research and Therapy</i> , 2019, 21, 195.	1.6	1
15	Update on the Genetics of Systemic Lupus Erythematosus: Genome-Wide Association Studies and Beyond. <i>Cells</i> , 2019, 8, 1180.	1.8	93
16	Amino acid signatures of HLA Class-I and II molecules are strongly associated with SLE susceptibility and autoantibody production in Eastern Asians. <i>PLoS Genetics</i> , 2019, 15, e1008092.	1.5	36
17	Understanding HLA associations from SNP summary association statistics. <i>Scientific Reports</i> , 2019, 9, 1337.	1.6	9
18	Genetic variants differentially associated with rheumatoid arthritis and systemic lupus erythematosus reveal the disease-specific biology. <i>Scientific Reports</i> , 2019, 9, 2739.	1.6	13

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19	135â€¦Influence of genetic variants on avascular necrosis in patients with systemic lupus erythematosus. , 2019, , .		0
20	254â€¦Identification of damage clusters in systemic lupus erythematosus. , 2019, , .		0
21	136â€¦Influence of genetic risk variants on the clinical subphenotypes of systemic lupus erythematosus in a korean cohort. , 2019, , .		0
22	Massive false-positive geneâ€“gene interactions by Rothmanâ€™s additive model. Annals of the Rheumatic Diseases, 2019, 78, 437-439.	0.5	6
23	Association of CD8⁺ Tâ€™cells with bone erosion in patients with rheumatoid arthritis. International Journal of Rheumatic Diseases, 2018, 21, 440-446.	0.9	7
24	Genetic variants in systemic lupus erythematosus susceptibility loci, XKR6 and GLT1D1 are associated with childhood-onset SLE in a Korean cohort. Scientific Reports, 2018, 8, 9962.	1.6	25
25	A missense variant in NCF1 is associated with susceptibility to multiple autoimmune diseases. Nature Genetics, 2017, 49, 433-437.	9.4	143
26	Confirmation of five novel susceptibility loci for Systemic Lupus Erythematosus (SLE) and integrated network analysis of 82 SLE susceptibility loci. Human Molecular Genetics, 2017, 26, ddx026.	1.4	47
27	Update on the genetic architecture of rheumatoid arthritis. Nature Reviews Rheumatology, 2017, 13, 13-24.	3.5	102
28	Biological function integrated prediction of severe radiographic progression in rheumatoid arthritis: a nested case control study. Arthritis Research and Therapy, 2017, 19, 244.	1.6	11
29	Identification of a Systemic Lupus Erythematosus Risk Locus Spanning <i>ATG16L2</i>, <i>FCHSD2</i>, and <i>P2RY2</i> in Koreans. Arthritis and Rheumatology, 2016, 68, 1197-1209.	2.9	89
30	Association of Single Nucleotide Polymorphisms of PADI4 and HLA-DRB1 Alleles with Susceptibility to Rheumatoid Arthritis-Related Lung Diseases. Lung, 2016, 194, 745-753.	1.4	17
31	Association-heterogeneity mapping identifies an Asian-specific association of the GTF2I locus with rheumatoid arthritis. Scientific Reports, 2016, 6, 27563.	1.6	23
32	High-density genotyping of immune-related loci identifies new SLE risk variants in individuals with Asian ancestry. Nature Genetics, 2016, 48, 323-330.	9.4	219
33	Response to Intravenous Cyclophosphamide Treatment for Lupus Nephritis Associated with Polymorphisms in the <i>FCGR2B-FCRLA</i> Locus. Journal of Rheumatology, 2016, 43, 1045-1049.	1.0	15
34	Imputing Variants in HLA-DR Beta Genes Reveals That HLA-DRB1 Is Solely Associated with Rheumatoid Arthritis and Systemic Lupus Erythematosus. PLoS ONE, 2016, 11, e0150283.	1.1	20
35	An HLA-C amino-acid variant in addition to HLA-B*27 confers risk for ankylosing spondylitis in the Korean population. Arthritis Research and Therapy, 2015, 17, 342.	1.6	21
36	Interactions Between Amino Acidâ€“Defined Major Histocompatibility Complex Class II Variants and Smoking in Seropositive Rheumatoid Arthritis. Arthritis and Rheumatology, 2015, 67, 2611-2623.	2.9	58

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37	High-density genotyping of immune loci in Koreans and Europeans identifies eight new rheumatoid arthritis risk loci. <i>Annals of the Rheumatic Diseases</i> , 2015, 74, e13-e13.	0.5	100
38	Construction and Application of a Korean Reference Panel for Imputing Classical Alleles and Amino Acids of Human Leukocyte Antigen Genes. <i>PLoS ONE</i> , 2014, 9, e112546.	1.1	27
39	The HLA-DR β 1 amino acid positions 11-13-26 explain the majority of SLE-MHC associations. <i>Nature Communications</i> , 2014, 5, 5902.	5.8	80
40	Development of rheumatoid arthritis specific HLA-DRB1 genotyping microarray. <i>Biochip Journal</i> , 2014, 8, 187-198.	2.5	1
41	Risk for ACPA-positive rheumatoid arthritis is driven by shared HLA amino acid polymorphisms in Asian and European populations. <i>Human Molecular Genetics</i> , 2014, 23, 6916-6926.	1.4	135
42	Ethnic specificity of lupus-associated loci identified in a genome-wide association study in Korean women. <i>Annals of the Rheumatic Diseases</i> , 2014, 73, 1240-1245.	0.5	61
43	Variation in the ICAM1-ICAM4-ICAM5 locus is associated with systemic lupus erythematosus susceptibility in multiple ancestries. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 1809-1814.	0.5	60