## Joel Gelernter

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4343116/publications.pdf

Version: 2024-02-01

		2100	5987
556	38,933	100	160
papers	citations	h-index	g-index
635	635	635	28148

times ranked

citing authors

docs citations

all docs

#	Article	IF	CITATIONS
1	Discovery of the first genome-wide significant risk loci for attention deficit/hyperactivity disorder. Nature Genetics, 2019, 51, 63-75.	21.4	1,594
2	Synonymous mutations in the human dopamine receptor D2 (DRD2) affect mRNA stability and synthesis of the receptor. Human Molecular Genetics, 2003, 12, 205-216.	2.9	800
3	Social supports and serotonin transporter gene moderate depression in maltreated children. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 17316-17321.	7.1	757
4	Brain-Derived Neurotrophic Factor–5-HTTLPR Gene Interactions and Environmental Modifiers of Depression in Children. Biological Psychiatry, 2006, 59, 673-680.	1.3	655
5	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. Nature Genetics, 2021, 53, 817-829.	21.4	629
6	A Functional Polymorphism of the $\hat{l}$ 4-Opioid Receptor Gene is Associated with Naltrexone Response in Alcohol-Dependent Patients. Neuropsychopharmacology, 2003, 28, 1546-1552.	5.4	594
7	Psychiatric Genomics: An Update and an Agenda. American Journal of Psychiatry, 2018, 175, 15-27.	7.2	518
8	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. Nature Neuroscience, 2018, 21, 1656-1669.	14.8	490
9	GWAS of lifetime cannabis use reveals new risk loci, genetic overlap with psychiatric traits, and a causal effect of schizophrenia liability. Nature Neuroscience, 2018, 21, 1161-1170.	14.8	436
10	Serotonin transporter protein (SLC6A4) allele and haplotype frequencies and linkage disequilibria in African- and European-American and Japanese populations and in alcohol-dependent subjects. Human Genetics, 1997, 101, 243-246.	3.8	393
11	The Serotonin Transporter Genotype and Social Support and Moderation of Posttraumatic Stress Disorder and Depression in Hurricane-Exposed Adults. American Journal of Psychiatry, 2007, 164, 1693-1699.	7.2	371
12	International meta-analysis of PTSD genome-wide association studies identifies sex- and ancestry-specific genetic risk loci. Nature Communications, 2019, 10, 4558.	12.8	363
13	Genome-wide association study of alcohol dependence:significant findings in African- and European-Americans including novel risk loci. Molecular Psychiatry, 2014, 19, 41-49.	7.9	353
14	<i>DCDC2</i> is associated with reading disability and modulates neuronal development in the brain. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 17053-17058.	7.1	351
15	Genome-wide association study of alcohol consumption and use disorder in 274,424 individuals from multiple populations. Nature Communications, 2019, 10, 1499.	12.8	346
16	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
17	Interaction of FKBP5 with Childhood Adversity on Risk for Post-Traumatic Stress Disorder. Neuropsychopharmacology, 2010, 35, 1684-1692.	5.4	299
18	A Prospective Cohort Study Investigating Factors Associated With Depression During Medical Internship. Archives of General Psychiatry, 2010, 67, 557.	12.3	282

