

Su-Hua Chang

List of Publications by Year in descending order

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Version: 2024-02-01

37
papers

1,201
citations

623734

14
h-index

395702

33
g-index

39
all docs

39
docs citations

39
times ranked

2603
citing authors

#	ARTICLE	IF	CITATIONS
1	The neural correlates of reward-related processing in major depressive disorder: A meta-analysis of functional magnetic resonance imaging studies. <i>Journal of Affective Disorders</i> , 2013, 151, 531-539.	4.1	292
2	Peripheral cytokine levels and response to antidepressant treatment in depression: a systematic review and meta-analysis. <i>Molecular Psychiatry</i> , 2020, 25, 339-350.	7.9	228
3	Molecular genetic studies of ADHD and its candidate genes: A review. <i>Psychiatry Research</i> , 2014, 219, 10-24.	3.3	118
4	Mutations in apoptosis-inducing factor cause X-linked recessive auditory neuropathy spectrum disorder. <i>Journal of Medical Genetics</i> , 2015, 52, 523-531.	3.2	92
5	BDgene: A Genetic Database for Bipolar Disorder and Its Overlap With Schizophrenia and Major Depressive Disorder. <i>Biological Psychiatry</i> , 2013, 74, 727-733.	1.3	47
6	Network-Based Analysis of Schizophrenia Genome-Wide Association Data to Detect the Joint Functional Association Signals. <i>PLoS ONE</i> , 2015, 10, e0133404.	2.5	38
7	Association of Interleukin-10 Polymorphisms with Schizophrenia: A Meta-Analysis. <i>PLoS ONE</i> , 2014, 9, e90407.	2.5	37
8	Neural substrates of smoking and reward cue reactivity in smokers: a meta-analysis of fMRI studies. <i>Translational Psychiatry</i> , 2020, 10, 97.	4.8	37
9	Shared polygenic risk for ADHD, executive dysfunction and other psychiatric disorders. <i>Translational Psychiatry</i> , 2020, 10, 182.	4.8	32
10	Integrating genome-wide association study and expression quantitative trait loci data identifies NEGR1 as a causal risk gene of major depression disorder. <i>Journal of Affective Disorders</i> , 2020, 265, 679-686.	4.1	27
11	Genome-wide association study of alcohol dependence in male Han Chinese and cross-ethnic polygenic risk score comparison. <i>Translational Psychiatry</i> , 2019, 9, 249.	4.8	21
12	Identification of novel risk loci with shared effects on alcoholism, heroin, and methamphetamine dependence. <i>Molecular Psychiatry</i> , 2021, 26, 1152-1161.	7.9	21
13	An overview of posttraumatic stress disorder genetic studies by analyzing and integrating genetic data into genetic database PTSDgene. <i>Neuroscience and Biobehavioral Reviews</i> , 2017, 83, 647-656.	6.1	17
14	Common variants near IKZF1 are associated with primary Sjögren's syndrome in Han Chinese. <i>PLoS ONE</i> , 2017, 12, e0177320.	2.5	17
15	A robust and powerful two-step testing procedure for local ancestry adjusted allelic association analysis in admixed populations. <i>Genetic Epidemiology</i> , 2018, 42, 288-302.	1.3	17
16	Precision Medicine: What Challenges Are We Facing?. <i>Genomics, Proteomics and Bioinformatics</i> , 2016, 14, 253-261.	6.9	15
17	Polygenic evidence and overlapped brain functional connectivities for the association between chronic pain and sleep disturbance. <i>Translational Psychiatry</i> , 2020, 10, 252.	4.8	15
18	GABRA2 rs279858-linked variants are associated with disrupted structural connectome of reward circuits in heroin abusers. <i>Translational Psychiatry</i> , 2018, 8, 138.	4.8	14

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19	The Association with Quantitative Response to Attention-Deficit/Hyperactivity Disorder Medication of the Previously Identified Neurodevelopmental Network Genes. <i>Journal of Child and Adolescent Psychopharmacology</i> , 2020, 30, 348-354.	1.3	12
20	Prioritization of candidate genes for attention deficit hyperactivity disorder by computational analysis of multiple data sources. <i>Protein and Cell</i> , 2012, 3, 526-534.	11.0	11
21	The effect of perceived stress on cognition is mediated by personality and the underlying neural mechanism. <i>Translational Psychiatry</i> , 2022, 12, 199.	4.8	11
22	Genome-wide meta-analysis of alcohol use disorder in East Asians. <i>Neuropsychopharmacology</i> , 2022, 47, 1791-1797.	5.4	10
23	Genetic variant for behavioral regulation factor of executive function and its possible brain mechanism in attention deficit hyperactivity disorder. <i>Scientific Reports</i> , 2018, 8, 7620.	3.3	9
24	Risk factors and an early prediction model for persistent methamphetamine-related psychiatric symptoms. <i>Addiction Biology</i> , 2020, 25, e12709.	2.6	8
25	Career choice and influential factors among medical students majoring in psychiatry in China. <i>BMC Medical Education</i> , 2021, 21, 183.	2.4	8
26	Synthetic biology: From the first synthetic cell to see its current situation and future development. <i>Science Bulletin</i> , 2011, 56, 229-237.	1.7	7
27	Genetic analysis for cognitive flexibility in the trail-making test in attention deficit hyperactivity disorder patients from single nucleotide polymorphism, gene to pathway level. <i>World Journal of Biological Psychiatry</i> , 2019, 20, 476-485.	2.6	7
28	A combined analysis of genome-wide expression profiling of bipolar disorder in human prefrontal cortex. <i>Journal of Psychiatric Research</i> , 2016, 82, 23-29.	3.1	6
29	Associations of DNA Methylation With Behavioral Problems, Gray Matter Volumes, and Negative Life Events Across Adolescence: Evidence From the Longitudinal IMAGEN Study. <i>Biological Psychiatry</i> , 2023, 93, 342-351.	1.3	6
30	Pathway-based analysis for genome-wide association study data of bipolar disorder provides new insights for genetic study. <i>Protein and Cell</i> , 2015, 6, 912-915.	11.0	5
31	Identification of Functional CircRNA-miRNA-mRNA Regulatory Network in Dorsolateral Prefrontal Cortex Neurons of Patients With Cocaine Use Disorder. <i>Frontiers in Molecular Neuroscience</i> , 2022, 15, 839233.	2.9	5
32	Problem-Based Learning Could Tackle the Issue of Insufficient Education and Adherence in People Living With HIV/AIDS. <i>Frontiers in Pharmacology</i> , 2019, 10, 901.	3.5	4
33	Common and distinguishing genetic factors for substance use behavior and disorder: an integrated analysis of genomic and transcriptomic studies from both human and animal studies. <i>Addiction</i> , 2022, 117, 2515-2529.	3.3	3
34	Connectivity-Based Brain Network Supports Restricted and Repetitive Behaviors in Autism Spectrum Disorder Across Development. <i>Frontiers in Psychiatry</i> , 2022, 13, 874090.	2.6	2
35	The Impact of Quarantine on Pain Sensation among the General Population in China during the COVID-19 Pandemic. <i>Brain Sciences</i> , 2022, 12, 79.	2.3	1
36	Genome-wide association meta-analyses identify novel genetic risk loci and polygenic phenotype associations for heroin, methamphetamine and alcohol dependences. <i>Clinical and Translational Medicine</i> , 2022, 12, e659.	4.0	1

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37	DNA Methylation Markers and Prediction Model for Depression and Their Contribution for Breast Cancer Risk. <i>Frontiers in Molecular Neuroscience</i> , 2022, 15, 845212.	2.9	0