Xiaoming Jia

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Exome sequencing in bipolar disorder identifies AKAP11 as a risk gene shared with schizophrenia. Nature Genetics, 2022, 54, 541-547.	21.4	65
2	Household paired design reduces variance and increases power in multi-city gut microbiome study in multiple sclerosis. Multiple Sclerosis Journal, 2021, 27, 366-379.	3.0	24
3	Accurate imputation of human leukocyte antigens with CookHLA. Nature Communications, 2021, 12, 1264.	12.8	21
4	Investigating rare pathogenic/likely pathogenic exonic variation in bipolar disorder. Molecular Psychiatry, 2021, 26, 5239-5250.	7.9	15
5	Genome sequencing uncovers phenocopies in primary progressive multiple sclerosis. Annals of Neurology, 2018, 84, 51-63.	5.3	38
6	Gut microbiota from multiple sclerosis patients enables spontaneous autoimmune encephalomyelitis in mice. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 10719-10724.	7.1	666
7	Common coding variants in the HLA-DQB1 region confer susceptibility to age-related macular degeneration. European Journal of Human Genetics, 2016, 24, 1049-1055.	2.8	13
8	Coding Variants at Hexa-allelic Amino Acid 13 of HLA-DRB1 Explain Independent SNP Associations with Follicular Lymphoma Risk. American Journal of Human Genetics, 2013, 93, 167-172.	6.2	26
9	Imputing Amino Acid Polymorphisms in Human Leukocyte Antigens. PLoS ONE, 2013, 8, e64683.	2.5	538
10	Fine-mapping classical HLA variation associated with durable host control of HIV-1 infection in African Americans. Human Molecular Genetics, 2012, 21, 4334-4347.	2.9	61
11	Five amino acids in three HLA proteins explain most of the association between MHC and seropositive rheumatoid arthritis. Nature Genetics, 2012, 44, 291-296.	21.4	768
12	HLA-A*3101 and Carbamazepine-Induced Hypersensitivity Reactions in Europeans. New England Journal of Medicine, 2011, 364, 1134-1143.	27.0	815
13	Next-generation sequencing for HLA typing of class I loci. BMC Genomics, 2011, 12, 42.	2.8	135
14	Integrating common and rare genetic variation in diverse human populations. Nature, 2010, 467, 52-58.	27.8	2,625
15	The Major Genetic Determinants of HIV-1 Control Affect HLA Class I Peptide Presentation. Science, 2010, 330, 1551-1557.	12.6	1,054
16	Meta-analysis of genome scans and replication identify CD6, IRF8 and TNFRSF1A as new multiple sclerosis susceptibility loci. Nature Genetics, 2009, 41, 776-782.	21.4	729
17	Practical aspects of imputation-driven meta-analysis of genome-wide association studies. Human Molecular Genetics, 2008, 17, R122-R128.	2.9	475