

Michael O'donovan

List of Publications by Year in descending order

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Version: 2024-02-01

40
papers

11,232
citations

159585

30
h-index

254184

43
g-index

44
all docs

44
docs citations

44
times ranked

17019
citing authors

#	ARTICLE	IF	CITATIONS
1	Facial Emotion Recognition in Psychosis and Associations With Polygenic Risk for Schizophrenia: Findings From the Multi-Center EU-GEI Case-Control Study. <i>Schizophrenia Bulletin</i> , 2022, 48, 1104-1114.	4.3	9
2	Jumping to conclusions, general intelligence, and psychosis liability: findings from the multi-centre EU-GEI case-control study. <i>Psychological Medicine</i> , 2021, 51, 623-633.	4.5	34
3	Genome-wide Association Analysis of Parkinson's Disease and Schizophrenia Reveals Shared Genetic Architecture and Identifies Novel Risk Loci. <i>Biological Psychiatry</i> , 2021, 89, 227-235.	1.3	53
4	Examining sex differences in neurodevelopmental and psychiatric genetic risk in anxiety and depression. <i>PLoS ONE</i> , 2021, 16, e0248254.	2.5	4
5	Investigating rare pathogenic/likely pathogenic exonic variation in bipolar disorder. <i>Molecular Psychiatry</i> , 2021, 26, 5239-5250.	7.9	15
6	Characterization of Single Gene Copy Number Variants in Schizophrenia. <i>Biological Psychiatry</i> , 2020, 87, 736-744.	1.3	10
7	A Population-Based Cohort Study Examining the Incidence and Impact of Psychotic Experiences From Childhood to Adulthood, and Prediction of Psychotic Disorder. <i>American Journal of Psychiatry</i> , 2020, 177, 308-317.	7.2	98
8	A brief report: de novo copy number variants in children with attention deficit hyperactivity disorder. <i>Translational Psychiatry</i> , 2020, 10, 135.	4.8	18
9	Estimating Exposome Score for Schizophrenia Using Predictive Modeling Approach in Two Independent Samples: The Results From the EUGEI Study. <i>Schizophrenia Bulletin</i> , 2019, 45, 960-965.	4.3	46
10	Association of copy number variation across the genome with neuropsychiatric traits in the general population. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018, 177, 489-502.	1.7	26
11	Genome-wide analysis of self-reported risk-taking behaviour and cross-disorder genetic correlations in the UK Biobank cohort. <i>Translational Psychiatry</i> , 2018, 8, 39.	4.8	57
12	Genetics of self-reported risk-taking behaviour, trans-ethnic consistency and relevance to brain gene expression. <i>Translational Psychiatry</i> , 2018, 8, 178.	4.8	29
13	Genome-wide analyses for personality traits identify six genomic loci and show correlations with psychiatric disorders. <i>Nature Genetics</i> , 2017, 49, 152-156.	21.4	350
14	Pleiotropic Effects of Trait-Associated Genetic Variation on DNA Methylation: Utility for Refining GWAS Loci. <i>American Journal of Human Genetics</i> , 2017, 100, 954-959.	6.2	77
15	Genome-wide analysis in UK Biobank identifies four loci associated with mood instability and genetic correlation with major depressive disorder, anxiety disorder and schizophrenia. <i>Translational Psychiatry</i> , 2017, 7, 1264.	4.8	69
16	Leveraging Genomic Annotations and Pleiotropic Enrichment for Improved Replication Rates in Schizophrenia GWAS. <i>PLoS Genetics</i> , 2016, 12, e1005803.	3.5	34
17	Shared Genetic Influences Between Attention-Deficit/Hyperactivity Disorder (ADHD) Traits in Children and Clinical ADHD. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2015, 54, 322-327.	0.5	75
18	Novel genetic advances in schizophrenia: an interview with Michael O'Donovan. <i>BMC Medicine</i> , 2015, 13, 181.	5.5	1

