Shahram Bahrami

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/6309609/publications.pdf

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27	892	566801 15	552369 26
papers	citations	h-index	g-index

32	32	32	1423
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Genome-wide analysis reveals extensive genetic overlap between schizophrenia, bipolar disorder, and intelligence. Molecular Psychiatry, 2020, 25, 844-853.	4.1	156
2	Discovery of shared genomic loci using the conditional false discovery rate approach. Human Genetics, 2020, 139, 85-94.	1.8	109
3	Shared Genetic Loci Between Body Mass Index and Major Psychiatric Disorders. JAMA Psychiatry, 2020, 77, 503.	6.0	82
4	Genome-wide Association Analysis of Parkinsonâ€™s Disease and Schizophrenia Reveals Shared Genetic Architecture and Identifies Novel Risk Loci. Biological Psychiatry, 2021, 89, 227-235.	0.7	53
5	Meta-analysis of Alzheimerâ€™s disease on 9,751 samples from Norway and IGAP study identifies four risk loci. Scientific Reports, 2018, 8, 18088.	1.6	47
6	Genetic Association Between Schizophrenia and Cortical Brain Surface Area and Thickness. JAMA Psychiatry, 2021, 78, 1020.	6.0	43
7	Identification of genetic overlap and novel risk loci for attention-deficit/hyperactivity disorder and bipolar disorder. Molecular Psychiatry, 2021, 26, 4055-4065.	4.1	31
8	Polygenic overlap and shared genetic loci between loneliness, severe mental disorders, and cardiovascular disease risk factors suggest shared molecular mechanisms. Translational Psychiatry, 2021, 11, 3.	2.4	29
9	Dissecting the shared genetic basis of migraine and mental disorders using novel statistical tools. Brain, 2022, 145, 142-153.	3.7	27
10	The genetic architecture of human brainstem structures and their involvement in common brain disorders. Nature Communications, 2020, 11, 4016.	5.8	26
11	The genetic architecture of the human thalamus and its overlap with ten common brain disorders. Nature Communications, 2021, 12, 2909.	5.8	25
12	Exploring lithiumâ€™s transcriptional mechanisms of action in bipolar disorder: a multi-step study. Neuropsychopharmacology, 2020, 45, 947-955.	2.8	24
13	Characterizing the Genetic Overlap Between Psychiatric Disorders and Sleep-Related Phenotypes. Biological Psychiatry, 2021, 90, 621-631.	0.7	24
14	Genetic loci shared between major depression and intelligence with mixed directions of effect. Nature Human Behaviour, 2021, 5, 795-801.	6.2	23
15	Oxytocin receptor expression patterns in the human brain across development. Neuropsychopharmacology, 2022, 47, 1550-1560.	2.8	23
16	Shared genetic loci between depression and cardiometabolic traits. PLoS Genetics, 2022, 18, e1010161.	1.5	18
17	Shared Genetics of Multiple System Atrophy and Inflammatory Bowel Disease. Movement Disorders, 2021, 36, 449-459.	2.2	16
18	Extensive bidirectional genetic overlap between bipolar disorder and cardiovascular disease phenotypes. Translational Psychiatry, 2021, 11, 407.	2.4	16

#	Article	IF	Citations
19	Genomeâ€wide analysis reveals genetic overlap between alcohol use behaviours, schizophrenia and bipolar disorder and identifies novel shared risk loci. Addiction, 2022, 117, 600-610.	1.7	16
20	Characterising the shared genetic determinants of bipolar disorder, schizophrenia and risk-taking. Translational Psychiatry, 2021, 11, 466.	2.4	15
21	Identification of Genetic Loci Shared Between Attention-Deficit/Hyperactivity Disorder, Intelligence, and Educational Attainment. Biological Psychiatry, 2020, 87, 1052-1062.	0.7	13
22	Distributed genetic architecture across the hippocampal formation implies common neuropathology across brain disorders. Nature Communications, 2022, 13, .	5.8	12
23	Shared genetic architecture between neuroticism, coronary artery disease and cardiovascular risk factors. Translational Psychiatry, 2021, 11, 368.	2.4	10
24	The shared genetic basis of mood instability and psychiatric disorders: A crossâ€trait genomeâ€wide association analysis. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2022, 189, 207-218.	1.1	10
25	Phenotype-specific differences in polygenicity and effect size distribution across functional annotation categories revealed by AI-MiXeR. Bioinformatics, 2020, 36, 4749-4756.	1.8	6
26	Genetic variants associated with cardiometabolic abnormalities during treatment with selective serotonin reuptake inhibitors: a genome-wide association study. Pharmacogenomics Journal, 2021, 21, 574-585.	0.9	5
27	Using Polygenic Hazard Scores to Predict Age at Onset of Alzheimerâ€™s Disease in Nordic Populations. Journal of Alzheimer's Disease, 2022, 88, 1533-1544.	1.2	3