#	Article	IF	CITATIONS
19	Child Abuse, Depression, and Methylation in Genes Involved With Stress, Neural Plasticity, and Brain Circuitry. Journal of the American Academy of Child and Adolescent Psychiatry, 2014, 53, 417-424.e5.	0.5	273
20	Genetics of two $\hat{l}\frac{1}{4}$ opioid receptor gene (OPRM1) exon I polymorphisms: population studies, and allele frequencies in alcohol- and drug-dependent subjects. Molecular Psychiatry, 1999, 4, 476-483.	7.9	266
21	Prediction of Dopamine Transporter Binding Availability by Genotype: A Preliminary Report. American Journal of Psychiatry, 2000, 157, 1700-1703.	7.2	263
22	Allelic and haplotypic association of GABRA2 with alcohol dependence. American Journal of Medical Genetics Part A, 2004, 129B, 104-109.	2.4	263
23	Interaction of Childhood Maltreatment with the Corticotropin-Releasing Hormone Receptor Gene: Effects on Hypothalamic-Pituitary-Adrenal Axis Reactivity. Biological Psychiatry, 2009, 66, 681-685.	1.3	254
24	A Quantitative-Trait Analysis of Human Plasma–Dopamine β-Hydroxylase Activity: Evidence for a Major Functional Polymorphism at the DBH Locus. American Journal of Human Genetics, 2001, 68, 515-522.	6.2	253
25	Genome-wide meta-analysis of problematic alcohol use in 435,563 individuals yields insights into biology and relationships with other traits. Nature Neuroscience, 2020, 23, 809-818.	14.8	242
26	Addictions Biology: Haplotype-Based Analysis for 130 Candidate Genes on a Single Array. Alcohol and Alcoholism, 2008, 43, 505-515.	1.6	222
27	Interactive Effect of Stressful Life Events and the Serotonin Transporter 5-HTTLPR Genotype on Posttraumatic Stress Disorder Diagnosis in 2 Independent Populations. Archives of General Psychiatry, 2009, 66, 1201.	12.3	221
28	Child Abuse and Epigenetic Mechanisms of Disease Risk. American Journal of Preventive Medicine, 2013, 44, 101-107.	3.0	212
29	NO Association Between an Allele at the D2 Dopamine Receptor Gene (DRD2) and Alcoholism. JAMA - Journal of the American Medical Association, 1991, 266, 1801.	7.4	209
30	Bi-ancestral depression GWAS in the Million Veteran Program and meta-analysis in >1.2 million individuals highlight new therapeutic directions. Nature Neuroscience, 2021, 24, 954-963.	14.8	207
31	Gene-by-Environment (Serotonin Transporter and Childhood Maltreatment) Interaction for Anxiety Sensitivity, an Intermediate Phenotype for Anxiety Disorders. Neuropsychopharmacology, 2008, 33, 312-319.	5.4	205
32	Adverse childhood events as risk factors for substance dependence: Partial mediation by mood and anxiety disorders. Addictive Behaviors, 2010, 35, 7-13.	3.0	203
33	A large-scale genome-wide association study meta-analysis of cannabis use disorder. Lancet Psychiatry,the, 2020, 7, 1032-1045.	7.4	200
34	COMT Polymorphisms and Anxiety-Related Personality Traits. Neuropsychopharmacology, 2005, 30, 2092-2102.	5.4	199
35	The A1 Allele at the D2 Dopamine Receptor Gene and Alcoholism. JAMA - Journal of the American Medical Association, 1993, 269, 1673.	7.4	198
36	Genetic Association between Dopamine Transporter Protein Alleles and Cocaine-Induced Paranoia. Neuropsychopharmacology, 1994, 11, 195-200.	5.4	198

