

Shizhong Han

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10991196/publications.pdf>

Version: 2024-02-01

27
papers

1,740
citations

394421

19
h-index

526287

27
g-index

27
all docs

27
docs citations

27
times ranked

3662
citing authors

#	ARTICLE	IF	CITATIONS
1	Apolipoprotein E gene polymorphism, posttraumatic stress disorder, and cognitive function in older U.S. veterans: Results from the National Health and Resilience in Veterans Study. <i>Depression and Anxiety</i> , 2019, 36, 834-845.	4.1	17
2	Attachment style moderates effects of <i>FKBP5</i> polymorphisms and childhood abuse on post-traumatic stress symptoms: Results from the National Health and Resilience in Veterans Study. <i>World Journal of Biological Psychiatry</i> , 2019, 20, 289-300.	2.6	21
3	Beyond genome-wide significance: integrative approaches to the interpretation and extension of GWAS findings for alcohol use disorder. <i>Addiction Biology</i> , 2019, 24, 275-289.	2.6	15
4	Genetic Risk Variants Associated With Comorbid Alcohol Dependence and Major Depression. <i>JAMA Psychiatry</i> , 2017, 74, 1234.	11.0	74
5	COMBAT: A Combined Association Test for Genes Using Summary Statistics. <i>Genetics</i> , 2017, 207, 883-891.	2.9	39
6	Polygenic Scores for Major Depressive Disorder and Risk of Alcohol Dependence. <i>JAMA Psychiatry</i> , 2017, 74, 1153.	11.0	73
7	Sex-specific linkage scans in opioid dependence. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 261-268.	1.7	10
8	FKBP5 polymorphisms, childhood abuse, and PTSD symptoms: Results from the National Health and Resilience in Veterans Study. <i>Psychoneuroendocrinology</i> , 2016, 69, 98-105.	2.7	66
9	GLITTER: a web-based application for gene link inspection through tissue-specific coexpression. <i>Scientific Reports</i> , 2016, 6, 33460.	3.3	3
10	FLAGS: A Flexible and Adaptive Association Test for Gene Sets Using Summary Statistics. <i>Genetics</i> , 2016, 202, 919-929.	2.9	11
11	Ordered subset linkage analysis based on admixture proportion identifies new linkage evidence for alcohol dependence in African-Americans. <i>Human Genetics</i> , 2013, 132, 397-403.	3.8	16
12	Integrating GWASs and Human Protein Interaction Networks Identifies a Gene Subnetwork Underlying Alcohol Dependence. <i>American Journal of Human Genetics</i> , 2013, 93, 1027-1034.	6.2	72
13	Distinct Loci in the <i>CHRNA5</i> / <i>CHRNA3</i> / <i>CHRN4</i> Gene Cluster Are Associated With Onset of Regular Smoking. <i>Genetic Epidemiology</i> , 2013, 37, 846-859.	1.3	32
14	Increased Genetic Vulnerability to Smoking at <i>CHRNA5</i> in Early-Onset Smokers. <i>Archives of General Psychiatry</i> , 2012, 69, 854.	12.3	71
15	Linkage Analysis Followed by Association Show <i>NRG1</i> Associated with Cannabis Dependence in African Americans. <i>Biological Psychiatry</i> , 2012, 72, 637-644.	1.3	46
16	Autosomal linkage scan for loci predisposing to comorbid dependence on multiple substances. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2012, 159B, 361-369.	1.7	14
17	Association of <i>CHRNA4</i> polymorphisms with smoking behavior in two populations. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2011, 156, 421-429.	1.7	27
18	Multi-locus Test Conditional on Confirmed Effects Leads to Increased Power in Genome-wide Association Studies. <i>PLoS ONE</i> , 2010, 5, e15006.	2.5	14

#	ARTICLE	IF	CITATIONS
19	Multiple Independent Loci at Chromosome 15q25.1 Affect Smoking Quantity: a Meta-Analysis and Comparison with Lung Cancer and COPD. <i>PLoS Genetics</i> , 2010, 6, e1001053.	3.5	332
20	Meta-Analysis of 15 Genome-Wide Linkage Scans of Smoking Behavior. <i>Biological Psychiatry</i> , 2010, 67, 12-19.	1.3	61
21	Evaluation of imputation-based association in and around the integrin- α M (ITGAM) gene and replication of robust association between a non-synonymous functional variant within ITGAM and systemic lupus erythematosus (SLE). <i>Human Molecular Genetics</i> , 2009, 18, 1171-1180.	2.9	100
22	A nonsynonymous functional variant in integrin- α M (encoded by ITGAM) is associated with systemic lupus erythematosus. <i>Nature Genetics</i> , 2008, 40, 152-154.	21.4	277
23	Osteopontin and Systemic Lupus Erythematosus Association: A Probable Gene-Gender Interaction. <i>PLoS ONE</i> , 2008, 3, e0001757.	2.5	79
24	Different roles of MTHFR C677T and A1298C polymorphisms in colorectal adenoma and colorectal cancer: a meta-analysis. <i>Journal of Human Genetics</i> , 2007, 52, 73-85.	2.3	83
25	CTLA4 Polymorphisms and Ophthalmopathy in Graves's Disease Patients: Association Study and Meta-Analysis. <i>Human Immunology</i> , 2006, 67, 618-626.	2.4	48
26	DNA repair gene XRCC3 polymorphisms and cancer risk: a meta-analysis of 48 case-control studies. <i>European Journal of Human Genetics</i> , 2006, 14, 1136-1144.	2.8	85
27	Meta-analysis of the association of CTLA-4 exon-1 +49A/G polymorphism with rheumatoid arthritis. <i>Human Genetics</i> , 2005, 118, 123-132.	3.8	54