Max Lam

List of Publications by Year
in descending order

25 papers	6,252 citations	471509 17 h-index	552781 26 g-index
39 all docs	39 docs citations	39 times ranked	9020 citing authors

#	Article	IF	Citations
1	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	27.8	929
2	Improving polygenic prediction in ancestrally diverse populations. Nature Genetics, 2022, 54, 573-580.	21.4	209
3	Novel ultra-rare exonic variants identified in a founder population implicate cadherins in schizophrenia. Neuron, 2021, 109, 1465-1478.e4.	8.1	21
4	Identifying nootropic drug targets via large-scale cognitive GWAS and transcriptomics. Neuropsychopharmacology, 2021, 46, 1788-1801.	5.4	12
5	Genome wide study of tardive dyskinesia in schizophrenia. Translational Psychiatry, 2021, 11, 351.	4.8	13
6	Large-scale evaluation of the Positive and Negative Syndrome Scale (PANSS) symptom architecture in schizophrenia. Asian Journal of Psychiatry, 2021, 62, 102732.	2.0	29
7	RICOPILI: Rapid Imputation for COnsortias PIpeLIne. Bioinformatics, 2020, 36, 930-933.	4.1	201
8	Genetic liability in individuals at ultra-high risk of psychosis: A comparison study of 9 psychiatric traits. PLoS ONE, 2020, 15, e0243104.	2.5	3
9	Pleiotropic Meta-Analysis of Cognition, Education, and Schizophrenia Differentiates Roles of Early Neurodevelopmental and Adult Synaptic Pathways. American Journal of Human Genetics, 2019, 105, 334-350.	6.2	86
10	Genome-wide Association Studies in Ancestrally Diverse Populations: Opportunities, Methods, Pitfalls, and Recommendations. Cell, 2019, 179, 589-603.	28.9	428
11	Comparative genetic architectures of schizophrenia in East Asian and European populations. Nature Genetics, 2019, 51, 1670-1678.	21.4	440
12	Screening Human Embryos for Polygenic Traits Has Limited Utility. Cell, 2019, 179, 1424-1435.e8.	28.9	78
13	Study of 300,486 individuals identifies 148 independent genetic loci influencing general cognitive function. Nature Communications, 2018, 9, 2098.	12.8	484
14	Factor structure of the positive and negative syndrome scale (PANSS) in people at ultra high risk (UHR) for psychosis. Schizophrenia Research, 2018, 201, 85-90.	2.0	11
15	Genome-wide association meta-analysis in 269,867 individuals identifies new genetic and functional links to intelligence. Nature Genetics, 2018, 50, 912-919.	21.4	893
16	Gene discovery and polygenic prediction from a genome-wide association study of educational attainment in 1.1 million individuals. Nature Genetics, 2018, 50, 1112-1121.	21.4	1,835
17	Longitudinal Cognitive Changes in Young Individuals at Ultrahigh Risk for Psychosis. JAMA Psychiatry, 2018, 75, 929.	11.0	54
18	Establishing the Brief Assessment of Cognition - Short form. Journal of Psychiatric Research, 2017, 93, 1-11.	3.1	6

#	Article	IF	Citations
19	Identification of Genetic Loci Jointly Influencing Schizophrenia Risk and the Cognitive Traits of Verbal-Numerical Reasoning, Reaction Time, and General Cognitive Function. JAMA Psychiatry, 2017, 74, 1065.	11.0	123
20	Large-Scale Cognitive GWAS Meta-Analysis Reveals Tissue-Specific Neural Expression and Potential Nootropic Drug Targets. Cell Reports, 2017, 21, 2597-2613.	6.4	103
21	Baseline social amotivation predicts 1-year functioning in UHR subjects: A validation and prospective investigation. European Neuropsychopharmacology, 2015, 25, 2187-2196.	0.7	19
22	Impact of psychiatric comorbidity in individuals at Ultra High Risk of psychosis â€" Findings from the Longitudinal Youth at Risk Study (LYRIKS). Schizophrenia Research, 2015, 164, 8-14.	2.0	94
23	The Continuous Performance Test, Identical Pairs: norms, reliability and performance in healthy controls and patients with schizophrenia in Singapore. Schizophrenia Research, 2014, 156, 233-240.	2.0	10
24	The Longitudinal Youth at Risk Study (LYRIKS) â€" An Asian UHR perspective. Schizophrenia Research, 2013, 151, 279-283.	2.0	46
25	Brief Assessment of Cognition in Schizophrenia: Normative Data in an English-Speaking Ethnic Chinese Sample. Archives of Clinical Neuropsychology, 2013, 28, 845-858.	0.5	14