Oliver Pain

List of Publications by Year
in descending order

23	691	623734 14	677142 22
papers	citations	h-index	g-index

37	37	37	1592
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Evaluation of polygenic prediction methodology within a reference-standardized framework. PLoS Genetics, 2021, 17, e1009021.	3.5	99
2	Genomeâ€wide analysis of adolescent psychoticâ€like experiences shows genetic overlap with psychiatric disorders. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2018, 177, 416-425.	1.7	74
3	Novel Insight Into the Etiology of Autism Spectrum Disorder Gained by Integrating Expression Data With Genome-wide Association Statistics. Biological Psychiatry, 2019, 86, 265-273.	1.3	65
4	Delineating the Genetic Component of Gene Expression in Major Depression. Biological Psychiatry, 2021, 89, 627-636.	1.3	63
5	A transcriptome-wide association study implicates specific pre- and post-synaptic abnormalities in schizophrenia. Human Molecular Genetics, 2020, 29, 159-167.	2.9	54
6	Impute.me: An Open-Source, Non-profit Tool for Using Data From Direct-to-Consumer Genetic Testing to Calculate and Interpret Polygenic Risk Scores. Frontiers in Genetics, 2020, 11, 578.	2.3	47
7	Identifying the Common Genetic Basis of Antidepressant Response. Biological Psychiatry Global Open Science, 2022, 2, 115-126.	2.2	31
8	Genes and Gene Networks Implicated in Aggression Related Behaviour. Neurogenetics, 2014, 15, 255-266.	1.4	30
9	A systematic review of genome-wide research on psychotic experiences and negative symptom traits: new revelations and implications for psychiatry. Human Molecular Genetics, 2018, 27, R136-R152.	2.9	27
10	Multiple measures of depression to enhance validity of major depressive disorder in the UK Biobank. BJPsych Open, 2021, 7, e44.	0.7	27
11	Transcriptome-wide association study of treatment-resistant depression and depression subtypes for drug repurposing. Neuropsychopharmacology, 2021, 46, 1821-1829.	5.4	27
12	Cis-effects on gene expression in the human prenatal brain associated with genetic risk for neuropsychiatric disorders. Molecular Psychiatry, 2021, 26, 2082-2088.	7.9	23
13	Are your covariates under control? How normalization can re-introduce covariate effects. European Journal of Human Genetics, 2018, 26, 1194-1201.	2.8	21
14	A tool for translating polygenic scores onto the absolute scale using summary statistics. European Journal of Human Genetics, 2022, 30, 339-348.	2.8	18
15	Genetic overlap between psychotic experiences in the community across age and with psychiatric disorders. Translational Psychiatry, 2020, 10, 86.	4.8	15
16	Imputed gene expression risk scores: a functionally informed component of polygenic risk. Human Molecular Genetics, 2021, 30, 727-738.	2.9	11
17	Evidence for specificity of polygenic contributions to attainment in English, maths and science during adolescence. Scientific Reports, 2021, 11, 3851.	3.3	10
18	Transcriptome-wide association study reveals two genes that influence mismatch negativity. Cell Reports, 2021, 34, 108868.	6.4	8

#	Article	IF	Citations
19	Methylome-wide association study of early life stressors and adult mental health. Human Molecular Genetics, 2022, 31, 651-664.	2.9	7
20	Latent subtypes of manic and/or irritable episode symptoms in two population-based cohorts. British Journal of Psychiatry, 2022, 221, 722-731.	2.8	4
21	Exploring polygenicâ€environment and residualâ€environment interactions for depressive symptoms within the UK Biobank. Genetic Epidemiology, 2022, 46, 219-233.	1.3	4
22	Investigating an in silico approach for prioritizing antidepressant drug prescription based on drug-induced expression profiles and predicted gene expression. Pharmacogenomics Journal, 2021, 21, 85-93.	2.0	1
23	SA139GENETIC ASSOCIATION BETWEEN TOBACCO USE AND SPECIFIC PSYCHOTIC EXPERIENCES DURING ADOLESCENCE. European Neuropsychopharmacology, 2019, 29, S1265.	0.7	0