#	Article	IF	CITATIONS
37	Population studies of polymorphisms of the serotonin transporter protein gene. American Journal of Medical Genetics Part A, 1999, 88, 61-66.	2.4	193
38	Genome-Wide Association Study of Opioid Dependence: Multiple Associations Mapped to Calcium and Potassium Pathways. Biological Psychiatry, 2014, 76, 66-74.	1.3	192
39	CHRM2 gene predisposes to alcohol dependence, drug dependence and affective disorders: results from an extended case–control structured association study. Human Molecular Genetics, 2005, 14, 2421-2434.	2.9	191
40	Evidence for linkage disequilibrium between serotonin transporter protein gene (SLC6A4) and obsessive compulsive disorder. Molecular Psychiatry, 1998, 3, 270-273.	7.9	190
41	Genetic and Environmental Predictors of Early Alcohol Use. Biological Psychiatry, 2007, 61, 1228-1234.	1.3	189
42	GPA: A Statistical Approach to Prioritizing GWAS Results by Integrating Pleiotropy and Annotation. PLoS Genetics, 2014, 10, e1004787.	<b>3.</b> 5	189
43	Association Between Alcoholism and ??-Amino Butyric Acid ??2 Receptor Subtype in a Russian Population. Alcoholism: Clinical and Experimental Research, 2005, 29, 493-498.	2.4	188
44	Correlates of co-occurring ADHD in drug-dependent subjects: Prevalence and features of substance dependence and psychiatric disorders. Addictive Behaviors, 2008, 33, 1199-1207.	3.0	187
45	The A1 allele at the D2 dopamine receptor gene and alcoholism. A reappraisal. JAMA - Journal of the American Medical Association, 1993, 269, 1673-1677.	7.4	187
46	Reproducible Genetic Risk Loci for Anxiety: Results From â <sup>1</sup> / <sub>4</sub> 200,000 Participants in the Million Veteran Program. American Journal of Psychiatry, 2020, 177, 223-232.	7.2	185
47	Genome-wide association study of cocaine dependence and related traits: FAM53B identified as a risk gene. Molecular Psychiatry, 2014, 19, 717-723.	7.9	182
48	Î <sup>3</sup> -Aminobutyric Acid Type A Receptors and Alcoholism. Archives of General Psychiatry, 2006, 63, 957.	12.3	181
49	Diagnostic reliability of the Semi-structured Assessment for Drug Dependence and Alcoholism (SSADDA). Drug and Alcohol Dependence, 2005, 80, 303-312.	3.2	180
50	A haplotype at the DBH locus, associated with low plasma dopamine $\hat{l}^2$ -hydroxylase activity, also associates with cocaine-induced paranoia. Molecular Psychiatry, 2000, 5, 56-63.	7.9	179
51	Opioid Receptor Gene (OPRM1, OPRK1, and OPRD1) Variants and Response to Naltrexone Treatment for Alcohol Dependence: Results From the VA Cooperative Study. Alcoholism: Clinical and Experimental Research, 2007, 31, 070212174136005-???.	2.4	178
52	No association between an allele at the D2 dopamine receptor gene (DRD2) and alcoholism. JAMA - Journal of the American Medical Association, 1991, 266, 1801-1807.	7.4	175
53	D4 Dopamine-Receptor (DRD4) Alleles and Novelty Seeking in Substance-Dependent, Personality-Disorder, and Control Subjects. American Journal of Human Genetics, 1997, 61, 1144-1152.	6.2	174
54	Genome-Wide Association Studies of a Broad Spectrum of Antisocial Behavior. JAMA Psychiatry, 2017, 74, 1242.	11.0	174

#	Article	IF	CITATIONS
55	MAOA Genotype, Maltreatment, and Aggressive Behavior: The Changing Impact of Genotype at Varying Levels of Trauma. Biological Psychiatry, 2009, 65, 417-424.	1.3	173
56	The Variable Number of Tandem Repeats Polymorphism of the Dopamine Transporter Gene Is Not Associated with Significant Change in Dopamine Transporter Phenotype in Humans. Neuropsychopharmacology, 2001, 24, 553-560.	5.4	171
57	Brain derived neurotrophic factor ( <i>BDNF</i> ) gene variants and Alzheimer's disease, affective disorders, posttraumatic stress disorder, schizophrenia, and substance dependence. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2006, 141B, 387-393.	1.7	170
58	A Functional Neuropeptide Y Leu7Pro Polymorphism Associated With Alcohol Dependence in a Large Population Sample From the United States. Archives of General Psychiatry, 2002, 59, 825.	12.3	165
59	Possible association of a polymorphism of the tryptophan hydroxylase gene with suicidal behavior in depressed patients. American Journal of Psychiatry, 1997, 154, 1451-1453.	7.2	164
60	Topiramate Treatment for Heavy Drinkers: Moderation by a <i>GRIK1</i> Polymorphism. American Journal of Psychiatry, 2014, 171, 445-452.	7.2	164
61	Serotonin Transporter Protein Gene Polymorphism and Personality Measures in African American and European American Subjects. American Journal of Psychiatry, 1998, 155, 1332-1338.	7.2	163
62	Genome-wide Association Studies of Posttraumatic Stress Disorder in 2 Cohorts of US Army Soldiers. JAMA Psychiatry, 2016, 73, 695.	11.0	158
63	Haplotype spanning TTC12 and ANKK1, flanked by the DRD2 and NCAM1 loci, is strongly associated to nicotine dependence in two distinct American populations. Human Molecular Genetics, 2006, 15, 3498-3507.	2.9	156
64	Association between two $\hat{A}\mu$ -opioid receptor gene (OPRM1) haplotype blocks and drug or alcohol dependence. Human Molecular Genetics, 2006, 15, 807-819.	2.9	155
65	Strong Association of the Alcohol Dehydrogenase 1B Gene (ADH1B) with Alcohol Dependence and Alcohol-Induced Medical Diseases. Biological Psychiatry, 2011, 70, 504-512.	1.3	150
66	Genome-wide Association Study Identifies New Susceptibility Loci for Posttraumatic Stress Disorder. Biological Psychiatry, 2013, 74, 656-663.	1.3	150
67	Genome-wide Association Study of Cannabis Dependence Severity, Novel Risk Variants, and Shared Genetic Risks. JAMA Psychiatry, 2016, 73, 472.	11.0	148
68	D2 receptors binding potential is not affected by Taq1 polymorphism at the D2 receptor gene. Molecular Psychiatry, 1998, 3, 261-265.	7.9	146
69	Modification of the Association Between Serotonin Transporter Genotype and Risk of Posttraumatic Stress Disorder in Adults by County-Level Social Environment. American Journal of Epidemiology, 2009, 169, 704-711.	3.4	146
70	A Genetic Investigation of Sex Bias in the Prevalence of Attention-Deficit/Hyperactivity Disorder. Biological Psychiatry, 2018, 83, 1044-1053.	1.3	146
71	Genome-wide association study of post-traumatic stress disorder reexperiencing symptoms in >165,000 US veterans. Nature Neuroscience, 2019, 22, 1394-1401.	14.8	145
72	The OPRD1 and OPRK1 loci in alcohol or drug dependence: OPRD1 variation modulates substance dependence risk. Molecular Psychiatry, 2008, 13, 531-543.	7.9	143

