## Dongbing Lai

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1342768/publications.pdf

Version: 2024-02-01

361413 289244 2,075 55 20 citations h-index papers

g-index 64 64 64 3907 docs citations times ranked citing authors all docs

40

#	Article	IF	CITATIONS
1	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. Nature Neuroscience, 2018, 21, 1656-1669.	14.8	490
2	A large-scale genome-wide association study meta-analysis of cannabis use disorder. Lancet Psychiatry,the, 2020, 7, 1032-1045.	7.4	200
3	Multivariate analysis of 1.5 million people identifies genetic associations with traits related to self-regulation and addiction. Nature Neuroscience, 2021, 24, 1367-1376.	14.8	137
4	Genome-Wide Association Studies for Taxane-Induced Peripheral Neuropathy in ECOG-5103 and ECOG-1199. Clinical Cancer Research, 2015, 21, 5082-5091.	7.0	106
5	Meta-analysis of up to 622,409 individuals identifies 40 novel smoking behaviour associated genetic loci. Molecular Psychiatry, 2020, 25, 2392-2409.	7.9	83
6	Leveraging genome-wide data to investigate differences between opioid use vs. opioid dependence in 41,176 individuals from the Psychiatric Genomics Consortium. Molecular Psychiatry, 2020, 25, 1673-1687.	7.9	82
7	Genome-Wide Association Study for Anthracycline-Induced Congestive Heart Failure. Clinical Cancer Research, 2017, 23, 43-51.	7.0	73
8	Whole-Exome Sequencing in Familial Parkinson Disease. JAMA Neurology, 2016, 73, 68.	9.0	71
9	Exome Chip Meta-analysis Fine Maps Causal Variants and Elucidates the Genetic Architecture of Rare Coding Variants in Smoking and AlcoholÂUse. Biological Psychiatry, 2019, 85, 946-955.	1.3	69
10	Genomeâ€wide association studies of alcohol dependence, DSMâ€N criterion count and individual criteria. Genes, Brain and Behavior, 2019, 18, e12579.	2.2	56
11	Genome-Wide Association Study of Intracranial Aneurysm Identifies a New Association on Chromosome 7. Stroke, 2014, 45, 3194-3199.	2.0	52
12	Shared Genetic Risk Factors of Intracranial, Abdominal, and Thoracic Aneurysms. Journal of the American Heart Association, $2016, 5, \ldots$	3.7	45
13	Charcot-Marie-Tooth gene, SBF2, associated with taxane-induced peripheral neuropathy in African Americans. Oncotarget, 2016, 7, 82244-82253.	1.8	35
14	Multi-omics integration analysis identifies novel genes for alcoholism with potential overlap with neurodegenerative diseases. Nature Communications, 2021, 12, 5071.	12.8	34
15	Lessons Learned from Whole Exome Sequencing in Multiplex Families Affected by a Complex Genetic Disorder, Intracranial Aneurysm. PLoS ONE, 2015, 10, e0121104.	2.5	32
16	Genomewide Association Studies of <scp><i>LRRK2</i></scp> Modifiers of Parkinson's Disease. Annals of Neurology, 2021, 90, 76-88.	5.3	30
17	Shared genetic risk between eating disorderâ€and substanceâ€useâ€related phenotypes: Evidence from genomeâ€wide association studies. Addiction Biology, 2021, 26, e12880.	2.6	28
18	Allele-specific expression and high-throughput reporter assay reveal functional genetic variants associated with alcohol use disorders. Molecular Psychiatry, 2021, 26, 1142-1151.	7.9	26

