

Matthew B Mcqueen

List of Publications by Year in descending order

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Version: 2024-02-01

39
papers

8,254
citations

218677

26
h-index

302126

39
g-index

42
all docs

42
docs citations

42
times ranked

12763
citing authors

#	ARTICLE	IF	CITATIONS
1	Shared genetic risk between eating disorder and substance use-related phenotypes: Evidence from genome-wide association studies. <i>Addiction Biology</i> , 2021, 26, e12880.	2.6	28
2	Identification of 371 genetic variants for age at first sex and birth linked to externalising behaviour. <i>Nature Human Behaviour</i> , 2021, 5, 1717-1730.	12.0	62
3	A large-scale genome-wide association study meta-analysis of cannabis use disorder. <i>Lancet Psychiatry</i> , 2020, 7, 1032-1045.	7.4	200
4	Imputation of behavioral candidate gene repeat variants in 486,551 publicly-available UK Biobank individuals. <i>European Journal of Human Genetics</i> , 2019, 27, 963-969.	2.8	15
5	Association studies of up to 1.2 million individuals yield new insights into the genetic etiology of tobacco and alcohol use. <i>Nature Genetics</i> , 2019, 51, 237-244.	21.4	1,307
6	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
7	Chronic nicotinamide riboside supplementation is well-tolerated and elevates NAD+ in healthy middle-aged and older adults. <i>Nature Communications</i> , 2018, 9, 1286.	12.8	406
8	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. <i>Nature Neuroscience</i> , 2018, 21, 1656-1669.	14.8	490
9	Genetic influences on the human oral microbiome. <i>BMC Genomics</i> , 2017, 18, 659.	2.8	66
10	Oral trehalose supplementation improves resistance artery endothelial function in healthy middle-aged and older adults. <i>Aging</i> , 2016, 8, 1167-1183.	3.1	64
11	Is the Gene-Environment Interaction Paradigm Relevant to Genome-Wide Studies? The Case of Education and Body Mass Index. <i>Demography</i> , 2014, 51, 119-139.	2.5	54
12	Dietary Sodium Restriction Reverses Vascular Endothelial Dysfunction in Middle-Aged/Older Adults With Moderately Elevated Systolic Blood Pressure. <i>Journal of the American College of Cardiology</i> , 2013, 61, 335-343.	2.8	126
13	Comprehensive Research Synopsis and Systematic Meta-Analyses in Parkinson's Disease Genetics: The PDGene Database. <i>PLoS Genetics</i> , 2012, 8, e1002548.	3.5	495
14	Externalizing Behaviors are Associated with SNPs in the CHRNA5/CHRNA3/CHRNA4 Gene Cluster. <i>Behavior Genetics</i> , 2012, 42, 402-414.	2.1	28
15	Linkage disequilibrium mapping of the chromosome 6q21-22.31 bipolar I disorder susceptibility locus. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2010, 153B, 29-37.	1.7	9
16	Association of <i>CHRN</i> genes with dizziness to tobacco. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2010, 153B, 600-609.	1.7	37
17	Response to Dizziness Genes. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2010, 153B, n/a-n/a.	1.7	0
18	An omnibus test for family-based association studies with multiple SNPs and multiple phenotypes. <i>European Journal of Human Genetics</i> , 2010, 18, 720-725.	2.8	7

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19	Correspondence to Sand et al. "Critical Reappraisal of a Catechol-O-Methyltransferase Transversion Variant in Schizophrenia". <i>Biological Psychiatry</i> , 2010, 67, e45-e48.	1.3	2
20	Ethnicity, Body Mass, and Genome-Wide Data. <i>Biodemography and Social Biology</i> , 2010, 56, 123-136.	1.0	7
21	Genetic Association of the CHRNA6 and CHRN3 Genes with Tobacco Dependence in a Nationally Representative Sample. <i>Neuropsychopharmacology</i> , 2009, 34, 698-706.	5.4	90
22	Assessment of Alzheimer's disease case-control associations using family-based methods. <i>Neurogenetics</i> , 2009, 10, 19-25.	1.4	65
23	Response to Macgregor. <i>American Journal of Human Genetics</i> , 2008, 82, 799-800.	6.2	3
24	Systematic meta-analyses and field synopsis of genetic association studies in schizophrenia: the SzGene database. <i>Nature Genetics</i> , 2008, 40, 827-834.	21.4	961
25	The CHRNA5/A3/B4 Gene Cluster Variability as an Important Determinant of Early Alcohol and Tobacco Initiation in Young Adults. <i>Biological Psychiatry</i> , 2008, 63, 1039-1046.	1.3	174
26	Association of candidate genes with antisocial drug dependence in adolescents. <i>Drug and Alcohol Dependence</i> , 2008, 96, 90-98.	3.2	46
27	Evaluation of the Potential Excess of Statistically Significant Findings in Published Genetic Association Studies: Application to Alzheimer's Disease. <i>American Journal of Epidemiology</i> , 2008, 168, 855-865.	3.4	40
28	The neuronal nicotinic receptor subunit genes (CHRNA6 and CHRN3) are associated with subjective responses to tobacco. <i>Human Molecular Genetics</i> , 2007, 17, 724-734.	2.9	88
29	Genomewide Weighted Hypothesis Testing in Family-Based Association Studies, with an Application to a 100K Scan. <i>American Journal of Human Genetics</i> , 2007, 81, 607-614.	6.2	94
30	Exploring candidate gene associations with neuropsychological performance. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2007, 144B, 987-991.	1.7	15
31	On the parsing of statistical information in family-based association testing. <i>Nature Genetics</i> , 2007, 39, 281-282.	21.4	4
32	Systematic meta-analyses of Alzheimer disease genetic association studies: the AlzGene database. <i>Nature Genetics</i> , 2007, 39, 17-23.	21.4	1,626
33	A Common Genetic Variant Is Associated with Adult and Childhood Obesity. <i>Science</i> , 2006, 312, 279-283.	12.6	652
34	Variance Calculations for Identity-by-Descent Estimation. <i>American Journal of Human Genetics</i> , 2006, 78, 914-921.	6.2	3
35	Genomic screening and replication using the same data set in family-based association testing. <i>Nature Genetics</i> , 2005, 37, 683-691.	21.4	173
36	Genomic screening in family-based association testing. <i>BMC Genetics</i> , 2005, 6, S115.	2.7	3

#	ARTICLE	IF	CITATIONS
37	Comparison of linkage and association strategies for quantitative traits using the COGA dataset. BMC Genetics, 2005, 6, S96.	2.7	3
38	Combined Analysis from Eleven Linkage Studies of Bipolar Disorder Provides Strong Evidence of Susceptibility Loci on Chromosomes 6q and 8q. American Journal of Human Genetics, 2005, 77, 582-595.	6.2	218
39	A QTL genome scan of the metabolic syndrome and its component traits. BMC Genetics, 2003, 4, S96.	2.7	50