Daniel J Benjamin

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3660970/publications.pdf

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182225 340414 38 11,226 30 citations h-index papers

39 g-index 46 46 46 17775 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Polygenic prediction of educational attainment within and between families from genome-wide association analyses in 3 million individuals. Nature Genetics, 2022, 54, 437-449.	9.4	215
2	Within-sibship genome-wide association analyses decrease bias in estimates of direct genetic effects. Nature Genetics, 2022, 54, 581-592.	9.4	142
3	Mendelian imputation of parental genotypes improves estimates of direct genetic effects. Nature Genetics, 2022, 54, 897-905.	9.4	31
4	Genomic analysis of diet composition finds novel loci and associations with health and lifestyle. Molecular Psychiatry, 2021, 26, 2056-2069.	4.1	79
5	Resource profile and user guide of the Polygenic Index Repository. Nature Human Behaviour, 2021, 5, 1744-1758.	6.2	63
6	Problems with Using Polygenic Scores to Select Embryos. New England Journal of Medicine, 2021, 385, 78-86.	13.9	105
7	Consensus-based guidance for conducting and reporting multi-analyst studies. ELife, 2021, 10, .	2.8	22
8	Predicting mid-life capital formation with pre-school delay of gratification and life-course measures of self-regulation. Journal of Economic Behavior and Organization, 2020, 179, 743-756.	1.0	16
9	A consensus-based transparency checklist. Nature Human Behaviour, 2020, 4, 4-6.	6.2	79
10	Three Recommendations for Improving the Use of $\langle i \rangle p \langle i \rangle$ -Values. American Statistician, 2019, 73, 186-191.	0.9	152
11	Errors in probabilistic reasoning and judgment biases. Handbook of Behavioral Economics, 2019, 2, 69-186.	3.7	89
12	Genome-wide association analyses of risk tolerance and risky behaviors in over 1 million individuals identify hundreds of loci and shared genetic influences. Nature Genetics, 2019, 51, 245-257.	9.4	536
13	No Evidence That Experiencing Physical Warmth Promotes Interpersonal Warmth. Social Psychology, 2019, 50, 127-132.	0.3	31
14	Multi-trait analysis of genome-wide association summary statistics using MTAG. Nature Genetics, 2018, 50, 229-237.	9.4	700
15	Redefine statistical significance. Nature Human Behaviour, 2018, 2, 6-10.	6.2	1,763
16	Imprint of assortative mating on the human genome. Nature Human Behaviour, 2018, 2, 948-954.	6.2	97
17	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.	9.4	1,835
18	The relationship between the normalized gradient addition mechanism and quadratic voting. Public Choice, 2017, 172, 233-263.	1.0	1

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19	A MODEL OF NONBELIEF IN THE LAW OF LARGE NUMBERS. Journal of the European Economic Association, 2016, 14, 515-544.	1.9	56
20	Genetic variants associated with subjective well-being, depressive symptoms, and neuroticism identified through genome-wide analyses. Nature Genetics, 2016, 48, 624-633.	9.4	870
21	Genome-wide association study identifies 74 loci associated with educational attainment. Nature, 2016, 533, 539-542.	13.7	1,204
22	Genome-wide analysis identifies 12 loci influencing human reproductive behavior. Nature Genetics, 2016, 48, 1462-1472.	9.4	284
23	Polygenic risk scores for schizophrenia and bipolar disorder predict creativity. Nature Neuroscience, 2015, 18, 953-955.	7.1	351
24	Distributional Preferences, Reciprocity-Like Behavior, and Efficiency in Bilateral Exchange. American Economic Journal: Microeconomics, 2015, 7, 70-98.	0.7	9
25	Directional dominance on stature and cognition inÂdiverse human populations. Nature, 2015, 523, 459-462.	13.7	173
26	Beyond Happiness and Satisfaction: Toward Well-Being Indices Based on Stated Preference. American Economic Review, 2014, 104, 2698-2735.	4.0	185
27	Can Marginal Rates of Substitution Be Inferred from Happiness Data? Evidence from Residency Choices. American Economic Review, 2014, 104, 3498-3528.	4.0	118
28	Common genetic variants associated with cognitive performance identified using the proxy-phenotype method. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 13790-13794.	3.3	244
29	Replicability and Robustness of Genome-Wide-Association Studies for Behavioral Traits. Psychological Science, 2014, 25, 1975-1986.	1.8	92
30	Genetic Variation Associated with Differential Educational Attainment in Adults Has Anticipated Associations with School Performance in Children. PLoS ONE, 2014, 9, e100248.	1.1	31
31	GWAS of 126,559 Individuals Identifies Genetic Variants Associated with Educational Attainment. Science, 2013, 340, 1467-1471.	6.0	750
32	Aggregating Local Preferences to Guide Marginal Policy Adjustments. American Economic Review, 2013, 103, 605-610.	4.0	8
33	Molecular genetics and subjective well-being. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 9692-9697.	3.3	82
34	Why It Is Hard to Find Genes Associated With Social Science Traits: Theoretical and Empirical Considerations. American Journal of Public Health, 2013, 103, S152-S166.	1.5	52
35	The Molecular Genetic Architecture of Self-Employment. PLoS ONE, 2013, 8, e60542.	1.1	41
36	Most Reported Genetic Associations With General Intelligence Are Probably False Positives. Psychological Science, 2012, 23, 1314-1323.	1.8	221

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37	The genetic architecture of economic and political preferences. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 8026-8031.	3.3	225
38	The Promises and Pitfalls of Genoeconomics. Annual Review of Economics, 2012, 4, 627-662.	2.4	168