## Oliver Pain

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/325505/publications.pdf

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623734 677142 23 691 14 22 citations h-index g-index papers 37 37 37 1592 citing authors docs citations times ranked all docs

#	Article	IF	CITATIONS
1	Identifying the Common Genetic Basis of Antidepressant Response. Biological Psychiatry Global Open Science, 2022, 2, 115-126.	2.2	31
2	Methylome-wide association study of early life stressors and adult mental health. Human Molecular Genetics, 2022, 31, 651-664.	2.9	7
3	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
4	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
5	Exploring polygenicâ€environment and residualâ€environment interactions for depressive symptoms within the UK Biobank. Genetic Epidemiology, 2022, 46, 219-233.	1.3	4
6	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
7	Delineating the Genetic Component of Gene Expression in Major Depression. Biological Psychiatry, 2021, 89, 627-636.	1.3	63
8	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
9	Evidence for specificity of polygenic contributions to attainment in English, maths and science during adolescence. Scientific Reports, 2021, 11, 3851.	3.3	10
10	Imputed gene expression risk scores: a functionally informed component of polygenic risk. Human Molecular Genetics, 2021, 30, 727-738.	2.9	11
11	Multiple measures of depression to enhance validity of major depressive disorder in the UK Biobank. BJPsych Open, 2021, 7, e44.	0.7	27
12	Transcriptome-wide association study reveals two genes that influence mismatch negativity. Cell Reports, 2021, 34, 108868.	6.4	8
13	Evaluation of polygenic prediction methodology within a reference-standardized framework. PLoS Genetics, 2021, 17, e1009021.	3.5	99
14	Transcriptome-wide association study of treatment-resistant depression and depression subtypes for drug repurposing. Neuropsychopharmacology, 2021, 46, 1821-1829.	5.4	27
15	A transcriptome-wide association study implicates specific pre- and post-synaptic abnormalities in schizophrenia. Human Molecular Genetics, 2020, 29, 159-167.	2.9	54
16	Genetic overlap between psychotic experiences in the community across age and with psychiatric disorders. Translational Psychiatry, 2020, 10, 86.	4.8	15
17	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
18	SA139GENETIC ASSOCIATION BETWEEN TOBACCO USE AND SPECIFIC PSYCHOTIC EXPERIENCES DURING ADOLESCENCE. European Neuropsychopharmacology, 2019, 29, S1265.	0.7	0

#	Article	IF	CITATIONS
19	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
20	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
21	Are your covariates under control? How normalization can re-introduce covariate effects. European Journal of Human Genetics, 2018, 26, 1194-1201.	2.8	21
22	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
23	Genes and Gene Networks Implicated in Aggression Related Behaviour. Neurogenetics, 2014, 15, 255-266.	1.4	30