

Huiping Zhang

List of Publications by Year in descending order

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65
papers

2,716
citations

236925

25
h-index

189892

50
g-index

67
all docs

67
docs citations

67
times ranked

4698
citing authors

#	ARTICLE	IF	CITATIONS
1	Child Abuse, Depression, and Methylation in Genes Involved With Stress, Neural Plasticity, and Brain Circuitry. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2014, 53, 417-424.e5.	0.5	273
2	Prevalence of metabolic syndrome in mainland china: a meta-analysis of published studies. <i>BMC Public Health</i> , 2016, 16, 296.	2.9	240
3	Child Abuse and Epigenetic Mechanisms of Disease Risk. <i>American Journal of Preventive Medicine</i> , 2013, 44, 101-107.	3.0	212
4	Opioid Receptor Gene (OPRM1, OPRK1, and OPRD1) Variants and Response to Naltrexone Treatment for Alcohol Dependence: Results From the VA Cooperative Study. <i>Alcoholism: Clinical and Experimental Research</i> , 2007, 31, 070212174136005-???	2.4	178
5	Brain derived neurotrophic factor (<i>BDNF</i>) gene variants and Alzheimer's disease, affective disorders, posttraumatic stress disorder, schizophrenia, and substance dependence. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2006, 141B, 387-393.	1.7	170
6	Association between two μ -opioid receptor gene (OPRM1) haplotype blocks and drug or alcohol dependence. <i>Human Molecular Genetics</i> , 2006, 15, 807-819.	2.9	155
7	Sex-biased methylome and transcriptome in human prefrontal cortex. <i>Human Molecular Genetics</i> , 2014, 23, 1260-1270.	2.9	130
8	Array-Based Profiling of <sc>DNA</sc> Methylation Changes Associated with Alcohol Dependence. <i>Alcoholism: Clinical and Experimental Research</i> , 2013, 37, E108-15.	2.4	83
9	A large-scale meta-analysis of the association between the ANKK1/DRD2 Taq1A polymorphism and alcohol dependence. <i>Human Genetics</i> , 2013, 132, 347-358.	3.8	74
10	Change in FK506 binding protein 5 (<i>FKBP5</i>) methylation over time among preschoolers with adversity. <i>Development and Psychopathology</i> , 2017, 29, 1627-1634.	2.3	69
11	DNA co-methylation modules in postmortem prefrontal cortex tissues of European Australians with alcohol use disorders. <i>Scientific Reports</i> , 2016, 6, 19430.	3.3	68
12	Childhood Adversity Increases Risk for Nicotine Dependence and Interacts with $\alpha 5$ Nicotinic Acetylcholine Receptor Genotype Specifically in Males. <i>Neuropsychopharmacology</i> , 2012, 37, 669-676.	5.4	63
13	Hypermethylation of OPRM1 promoter region in European Americans with alcohol dependence. <i>Journal of Human Genetics</i> , 2012, 57, 670-675.	2.3	57
14	Population-specific effects of the Asn40Asp polymorphism at the μ -opioid receptor gene (OPRM1) on HPA-axis activation. <i>Pharmacogenetics and Genomics</i> , 2007, 17, 1031-1038.	1.5	56
15	Review: DNA methylation and alcohol use disorders: Progress and challenges. <i>American Journal on Addictions</i> , 2017, 26, 502-515.	1.4	49
16	Profiling of Childhood Adversity-Associated DNA Methylation Changes in Alcoholic Patients and Healthy Controls. <i>PLoS ONE</i> , 2013, 8, e65648.	2.5	41
17	Identification of methylation quantitative trait loci (mQTLs) influencing promoter DNA methylation of alcohol dependence risk genes. <i>Human Genetics</i> , 2014, 133, 1093-1104.	3.8	39
18	Differentially co-expressed genes in postmortem prefrontal cortex of individuals with alcohol use disorders: influence on alcohol metabolism-related pathways. <i>Human Genetics</i> , 2014, 133, 1383-1394.	3.8	36

