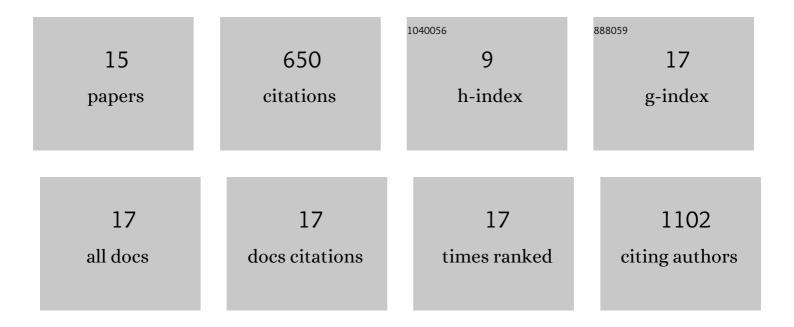
Kicheol Kim

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Gut microbiota–specific IgA ⁺ B cells traffic to the CNS in active multiple sclerosis. Science Immunology, 2020, 5, .	11.9	132
2	Aberrant oligodendroglial–vascular interactions disrupt the blood–brain barrier, triggering CNS inflammation. Nature Neuroscience, 2019, 22, 709-718.	14.8	131
3	A pathogenic and clonally expanded B cell transcriptome in active multiple sclerosis. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 22932-22943.	7.1	119
4	Selective Estrogen Receptor Modulators Enhance CNS Remyelination Independent of Estrogen Receptors. Journal of Neuroscience, 2019, 39, 2184-2194.	3.6	49
5	Population genetic study for 24 STR loci and Y indel (GlobalFilerâ,,¢ PCR Amplification kit and) Tj ETQq1 1 0.7843	14.rgBT /C 1.3	Overlock 10
6	Early complement genes are associated with visual system degeneration in multiple sclerosis. Brain, 2019, 142, 2722-2736.	7.6	30
7	Forensic and population genetic analyses of the GlobalFiler STR loci in the Mongolian population. Genes and Genomics, 2017, 39, 423-431.	1.4	22
8	Cell type-specific transcriptomics identifies neddylation as a novel therapeutic target in multiple sclerosis. Brain, 2021, 144, 450-461.	7.6	16
9	Distinctive waves of innate immune response in the retina in experimental autoimmune encephalomyelitis. JCI Insight, 2021, 6, .	5.0	14
10	Mononuclear cell transcriptome changes associated with dimethyl fumarate in MS. Neurology: Neuroimmunology and NeuroInflammation, 2018, 5, e470.	6.0	8
11	Evaluation of forensic genetic parameters of 12 STR loci in the Korean population using the InvestigatorⓇ HDplex kit. International Journal of Legal Medicine, 2017, 131, 1247-1249.	2.2	6
12	Association of mitochondrial haplogroup F with physical performance in Korean population. Genomics and Informatics, 2019, 17, e11.	0.8	6
13	Allele frequency data of 20 STR loci in 2000 Korean individuals. Forensic Science International: Genetics Supplement Series, 2017, 6, e65-e68.	0.3	2
14	Association of glutathione S-transferase M1 and T1 null/present polymorphism with physical performance in the Korean population. Genes and Genomics, 2019, 41, 71-78.	1.4	2
15	Comparative analysis of allele variation using allele frequencies according to sample size in Korean population. Genes and Genomics, 2021, 43, 1301-1305.	1.4	2