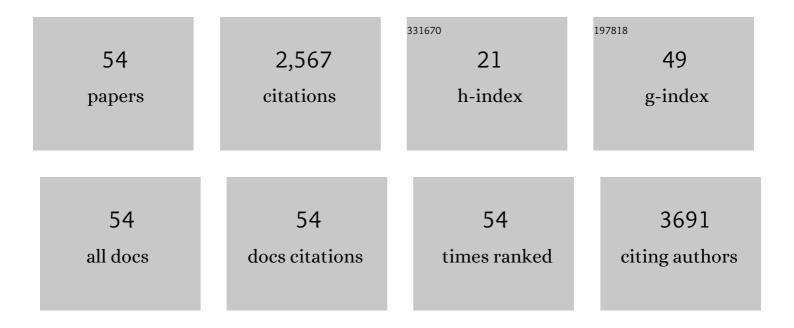
Ming D Li

List of Publications by Year in descending order

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MINCDL

#	Article	IF	CITATIONS
1	Determination of shared genetic etiology and possible causal relations between tobacco smoking and depression. Psychological Medicine, 2021, 51, 1870-1879.	4.5	15
2	Geneâ€based association analysis reveals involvement of LAMA5 and cell adhesion pathways in nicotine dependence in African―and Europeanâ€American samples. Addiction Biology, 2021, 26, e12898.	2.6	2
3	Genome-wide methylation and expression analyses reveal the epigenetic landscape of immune-related diseases for tobacco smoking. Clinical Epigenetics, 2021, 13, 215.	4.1	13
4	Demonstration of critical role of <i>GRIN3A</i> in nicotine dependence through both genetic association and molecular functional studies. Addiction Biology, 2020, 25, e12718.	2.6	8
5	Genetic and Epigenetic Analysis Revealing Variants in the NCAM1–TTC12–ANKK1–DRD2 Cluster Associated Significantly With Nicotine Dependence in Chinese Han Smokers. Nicotine and Tobacco Research, 2020, 22, 1301-1309.	2.6	11
6	Genome-wide DNA methylation analysis reveals significant impact of long-term ambient air pollution exposure on biological functions related to mitochondria and immune response. Environmental Pollution, 2020, 264, 114707.	7.5	32
7	Identification of 34 genes conferring genetic and pharmacological risk for the comorbidity of schizophrenia and smoking behaviors. Aging, 2020, 12, 2169-2225.	3.1	15
8	Identification and characterization of SEC24D as a susceptibility gene for hepatitis B virus infection. Scientific Reports, 2019, 9, 13425.	3.3	2
9	An Exome-Wide Association Study Identifies New Susceptibility Loci for Age of Smoking Initiation in African- and European-American Populations. Nicotine and Tobacco Research, 2019, 21, 707-713.	2.6	6
10	The Prevalence of Smoking and Its Associated Diseases. , 2018, , 1-11.		0
11	Tobacco Smoking Addiction: Epidemiology, Genetics, Mechanisms, and Treatment. , 2018, , .		7
12	Association and cis-mQTL analysis of variants in CHRNA3-A5, CHRNA7, CHRNB2, and CHRNB4 in relation to nicotine dependence in a Chinese Han population. Translational Psychiatry, 2018, 8, 83.	4.8	21
13	DNA Methylation Analysis Reveals a Strong Connection Between Tobacco Smoking and Cancer Pathogenesis. , 2018, , 303-317.		0
14	Implication of Genes for the N-Methyl-d-Aspartate (NMDA) Receptor in Substance Addictions. Molecular Neurobiology, 2018, 55, 7567-7578.	4.0	14
15	Modulatory Effects of Nicotine on neuroHIV/neuroAIDS. Journal of NeuroImmune Pharmacology, 2018, 13, 467-478.	4.1	7
16	Establishment of a Strong Link Between Smoking and Cancer Pathogenesis through DNA Methylation Analysis. Scientific Reports, 2017, 7, 1811.	3.3	59
17	HIV-1 Proteins Influence Novelty-Seeking Behavior and Alter Region-Specific Transcriptional Responses to Chronic Nicotine Treatment in HIV-1Tg Rats. Nicotine and Tobacco Research, 2017, 19, 1024-1032.	2.6	8
18	Significant association of the CHRNB3-CHRNA6 gene cluster with nicotine dependence in the Chinese Han population. Scientific Reports, 2017, 7, 9745.	3.3	11