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19	Epigenetic and pharmacological regulation of 5HT3 receptors controls compulsive ethanol seeking in mice. <i>European Journal of Neuroscience</i> , 2014, 39, 999-1008.	2.6	33
20	Ethanol Upregulates NMDA Receptor Subunit Gene Expression in Human Embryonic Stem Cell-Derived Cortical Neurons. <i>PLoS ONE</i> , 2015, 10, e0134907.	2.5	33
21	Pro-Opiomelanocortin Gene Variation Related to Alcohol or Drug Dependence: Evidence and Replications Across Family- and Population-based Studies. <i>Biological Psychiatry</i> , 2009, 66, 128-136.	1.3	31
22	Time Series Analyses of Hand, Foot and Mouth Disease Integrating Weather Variables. <i>PLoS ONE</i> , 2015, 10, e0117296.	2.5	31
23	Genomewide Study of Epigenetic Biomarkers of Opioid Dependence in European- American Women. <i>Scientific Reports</i> , 2019, 9, 4660.	3.3	30
24	Cognitive Flexibility is Associated with KIBRA Variant and Modulated by Recent Tobacco Use. <i>Neuropsychopharmacology</i> , 2009, 34, 2508-2516.	5.4	28
25	Functional impact of a single-nucleotide polymorphism in the OPRD1 promoter region. <i>Journal of Human Genetics</i> , 2010, 55, 278-284.	2.3	27
26	Variation in the Nicotinic Acetylcholine Receptor Gene Cluster CHRNA5-CHRNA3-CHRNA4 and Its Interaction with Recent Tobacco Use Influence Cognitive Flexibility. <i>Neuropsychopharmacology</i> , 2010, 35, 2211-2224.	5.4	26
27	Ethanol-induced Htr3a Promoter Methylation Changes in Mouse Blood and Brain. <i>Alcoholism: Clinical and Experimental Research</i> , 2013, 37, E101-7.	2.4	26
28	Smoking Prevalence and Associated Factors as well as Attitudes and Perceptions towards Tobacco Control in Northeast China. <i>International Journal of Environmental Research and Public Health</i> , 2015, 12, 8606-8618.	2.6	26
29	Multiple OPR genes influence personality traits in substance dependent and healthy subjects in two American populations. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008, 147B, 1028-1039.	1.7	25
30	CHRM2 variation predisposes to personality traits of agreeableness and conscientiousness. <i>Human Molecular Genetics</i> , 2007, 16, 1557-1568.	2.9	23
31	Exploration of alcohol use disorder-associated brain miRNA-mRNA regulatory networks. <i>Translational Psychiatry</i> , 2021, 11, 504.	4.8	23
32	Identification of Risk Factors Affecting Impaired Fasting Glucose and Diabetes in Adult Patients from Northeast China. <i>International Journal of Environmental Research and Public Health</i> , 2015, 12, 12662-12678.	2.6	21
33	Loss-of-function Mutation in PMVK Causes Autosomal Dominant Disseminated Superficial Porokeratosis. <i>Scientific Reports</i> , 2016, 6, 24226.	3.3	21
34	Genome-wide association study of body mass index in subjects with alcohol dependence. <i>Addiction Biology</i> , 2017, 22, 535-549.	2.6	21
35	Salivary microRNAs identified by small RNA sequencing and machine learning as potential biomarkers of alcohol dependence. <i>Epigenomics</i> , 2019, 11, 739-749.	2.1	19
36	ADH7 variation modulates extraversion and conscientiousness in substance-dependent subjects. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008, 147B, 179-186.	1.7	18

