

Douglas F Levinson

List of Publications by Year in descending order

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Version: 2024-02-01

45
papers

11,687
citations

186265

28
h-index

243625

44
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48
docs citations

48
times ranked

18855
citing authors

#	ARTICLE	IF	CITATIONS
1	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. <i>Biological Psychiatry</i> , 2022, 91, 102-117.	1.3	61
2	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. <i>Biological Psychiatry</i> , 2022, 91, 313-327.	1.3	114
3	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. <i>Nature</i> , 2022, 604, 502-508.	27.8	929
4	Cross-platform validation of neurotransmitter release impairments in schizophrenia patient-derived <i>NRXN1</i> -mutant neurons. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	49
5	Narcolepsy and psychosis: A systematic review. <i>Acta Psychiatrica Scandinavica</i> , 2021, 144, 28-41.	4.5	17
6	The association between genetically determined ABO blood types and major depressive disorder. <i>Psychiatry Research</i> , 2021, 299, 113837.	3.3	4
7	Genome-wide analyses of smoking behaviors in schizophrenia: Findings from the Psychiatric Genomics Consortium. <i>Journal of Psychiatric Research</i> , 2021, 137, 215-224.	3.1	10
8	Transcriptome-based polygenic score links depression-related corticolimbic gene expression changes to sex-specific brain morphology and depression risk. <i>Neuropsychopharmacology</i> , 2021, 46, 2304-2311.	5.4	5
9	Increased activation product of complement 4 protein in plasma of individuals with schizophrenia. <i>Translational Psychiatry</i> , 2021, 11, 486.	4.8	10
10	Machine learning reveals bilateral distribution of somatic L1 insertions in human neurons and glia. <i>Nature Neuroscience</i> , 2021, 24, 186-196.	14.8	22
11	Polygenic burden could explain high rates of affective disorders in a community with restricted founder population. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2021, 186, 367-375.	1.7	0
12	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. <i>Biological Psychiatry</i> , 2020, 87, 419-430.	1.3	27
13	Developmental and symptom profiles in early-onset psychosis. <i>Schizophrenia Research</i> , 2020, 216, 470-478.	2.0	8
14	Cohort profile: the Australian genetics of depression study. <i>BMJ Open</i> , 2020, 10, e032580.	1.9	40
15	Minimal phenotyping yields genome-wide association signals of low specificity for major depression. <i>Nature Genetics</i> , 2020, 52, 437-447.	21.4	207
16	Quantifying between-cohort and between-sex genetic heterogeneity in major depressive disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019, 180, 439-447.	1.7	35
17	CWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. <i>American Journal of Psychiatry</i> , 2019, 176, 651-660.	7.2	186
18	Genome-wide Burden of Rare Short Deletions Is Enriched in Major Depressive Disorder in Four Cohorts. <i>Biological Psychiatry</i> , 2019, 85, 1065-1073.	1.3	25