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19	Relationship between Personality Traits and Nicotine Dependence in Male and Female Smokers of African-American and European-American Samples. Frontiers in Psychiatry, 2017, 8, 122.	2.6	20
20	Prevalence of Cigarette Smoking and Nicotine Dependence in Men and Women Residing in Two Provinces in China. Frontiers in Psychiatry, 2017, 8, 254.	2.6	29
21	Expression profile of nicotinic acetylcholine receptor subunits in the brain of HIV-1 transgenic rats given chronic nicotine treatment. Journal of NeuroVirology, 2016, 22, 626-633.	2.1	6
22	Modulation Effect of HIV-1 Viral Proteins and Nicotine on Expression of the Immune-Related Genes in Brain of the HIV-1 Transgenic Rats. Journal of NeuroImmune Pharmacology, 2016, 11, 562-571.	4.1	17
23	Association of STAT4 polymorphisms with hepatitis B virus infection and clearance in Chinese Han population. Amino Acids, 2016, 48, 2589-2598.	2.7	15
24	Promoting Global Health — Prevention and Treatment of Substance Abuse and HIV in Asia. Journal of NeuroImmune Pharmacology, 2016, 11, 379-382.	4.1	0
25	Interactive Effects of Ethanol and <scp>HIV</scp> â€1 Proteins on Noveltyâ€6eeking Behaviors and Addictionâ€Related Gene Expression. Alcoholism: Clinical and Experimental Research, 2016, 40, 2102-2113.	2.4	9
26	Meta-Analysis of the COMT Val158Met Polymorphism in Major Depressive Disorder: Effect of Ethnicity. Journal of NeuroImmune Pharmacology, 2016, 11, 434-445.	4.1	38
27	Meta-Analysis Reveals Significant Association of the 3′-UTR VNTR in <i>SLC6A3</i> with Alcohol Dependence. Alcoholism: Clinical and Experimental Research, 2016, 40, 1443-1453.	2.4	19
28	Contribution of Variants in CHRNA5/A3/B4 Gene Cluster on Chromosome 15 to Tobacco Smoking: From Genetic Association to Mechanism. Molecular Neurobiology, 2016, 53, 472-484.	4.0	49
29	Novelty Seeking and Drug Addiction in Humans and Animals: From Behavior to Molecules. Journal of NeuroImmune Pharmacology, 2016, 11, 456-470.	4.1	112
30	Nicotine mediates expression of genes related to antioxidant capacity and oxidative stress response in HIV-1 transgenic rat brain. Journal of NeuroVirology, 2016, 22, 114-124.	2.1	12
31	Ethnic-Specific Genetic Association of Variants in the Corticotropin-Releasing Hormone Receptor 1 Gene with Nicotine Dependence. BioMed Research International, 2015, 2015, 1-7.	1.9	5
32	Updated Findings of the Association and Functional Studies of DRD2/ANKK1 Variants with Addictions. Molecular Neurobiology, 2015, 51, 281-299.	4.0	25
33	Nicotine attenuates the effect of HIV-1 proteins on the neural circuits of working and contextual memories. Molecular Brain, 2015, 8, 43.	2.6	21
34	Fine mapping analysis of HLA-DP/DQ gene clusters on chromosome 6 reveals multiple susceptibility loci for HBV infection. Amino Acids, 2015, 47, 2623-2634.	2.7	10
35	Significant associations of CHRNA2 and CHRNA6 with nicotine dependence in European American and African American populations. Human Genetics, 2014, 133, 575-586.	3.8	39
36	Association and interaction analyses of 5-HT3 receptor and serotonin transporter genes with alcohol, cocaine, and nicotine dependence using the SAGE data. Human Genetics, 2014, 133, 905-918.	3.8	28

