Robert M Maier

List of Publications by Year
in descending order

13 papers	4,046 citations	840776 11 h-index	996975 15 g-index
20 all docs	20 docs citations	20 times ranked	9274 citing authors

#	Article	IF	Citations
1	An integrative skeletal and paleogenomic analysis of stature variation suggests relatively reduced health for early European farmers. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e2106743119.	7.1	21
2	Response to Comment on â€œLarge-scale GWAS reveals insights into the genetic architecture of same-sex sexual behaviorâ€• Science, 2021, 371, .	12.6	5
3	Genomic evidence consistent with antagonistic pleiotropy may help explain the evolutionary maintenance of same-sex sexual behaviour in humans. Nature Human Behaviour, 2021, 5, 1251-1258.	12.0	27
4	The Genetics of the Mood Disorder Spectrum: Genome-wide Association Analyses of More Than 185,000 Cases and 439,000 Controls. Biological Psychiatry, 2020, 88, 169-184.	1.3	137
5	No statistical evidence for an effect of CCR5-â^†32 on lifespan in the UK Biobank cohort. Nature Medicine, 2020, 26, 178-180.	30.7	16
6	Genetic contributions to variation in human stature in prehistoric Europe. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 21484-21492.	7.1	64
7	Large-scale GWAS reveals insights into the genetic architecture of same-sex sexual behavior. Science, 2019, 365, .	12.6	245
8	Polygenic adaptation on height is overestimated due to uncorrected stratification in genome-wide association studies. ELife, 2019, 8, .	6.0	276
9	Genome studies must account for historyâ€"Response. Science, 2019, 366, 1461-1462.	12.6	4
10	Improving genetic prediction by leveraging genetic correlations among human diseases and traits. Nature Communications, 2018, 9, 989.	12.8	136
11	Causal associations between risk factors and common diseases inferred from GWAS summary data. Nature Communications, 2018, 9, 224.	12.8	629
12	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. Nature Genetics, 2018, 50, 668-681.	21.4	2,224
13	Joint Analysis of Psychiatric Disorders Increases Accuracy of Risk Prediction for Schizophrenia, Bipolar Disorder, and Major Depressive Disorder. American Journal of Human Genetics, 2015, 96, 283-294.	6.2	225