

Brendan Bulik-Sullivan

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11155749/publications.pdf>

Version: 2024-02-01

17
papers

11,315
citations

516710

16
h-index

888059

17
g-index

22
all docs

22
docs citations

22
times ranked

19185
citing authors

#	ARTICLE	IF	CITATIONS
1	An atlas of genetic correlations across human diseases and traits. <i>Nature Genetics</i> , 2015, 47, 1236-1241.	21.4	3,145
2	Partitioning heritability by functional annotation using genome-wide association summary statistics. <i>Nature Genetics</i> , 2015, 47, 1228-1235.	21.4	2,045
3	Modeling Linkage Disequilibrium Increases Accuracy of Polygenic Risk Scores. <i>American Journal of Human Genetics</i> , 2015, 97, 576-592.	6.2	1,098
4	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , 2018, 360, .	12.6	1,085
5	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. <i>Nature Genetics</i> , 2017, 49, 27-35.	21.4	838
6	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. <i>Cell</i> , 2018, 173, 1705-1715.e16.	28.9	623
7	Partitioning Heritability of Regulatory and Cell-Type-Specific Variants across 11 Common Diseases. <i>American Journal of Human Genetics</i> , 2014, 95, 535-552.	6.2	569
8	Significant Locus and Metabolic Genetic Correlations Revealed in Genome-Wide Association Study of Anorexia Nervosa. <i>American Journal of Psychiatry</i> , 2017, 174, 850-858.	7.2	410
9	Linkage disequilibrium-dependent architecture of human complex traits shows action of negative selection. <i>Nature Genetics</i> , 2017, 49, 1421-1427.	21.4	400
10	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. <i>Nature Genetics</i> , 2015, 47, 1294-1303.	21.4	357
11	Genetic risk for autism spectrum disorders and neuropsychiatric variation in the general population. <i>Nature Genetics</i> , 2016, 48, 552-555.	21.4	326
12	Estimation of Genetic Correlation via Linkage Disequilibrium Score Regression and Genomic Restricted Maximum Likelihood. <i>American Journal of Human Genetics</i> , 2018, 102, 1185-1194.	6.2	119
13	Shared genetic aetiology of puberty timing between sexes and with health-related outcomes. <i>Nature Communications</i> , 2015, 6, 8842.	12.8	100
14	Functional Architectures of Local and Distal Regulation of Gene Expression in Multiple Human Tissues. <i>American Journal of Human Genetics</i> , 2017, 100, 605-616.	6.2	76
15	Quantifying the Genetic Correlation between Multiple Cancer Types. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 1427-1435.	2.5	48
16	Prioritization of Genetic Variants in the micro RNA Regulome as Functional Candidates in Genome-Wide Association Studies. <i>Human Mutation</i> , 2013, 34, 1049-1056.	2.5	33
17	Large-Scale Genomic Analyses Link Reproductive Aging to Hypothalamic Signaling, Breast Cancer Susceptibility, and BRCA1-Mediated DNA Repair. <i>Obstetrical and Gynecological Survey</i> , 2015, 70, 758-762.	0.4	0