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19	Genetic Correlation Profile of Schizophrenia Mirrors Epidemiological Results and Suggests Link Between Polygenic and Rare Variant (22q11.2) Cases of Schizophrenia. <i>Schizophrenia Bulletin</i> , 2018, 44, 1350-1361.	4.3	26
20	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. <i>Nature Genetics</i> , 2018, 50, 668-681.	21.4	2,224
21	An overview of medical risk factors for childhood psychosis: Implications for research and treatment. <i>Schizophrenia Research</i> , 2018, 192, 39-49.	2.0	67
22	Does Childhood Trauma Moderate Polygenic Risk for Depression? A Meta-analysis of 5765 Subjects From the Psychiatric Genomics Consortium. <i>Biological Psychiatry</i> , 2018, 84, 138-147.	1.3	87
23	Genome-wide association study of seasonal affective disorder. <i>Translational Psychiatry</i> , 2018, 8, 190.	4.8	18
24	1q21.1 microduplication: large verbalâ€“nonverbal performance discrepancy and ddPCR assays of HYDIN/HYDIN2 copy number. <i>Npj Genomic Medicine</i> , 2018, 3, 24.	3.8	7
25	Transdifferentiation of human adult peripheral blood T cells into neurons. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 6470-6475.	7.1	71
26	Genome-wide Association for Major Depression Through Age at Onset Stratification: Major Depressive Disorder Working Group of the Psychiatric Genomics Consortium. <i>Biological Psychiatry</i> , 2017, 81, 325-335.	1.3	175
27	Allele-specific expression reveals interactions between genetic variation and environment. <i>Nature Methods</i> , 2017, 14, 699-702.	19.0	135
28	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. <i>Biological Psychiatry</i> , 2017, 82, 322-329.	1.3	84
29	Polygenic Scores for Major Depressive Disorder and Risk of Alcohol Dependence. <i>JAMA Psychiatry</i> , 2017, 74, 1153.	11.0	73
30	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
31	Parental Origin of Interstitial Duplications at 15q11.2-q13.3 in Schizophrenia and Neurodevelopmental Disorders. <i>PLoS Genetics</i> , 2016, 12, e1005993.	3.5	51
32	Impact of the X Chromosome and sex on regulatory variation. <i>Genome Research</i> , 2016, 26, 768-777.	5.5	88
33	Genome-wide association study identifies 74 loci associated with educational attainment. <i>Nature</i> , 2016, 533, 539-542.	27.8	1,204
34	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
35	Winner's Curse Correction and Variable Thresholding Improve Performance of Polygenic Risk Modeling Based on Genome-Wide Association Study Summary-Level Data. <i>PLoS Genetics</i> , 2016, 12, e1006493.	3.5	98
36	Joint Analysis of Psychiatric Disorders Increases Accuracy of Risk Prediction for Schizophrenia, Bipolar Disorder, and Major Depressive Disorder. <i>American Journal of Human Genetics</i> , 2015, 96, 283-294.	6.2	225

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37	EIF3G is associated with narcolepsy across ethnicities. <i>European Journal of Human Genetics</i> , 2015, 23, 1573-1580.	2.8	21
38	A Rare Functional Noncoding Variant at the GWAS-Implicated MIR137/MIR2682 Locus Might Confer Risk to Schizophrenia and Bipolar Disorder. <i>American Journal of Human Genetics</i> , 2014, 95, 744-753.	6.2	91
39	Reciprocal Duplication of the Williams-Beuren Syndrome Deletion on Chromosome 7q11.23 Is Associated with Schizophrenia. <i>Biological Psychiatry</i> , 2014, 75, 371-377.	1.3	66
40	A mega-analysis of genome-wide association studies for major depressive disorder. <i>Molecular Psychiatry</i> , 2013, 18, 497-511.	7.9	1,002
41	Genome-Wide Association Study of Multiplex Schizophrenia Pedigrees. <i>American Journal of Psychiatry</i> , 2012, 169, 963-973.	7.2	61
42	Copy Number Variants in Schizophrenia: Confirmation of Five Previous Findings and New Evidence for 3q29 Microdeletions and VIPR2 Duplications. <i>American Journal of Psychiatry</i> , 2011, 168, 302-316.	7.2	398
43	Association analyses of 249,796 individuals reveal 18 new loci associated with body mass index. <i>Nature Genetics</i> , 2010, 42, 937-948.	21.4	2,634
44	Genetics of Recurrent Early-Onset Major Depression (GenRED): Significant Linkage on Chromosome 15q25-q26 After Fine Mapping With Single Nucleotide Polymorphism Markers. <i>American Journal of Psychiatry</i> , 2007, 164, 259-264.	7.2	48
45	Association Study of the Dystrobrevin-Binding Gene With Schizophrenia in Australian and Indian Samples. <i>Twin Research and Human Genetics</i> , 2006, 9, 531-539.	0.6	21