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37	Prevalence and Associated Factors of Passive Smoking among Women in Jilin Province, China: A Cross-Sectional Study. <i>International Journal of Environmental Research and Public Health</i> , 2015, 12, 13970-13980.	2.6	18
38	Genome-Wide DNA Methylation Changes Associated with Intermittent Explosive Disorder: A Gene-Based Functional Enrichment Analysis. <i>International Journal of Neuropsychopharmacology</i> , 2018, 21, 12-20.	2.1	17
39	Alcohol and nicotine codependence-associated DNA methylation changes in promoter regions of addiction-related genes. <i>Scientific Reports</i> , 2017, 7, 41816.	3.3	15
40	Prefrontal cortex eQTLs/mQTLs enriched in genetic variants associated with alcohol use disorder and other diseases. <i>Epigenomics</i> , 2020, 12, 789-800.	2.1	15
41	BDNF Variants, Premorbid Educational Attainment, and Disease Characteristics in Alzheimer's Disease: An Exploratory Study. <i>Journal of Alzheimer's Disease</i> , 2009, 17, 887-898.	2.6	13
42	Variation in regulator of G-protein signaling 17 gene (RGS17) is associated with multiple substance dependence diagnoses. <i>Behavioral and Brain Functions</i> , 2012, 8, 23.	3.3	13
43	Identification of POMC Exonic Variants Associated with Substance Dependence and Body Mass Index. <i>PLoS ONE</i> , 2012, 7, e45300.	2.5	13
44	Differential Expression of miR-130a in Postmortem Prefrontal Cortex of Subjects with Alcohol Use Disorders. <i>Journal of Addiction Research & Therapy</i> , 2013, 04, .	0.2	13
45	<i>ADH1A</i> variation predisposes to personality traits and substance dependence. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2010, 153B, 376-386.	1.7	12
46	Association of functional dopamine-beta-hydroxylase (DBH) 19bp insertion/deletion polymorphism with smoking severity in male schizophrenic smokers. <i>Schizophrenia Research</i> , 2012, 141, 48-53.	2.0	12
47	Diverse types of genomic evidence converge on alcohol use disorder risk genes. <i>Journal of Medical Genetics</i> , 2020, 57, 733-743.	3.2	10
48	Atomoxetine in abstinent cocaine users: Cognitive, subjective and cardiovascular effects. <i>Pharmacology Biochemistry and Behavior</i> , 2017, 159, 55-61.	2.9	9
49	An analysis of the effect of mu-opioid receptor gene (OPRM1) promoter region DNA methylation on the response of naltrexone treatment of alcohol dependence. <i>Pharmacogenomics Journal</i> , 2020, 20, 672-680.	2.0	9
50	Association of ATP-binding cassette transporter A1 gene polymorphisms with plasma lipid variability and coronary heart disease risk. <i>International Journal of Clinical and Experimental Pathology</i> , 2015, 8, 13441-9.	0.5	9
51	Effect of Prenatal Opioid Exposure on the Human Placental Methylome. <i>Biomedicines</i> , 2022, 10, 1150.	3.2	9
52	Variation in <i>NGFB</i> is associated with primary affective disorders in women. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2011, 156, 401-412.	1.7	8
53	Internet addiction in college students and its relationship with cigarette smoking and alcohol use in Northeast China. <i>Asia-Pacific Psychiatry</i> , 2017, 9, e12281.	2.2	8
54	RNA m6A Modification Changes in Postmortem Nucleus Accumbens of Subjects with Alcohol Use Disorder: A Pilot Study. <i>Genes</i> , 2022, 13, 958.	2.4	8

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55	Association between single nucleotide variants of vascular endothelial growth factor A and the risk of thyroid carcinoma and nodular goiter in a Han Chinese population. <i>Oncotarget</i> , 2017, 8, 15838-15845.	1.8	7
56	Analyzing Interaction of μ -, δ - and κ -opioid Receptor Gene Variants on Alcohol or Drug Dependence Using a Pattern Discovery-based Method. <i>Journal of Addiction Research & Therapy</i> , 2013, Suppl 7, 007.	0.2	7
57	Association between PLA2G12A Polymorphisms and Schizophrenia in a Han Chinese Population from Northeast China. <i>PLoS ONE</i> , 2016, 11, e0159584.	2.5	6
58	Association of NCOA3 polymorphisms with Dyslipidemia in the Chinese Han population. <i>Lipids in Health and Disease</i> , 2015, 14, 124.	3.0	5
59	A Case-Control Study of the Association between Polymorphisms in the Fibrinogen Alpha Chain Gene and Schizophrenia. <i>Disease Markers</i> , 2017, 2017, 1-5.	1.3	4
60	Genome-wide association study of cognitive flexibility assessed by the Wisconsin Card Sorting Test. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018, 177, 511-519.	1.7	4
61	Analysis of telomere length variation and Shelterin complex subunit gene expression changes in ethanol-exposed human embryonic stem cells. <i>Journal of Psychiatric Research</i> , 2021, 143, 543-549.	3.1	4
62	Association of NCOA2 gene polymorphisms with obesity and dyslipidemia in the Chinese Han population. <i>International Journal of Clinical and Experimental Pathology</i> , 2015, 8, 7341-9.	0.5	4
63	Atomoxetine in abstinent cocaine users: Sex differences. <i>Data in Brief</i> , 2017, 14, 566-572.	1.0	3
64	Profiling of schizophrenia-associated serum peptides by MALDI-TOF-MS. <i>Journal of Neural Transmission</i> , 2020, 127, 95-101.	2.8	2
65	Kininogen-1 as a protein biomarker for schizophrenia through mass spectrometry and genetic association analyses. <i>PeerJ</i> , 2019, 7, e7327.	2.0	2