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19	An inherited duplication at the gene p21 Protein-Activated Kinase 7 (PAK7) is a risk factor for psychosis. <i>Human Molecular Genetics</i> , 2014, 23, 3316-3326.	2.9	37
20	Cognitive mechanisms underlying reading and spelling development in five European orthographies. <i>Learning and Instruction</i> , 2014, 29, 65-77.	3.2	293
21	Predictors of developmental dyslexia in European orthographies with varying complexity. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2013, 54, 686-694.	5.2	307
22	A mega-analysis of genome-wide association studies for major depressive disorder. <i>Molecular Psychiatry</i> , 2013, 18, 497-511.	7.9	1,002
23	Investigating the Contribution of Common Genetic Variants to the Risk and Pathogenesis of ADHD. <i>American Journal of Psychiatry</i> , 2012, 169, 186-194.	7.2	174
24	Genetic architectures of psychiatric disorders: the emerging picture and its implications. <i>Nature Reviews Genetics</i> , 2012, 13, 537-551.	16.3	1,025
25	Common variants at ABCA7, MS4A6A/MS4A4E, EPHA1, CD33 and CD2AP are associated with Alzheimer's disease. <i>Nature Genetics</i> , 2011, 43, 429-435.	21.4	1,708
26	No evidence that extended tracts of homozygosity are associated with Alzheimer's disease. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2011, 156, 764-771.	1.7	17
27	Clinical and cognitive characteristics of children with attention-deficit hyperactivity disorder, with and without copy number variants. <i>British Journal of Psychiatry</i> , 2011, 199, 398-403.	2.8	28
28	Meta-Analysis of Genome-Wide Association Studies of Attention-Deficit/Hyperactivity Disorder. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2010, 49, 884-897.	0.5	423
29	Rare chromosomal deletions and duplications in attention-deficit hyperactivity disorder: a genome-wide analysis. <i>Lancet, The</i> , 2010, 376, 1401-1408.	13.7	485
30	Genome-wide association study identifies variants at CLU and PICALM associated with Alzheimer's disease. <i>Nature Genetics</i> , 2009, 41, 1088-1093.	21.4	2,697
31	Genome-Wide Association Identifies a Common Variant in the Reelin Gene That Increases the Risk of Schizophrenia Only in Women. <i>PLoS Genetics</i> , 2008, 4, e28.	3.5	302
32	Evidence for novel susceptibility genes for late-onset Alzheimer's disease from a genome-wide association study of putative functional variants. <i>Human Molecular Genetics</i> , 2007, 16, 865-873.	2.9	256
33	A Scan of Chromosome 10 Identifies a Novel Locus Showing Strong Association with Late-Onset Alzheimer Disease. <i>American Journal of Human Genetics</i> , 2006, 78, 78-88.	6.2	157
34	DAPK1 variants are associated with Alzheimer's disease and allele-specific expression. <i>Human Molecular Genetics</i> , 2006, 15, 2560-2568.	2.9	125
35	Genetic association of the APP binding protein 2 gene (APBB2) with late onset Alzheimer disease. <i>Human Mutation</i> , 2005, 25, 270-277.	2.5	36
36	The genetics of attention deficit hyperactivity disorder. <i>Human Molecular Genetics</i> , 2005, 14, R275-R282.	2.9	189

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37	Association of the Dopamine D4Receptor Gene 7-Repeat Allele With Neuropsychological Test Performance of Children With ADHD. <i>American Journal of Psychiatry</i> , 2004, 161, 133-138.	7.2	162
38	DNA pooling as a tool for large-scale association studies in complex traits. <i>Annals of Medicine</i> , 2004, 36, 146-152.	3.8	68
39	DNA Pooling: a tool for large-scale association studies. <i>Nature Reviews Genetics</i> , 2002, 3, 862-871.	16.3	534
40	Cheap, accurate and rapid allele frequency estimation of single nucleotide polymorphisms by primer extension and DHPLC in DNA pools. <i>Human Genetics</i> , 2000, 107, 488-493.	3.8	162