

Ronald de Vlaming

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/677673/publications.pdf>

Version: 2024-02-01

17
papers

4,824
citations

516710

16
h-index

794594

19
g-index

28
all docs

28
docs citations

28
times ranked

8207
citing authors

#	ARTICLE	IF	CITATIONS
1	Polygenic prediction of educational attainment within and between families from genome-wide association analyses in 3 million individuals. <i>Nature Genetics</i> , 2022, 54, 437-449.	21.4	215
2	Multivariate GWAS of psychiatric disorders and their cardinal symptoms reveal two dimensions of cross-cutting genetic liabilities. <i>Cell Genomics</i> , 2022, 2, 100140.	6.5	32
3	Genomic analysis of diet composition finds novel loci and associations with health and lifestyle. <i>Molecular Psychiatry</i> , 2021, 26, 2056-2069.	7.9	79
4	Multivariate analysis of 1.5 million people identifies genetic associations with traits related to self-regulation and addiction. <i>Nature Neuroscience</i> , 2021, 24, 1367-1376.	14.8	137
5	Multivariate analysis reveals shared genetic architecture of brain morphology and human behavior. <i>Communications Biology</i> , 2021, 4, 1180.	4.4	7
6	Mendelian randomization: the challenge of unobserved environmental confounds. <i>International Journal of Epidemiology</i> , 2019, 48, 665-671.	1.9	56
7	Genomic structural equation modelling provides insights into the multivariate genetic architecture of complex traits. <i>Nature Human Behaviour</i> , 2019, 3, 513-525.	12.0	511
8	Genome-wide association analyses of risk tolerance and risky behaviors in over 1 million individuals identify hundreds of loci and shared genetic influences. <i>Nature Genetics</i> , 2019, 51, 245-257.	21.4	536
9	Signatures of negative selection in the genetic architecture of human complex traits. <i>Nature Genetics</i> , 2018, 50, 746-753.	21.4	304
10	Genome-wide association study results for educational attainment aid in identifying genetic heterogeneity of schizophrenia. <i>Nature Communications</i> , 2018, 9, 3078.	12.8	64
11	Hidden heritability due to heterogeneity across seven populations. <i>Nature Human Behaviour</i> , 2017, 1, 757-765.	12.0	137
12	Meta-GWAS Accuracy and Power (MetaGAP) Calculator Shows that Hiding Heritability Is Partially Due to Imperfect Genetic Correlations across Studies. <i>PLoS Genetics</i> , 2017, 13, e1006495.	3.5	78
13	Genetic variants associated with subjective well-being, depressive symptoms, and neuroticism identified through genome-wide analyses. <i>Nature Genetics</i> , 2016, 48, 624-633.	21.4	870
14	Genome-wide association study identifies 74 loci associated with educational attainment. <i>Nature</i> , 2016, 533, 539-542.	27.8	1,204
15	Genome-wide analysis identifies 12 loci influencing human reproductive behavior. <i>Nature Genetics</i> , 2016, 48, 1462-1472.	21.4	284
16	Genetic variants linked to education predict longevity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 13366-13371.	7.1	110
17	The Current and Future Use of Ridge Regression for Prediction in Quantitative Genetics. <i>BioMed Research International</i> , 2015, 2015, 1-18.	1.9	67