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37	Acquisition and long-term retention of spatial learning in the human immunodeficiency virus-1 transgenic rat: effects of repeated nicotine treatment. Journal of NeuroVirology, 2013, 19, 157-165.	2.1	18
38	Nine Generations of Selection for High and Low Nicotine Intake in Outbred Sprague–Dawley Rats. Behavior Genetics, 2013, 43, 436-444.	2.1	14
39	Depressive Symptoms Among Heavy Cigarette Smokers: The Influence of Daily Rate, Gender, and Race. Nicotine and Tobacco Research, 2013, 15, 1714-1721.	2.6	21
40	Identification and Characterization of Poly(I:C)-induced Molecular Responses Attenuated by Nicotine in Mouse Macrophages. Molecular Pharmacology, 2013, 83, 61-72.	2.3	39
41	RNA Deep Sequencing Analysis Reveals That Nicotine Restores Impaired Gene Expression by Viral Proteins in the Brains of HIV-1 Transgenic Rats. PLoS ONE, 2013, 8, e68517.	2.5	16
42	Transcriptome Sequencing of Gene Expression in the Brain of the HIV-1 Transgenic Rat. PLoS ONE, 2013, 8, e59582.	2.5	35
43	Association and interaction analysis of variants in <i>CHRNA5/CHRNA3/CHRNB4</i> gene cluster with nicotine dependence in African and European Americans. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 745-756.	1.7	53
44	Significant association of glutamate receptor, ionotropic N-methyl-d-aspartate 3A (GRIN3A), with nicotine dependence in European- and African-American smokers. Human Genetics, 2010, 127, 503-512.	3.8	18
45	Multiple Independent Loci at Chromosome 15q25.1 Affect Smoking Quantity: a Meta-Analysis and Comparison with Lung Cancer and COPD. PLoS Genetics, 2010, 6, e1001053.	3.5	332
46	Associations of Variants in CHRNA5/A3/B4 Gene Cluster with Smoking Behaviors in a Korean Population. PLoS ONE, 2010, 5, e12183.	2.5	57
47	New insights into the genetics of addiction. Nature Reviews Genetics, 2009, 10, 225-231.	16.3	207
48	Identifying susceptibility loci for nicotine dependence: 2008 update based on recent genome-wide linkage analyses. Human Genetics, 2008, 123, 119-131.	3.8	89
49	Significant Association of Catechol-O-Methyltransferase (COMT) Haplotypes with Nicotine Dependence in Male and Female Smokers of Two Ethnic Populations. Neuropsychopharmacology, 2006, 31, 675-684.	5.4	141
50	Ethnic- and gender-specific association of the nicotinic acetylcholine receptor α4 subunit gene (CHRNA4) with nicotine dependence. Human Molecular Genetics, 2005, 14, 1211-1219.	2.9	182
51	Single- and Multilocus Allelic Variants within the GABAB Receptor Subunit 2 (GABAB2) Gene Are Significantly Associated with Nicotine Dependence. American Journal of Human Genetics, 2005, 76, 859-864.	6.2	99
52	Time-dependent changes in transcriptional profiles within five rat brain regions in response to nicotine treatment. Molecular Brain Research, 2004, 132, 168-180.	2.3	61
53	A metaâ€analysis of estimated genetic and environmental effects on smoking behavior in male and female adult twins. Addiction, 2003, 98, 23-31.	3.3	499
54	Identification of a Novel Functional Non-synonymous Single Nucleotide Polymorphism in Frizzled Class Receptor 6 Gene for Involvement in Depressive Symptoms. Frontiers in Molecular Neuroscience, 0, 15, .	2.9	1