

# Jae-Jung Kim

## List of Publications by Year in descending order

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Version: 2024-02-01

21  
papers

552  
citations

840585

11  
h-index

794469

19  
g-index

21  
all docs

21  
docs citations

21  
times ranked

1095  
citing authors

#	ARTICLE	IF	CITATIONS
1	Diagnosis of Central Sensitization and Its Effects on Postoperative Outcomes following Total Knee Arthroplasty: A Systematic Review and Meta-Analysis. <i>Diagnostics</i> , 2022, 12, 1248.	1.3	11
2	Association of an IGHV3-66 gene variant with Kawasaki disease. <i>Journal of Human Genetics</i> , 2021, 66, 475-489.	1.1	27
3	IgA Levels Are Associated with Coronary Artery Lesions in Kawasaki Disease. <i>Korean Circulation Journal</i> , 2021, 51, 267.	0.7	12
4	Identification of rare coding variants associated with Kawasaki disease by whole exome sequencing. <i>Genomics and Informatics</i> , 2021, 19, e38.	0.4	3
5	Identification of SAMD9L as a susceptibility locus for intravenous immunoglobulin resistance in Kawasaki disease by genome-wide association analysis. <i>Pharmacogenomics Journal</i> , 2020, 20, 80-86.	0.9	9
6	Association of the IL16 Asn1147Lys polymorphism with intravenous immunoglobulin resistance in Kawasaki disease. <i>Journal of Human Genetics</i> , 2020, 65, 421-426.	1.1	3
7	Assessment of the Clinical Heterogeneity of Kawasaki Disease Using Genetic Variants of <i>BLK</i> and <i>FCGR2A</i> . <i>Korean Circulation Journal</i> , 2019, 49, 99.	0.7	6
8	Identification of the TIFAB Gene as a Susceptibility Locus for Coronary Artery Aneurysm in Patients with Kawasaki Disease. <i>Pediatric Cardiology</i> , 2019, 40, 483-488.	0.6	14
9	<i>BCL2L1</i> Is Associated With Kawasaki Disease in Intravenous Immunoglobulin Responder Patients. <i>Circulation Genomic and Precision Medicine</i> , 2018, 11, e002020.	1.6	12
10	A genome-wide association analysis identifies NMNAT2 and HCP5 as susceptibility loci for Kawasaki disease. <i>Journal of Human Genetics</i> , 2017, 62, 1023-1029.	1.1	40
11	Male-specific association of the FCGR2A His167Arg polymorphism with Kawasaki disease. <i>PLoS ONE</i> , 2017, 12, e0184248.	1.1	33
12	Common Variants in the CRP Promoter are Associated with a High C-Reactive Protein Level in Kawasaki Disease. <i>Pediatric Cardiology</i> , 2015, 36, 438-444.	0.6	20
13	Identification of KCNN2 as a susceptibility locus for coronary artery aneurysms in Kawasaki disease using genome-wide association analysis. <i>Journal of Human Genetics</i> , 2013, 58, 521-525.	1.1	32
14	Assessment of Risk Factors for Korean Children with Kawasaki Disease. <i>Pediatric Cardiology</i> , 2012, 33, 513-520.	0.6	49
15	Exome sequencing and subsequent association studies identify five amino acid-altering variants influencing human height. <i>Human Genetics</i> , 2012, 131, 471-478.	1.8	33
16	A genome-wide association analysis reveals 1p31 and 2p13.3 as susceptibility loci for Kawasaki disease. <i>Human Genetics</i> , 2011, 129, 487-495.	1.8	79
17	Association of a c.1084A>G (p.Thr362Ala) Variant in the DCTN4 Gene with Wilson Disease. <i>Journal of Genetic Medicine</i> , 2011, 8, 53-57.	0.1	0
18	Development of SNP-based human identification system. <i>International Journal of Legal Medicine</i> , 2010, 124, 125-131.	1.2	33

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19	Identification of 15 loci influencing height in a Korean population. <i>Journal of Human Genetics</i> , 2010, 55, 27-31.	1.1	101
20	Genetic variants in the HLA-G region are associated with Kawasaki disease. <i>Human Immunology</i> , 2008, 69, 867-871.	1.2	35
21	Some Trials of L.F.A. for Comminuted Intertrochanteric Fractures (Hip). <i>The Journal of the Korean Orthopaedic Association</i> , 1974, 9, 374.	0.0	0