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19	Genomewide Metaâ€Analysis Validates a Role for <i>S1PR1</i> in Microtubule Targeting Agentâ€Induced Sensory Peripheral Neuropathy. Clinical Pharmacology and Therapeutics, 2020, 108, 625-634.	4.7	25
20	Exome Sequencing Identifies Candidate Genetic Modifiers of Syndromic and Familial Thoracic Aortic Aneurysm Severity. Journal of Cardiovascular Translational Research, 2017, 10, 423-432.	2.4	24
21	Impact of Genetic Ancestry on Outcomes in ECOG-ACRIN-5103. JCO Precision Oncology, 2017, 2017, 1-9.	3.0	23
22	A genome-wide association study of interhemispheric theta EEG coherence: implications for neural connectivity and alcohol use behavior. Molecular Psychiatry, 2021, 26, 5040-5052.	7.9	22
23	Calcium-Sensing Receptor Genotype and Response to Cinacalcet in Patients Undergoing Hemodialysis. Clinical Journal of the American Society of Nephrology: CJASN, 2017, 12, 1128-1138.	4.5	21
24	Metaâ€Analysis of Genetic Influences on Initial Alcohol Sensitivity. Alcoholism: Clinical and Experimental Research, 2018, 42, 2349-2359.	2.4	21
25	Genomeâ€wide association studies of the selfâ€rating of effects of ethanol (SRE). Addiction Biology, 2020, 25, e12800.	2.6	20
26	Polygenic contributions to alcohol use and alcohol use disorders across population-based and clinically ascertained samples. Psychological Medicine, 2021, 51, 1147-1156.	4.5	18
27	A multivariate finite mixture latent trajectory model with application to dementia studies. Journal of Applied Statistics, 2016, 43, 2503-2523.	1.3	16
28	Evaluating risk for alcohol use disorder: Polygenic risk scores and family history. Alcoholism: Clinical and Experimental Research, 2022, 46, 374-383.	2.4	16
29	Genomeâ€wide association study identifies loci associated with liability to alcohol and drug dependence that is associated with variability in rewardâ€related ventral striatum activity in African― and Europeanâ€Americans. Genes, Brain and Behavior, 2019, 18, e12580.	2.2	15
30	The Genetic Relationship Between Alcohol Consumption and Aspects of Problem Drinking in an Ascertained Sample. Alcoholism: Clinical and Experimental Research, 2019, 43, 1113-1125.	2.4	15
31	Polymorphisms in the bone morphogenetic protein 2 (BMP2) gene do not affect bone mineral density in white men or women. Osteoporosis International, 2006, 17, 587-592.	3.1	13
32	Mapping Pathways by Which Genetic Risk Influences Adolescent Externalizing Behavior: The Interplay Between Externalizing Polygenic Risk Scores, Parental Knowledge, and Peer Substance Use. Behavior Genetics, 2021, 51, 543-558.	2.1	13
33	Genetic Influences on Plasma Homocysteine Levels in African Americans and Yoruba Nigerians. Journal of Alzheimer's Disease, 2016, 49, 991-1003.	2.6	12
34	Genome-wide association analysis identifies new candidate risk loci for familial intracranial aneurysm in the French-Canadian population. Scientific Reports, 2018, 8, 4356.	3.3	12
35	Integration of evidence across human and model organism studies: A meeting report. Genes, Brain and Behavior, 2021, 20, e12738.	2.2	12
36	Genomeâ€wide admixture mapping of <scp>DSMâ€IV</scp> alcohol dependence, criterion count, and the selfâ€rating of the effects of ethanol in African American populations. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2021, 186, 151-161.	1.7	11

#	Article	lF	Citations
37	SIBLING family genes and bone mineral density: Association and allele-specific expression in humans. Bone, 2014, 64, 166-172.	2.9	10
38	Psychosocial moderation of polygenic risk for cannabis involvement: the role of trauma exposure and frequency of religious service attendance. Translational Psychiatry, 2019, 9, 269.	4.8	10
39	Gene-based polygenic risk scores analysis of alcohol use disorder in African Americans. Translational Psychiatry, 2022, 12, .	4.8	10
40	Steroid Pathway Genes and Neonatal Respiratory Distress After Betamethasone Use in Anticipated Preterm Birth. Reproductive Sciences, 2016, 23, 680-686.	2.5	9
41	Identification of Functional Genetic Variants Associated With Alcohol Dependence and Related Phenotypes Using a Highâ€Throughput Assay. Alcoholism: Clinical and Experimental Research, 2020, 44, 2494-2518.	2.4	7
42	Investigation of convergent and divergent genetic influences underlying schizophrenia and alcohol use disorder. Psychological Medicine, 2023, 53, 1196-1204.	4.5	7
43	High Polygenic Risk Scores Are Associated With Early Age of Onset of Alcohol Use Disorder in Adolescents and Young Adults at Risk. Biological Psychiatry Global Open Science, 2022, 2, 379-388.	2.2	7
44	Alcohol use disorder, psychiatric comorbidities, marriage and divorce in a high-risk sample Psychology of Addictive Behaviors, 2022, 36, 364-374.	2.1	7
45	Exome-chip association analysis of intracranial aneurysms. Neurology, 2020, 94, e481-e488.	1.1	5
46	Genes identified in rodent studies of alcohol intake are enriched for heritability of human substance use. Alcoholism: Clinical and Experimental Research, 2021, 45, 2485-2494.	2.4	5
47	Genome-wide association study of serum iron phenotypes in premenopausal women of European descent. Blood Cells, Molecules, and Diseases, 2016, 57, 50-53.	1.4	3
48	A regulatory variant of CHRM3 is associated with cannabis-induced hallucinations in European Americans. Translational Psychiatry, 2019, 9, 309.	4.8	3
49	Identification of a common polymorphism in COQ8B acting as a modifier of thoracic aortic aneurysm severity Human Genetics and Genomics Advances, 2021, 3, 100057.	1.7	3
50	Epigenetic changes on rat chromosome 4 contribute to disparate alcohol drinking behavior in alcohol-preferring and -nonpreferring rats. Alcohol, 2020, 89, 103-112.	1.7	2
51	Genome wide association study for anthracycline-induced congestive heart failure Journal of Clinical Oncology, 2016, 34, 1017-1017.	1.6	2
52	Examining social genetic effects on educational attainment via parental educational attainment, income, and parenting Journal of Family Psychology, 2022, 36, 1340-1350.	1.3	2
53	Testing influences of APOE and BDNF genes and heart failure on cognitive function. Heart and Lung: Journal of Acute and Critical Care, 2021, 50, 51-58.	1.6	1
54	P2â€509: HEART FAILURE PREDICTS COGNITIVE DYSFUNCTION AFTER ADJUSTING FOR ⟨i⟩APOE⟨/i⟩ FACTORS. Alzheimer's and Dementia, 2018, 14, P927.	0.8	0

#	Article	IF	CITATIONS
55	Associations of targeted genetic variants with Alzheimer's disease in African Americans and Nigerians. Alzheimer's and Dementia, 2020, 16, e042323.	0.8	O