

Hongbao Cao

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/761801/publications.pdf>

Version: 2024-02-01

24
papers

535
citations

933447

10
h-index

713466

21
g-index

26
all docs

26
docs citations

26
times ranked

466
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic evidence suggests posttraumatic stress disorder as a subtype of major depressive disorder. <i>Journal of Clinical Investigation</i> , 2022, 132, .	8.2	68
2	Classifying major mental disorders genetically. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2022, 112, 110410.	4.8	11
3	Shared genetic liability and causal effects between major depressive disorder and insomnia. <i>Human Molecular Genetics</i> , 2022, 31, 1336-1345.	2.9	22
4	Involvement of the long intergenic non-coding RNA LINC00461 in schizophrenia. <i>BMC Psychiatry</i> , 2022, 22, 59.	2.6	9
5	Causal Association and Shared Genetics Between Asthma and COVID-19. <i>Frontiers in Immunology</i> , 2022, 13, 705379.	4.8	16
6	Convergent lines of evidence supporting involvement of NFKB1 in schizophrenia. <i>Psychiatry Research</i> , 2022, 312, 114588.	3.3	8
7	Shared genetics between autism spectrum disorder and attention-deficit/hyperactivity disorder and their association with extraversion. <i>Psychiatry Research</i> , 2022, 314, 114679.	3.3	22
8	Causal links between major depressive disorder and insomnia: A Mendelian randomisation study. <i>Gene</i> , 2021, 768, 145271.	2.2	25
9	In silico Gene Set and Pathway Enrichment Analyses Highlight Involvement of Ion Transport in Cholinergic Pathways in Autism: Rationale for Nutritional Intervention. <i>Frontiers in Neuroscience</i> , 2021, 15, 648410.	2.8	4
10	Causal influences of neuroticism on mental health and cardiovascular disease. <i>Human Genetics</i> , 2021, 140, 1267-1281.	3.8	71
11	Schizophrenia Plays a Negative Role in the Pathological Development of Myocardial Infarction at Multiple Biological Levels. <i>Frontiers in Genetics</i> , 2021, 12, 607690.	2.3	2
12	Genetic mechanisms of COVID-19 and its association with smoking and alcohol consumption. <i>Briefings in Bioinformatics</i> , 2021, 22, .	6.5	31
13	A microfluidic array device for single cell capture and intracellular Ca ²⁺ response analysis induced by dynamic biochemical stimulus. <i>Bioscience Reports</i> , 2021, 41, .	2.4	3
14	Unraveling Risk Genes of COVID-19 by Multi-Omics Integrative Analyses. <i>Frontiers in Medicine</i> , 2021, 8, 738687.	2.6	22
15	Shared Genetic Liability and Causal Associations Between Major Depressive Disorder and Cardiovascular Diseases. <i>Frontiers in Cardiovascular Medicine</i> , 2021, 8, 735136.	2.4	25
16	Identifying common genome-wide risk genes for major psychiatric traits. <i>Human Genetics</i> , 2020, 139, 185-198.	3.8	40
17	Preeclampsia Drives Molecular Networks to Shift Toward Greater Vulnerability to the Development of Autism Spectrum Disorder. <i>Frontiers in Neurology</i> , 2020, 11, 590.	2.4	6
18	Development of two psychological experience questionnaires for screening violence-related mental health disorders of non-psychiatric inpatients. <i>Health and Quality of Life Outcomes</i> , 2020, 18, 151.	2.4	1

#	ARTICLE	IF	CITATIONS
19	TNFRSF12A and CD38 Contribute to a Vicious Circle for Chronic Obstructive Pulmonary Disease by Engaging Senescence Pathways. <i>Frontiers in Cell and Developmental Biology</i> , 2020, 8, 330.	3.7	2
20	Multi-trait analysis for genome-wide association study of five psychiatric disorders. <i>Translational Psychiatry</i> , 2020, 10, 209.	4.8	132
21	Improve cognition of depressive patients through the regulation of basal ganglia connectivity: Combined medication using Shuganjieyu capsule. <i>Journal of Psychiatric Research</i> , 2020, 123, 39-47.	3.1	7
22	<i>GPNMB</i> contributes to a vicious circle for chronic obstructive pulmonary disease. <i>Bioscience Reports</i> , 2020, 40, .	2.4	1
23	Variants and expression changes in PPAR-encoding genes display no significant association with schizophrenia. <i>Bioscience Reports</i> , 2020, 40, .	2.4	3
24	A core collection of pan-schizophrenia genes allows building cohort-specific signatures of affected brain. <i>Scientific Reports</i> , 2019, 9, 12671.	3.3	2