#	Article	IF	Citations
73	Variant Callers for Next-Generation Sequencing Data: A Comparison Study. PLoS ONE, 2013, 8, e75619.	2.5	139
74	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
75	Genome-wide association study of lifetime cannabis use based on a large meta-analytic sample of 32 330 subjects from the International Cannabis Consortium. Translational Psychiatry, 2016, 6, e769-e769.	4.8	136
76	Association of <i>OPRM1</i> Functional Coding Variant With Opioid Use Disorder. JAMA Psychiatry, 2020, 77, 1072.	11.0	135
77	Serotonin transporter gene promoter polymorphism predicts SSRI response in generalized social anxiety disorder. Psychopharmacology, 2006, 187, 68-72.	3.1	134
78	Influence of RGS2 on Anxiety-Related Temperament, Personality, and Brain Function. Archives of General Psychiatry, 2008, 65, 298.	12.3	134
79	Association between the cortisol response to opioid blockade and the Asn40Asp polymorphism at the μâ€opioid receptor locus (OPRM1). American Journal of Medical Genetics Part A, 2003, 118B, 60-65.	2.4	132
80	Strong protective effect of the aldehyde dehydrogenase gene (ALDH2) 504lys (*2) allele against alcoholism and alcohol-induced medical diseases in Asians. Human Genetics, 2012, 131, 725-737.	3.8	132
81	Sex-biased methylome and transcriptome in human prefrontal cortex. Human Molecular Genetics, 2014, 23, 1260-1270.	2.9	130
82	Dopamine $\hat{l}^2$ -hydroxylase: two polymorphisms in linkage disequilibrium at the structural gene DBH associate with biochemical phenotypic variation. Human Genetics, 1998, 102, 533-540.	3.8	127
83	Genomewide Linkage Scan for Opioid Dependence and Related Traits. American Journal of Human Genetics, 2006, 78, 759-769.	6.2	125
84	Reliability of DSM-IV diagnostic criteria using the semi-structured assessment for drug dependence and alcoholism (SSADDA). Drug and Alcohol Dependence, 2007, 91, 85-90.	3.2	124
85	Central Serotonin Transporter Availability Measured With [ <sup>123</sup> 1]β-CIT SPECT in Relation to Serotonin Transporter Genotype. American Journal of Psychiatry, 2004, 161, 525-531.	7.2	122
86	Genome-wide association analyses of post-traumatic stress disorder and its symptom subdomains in the Million Veteran Program. Nature Genetics, 2021, 53, 174-184.	21.4	121
87	The D4 dopamine receptor (DRD4) maps to distal 11p close to HRAS. Genomics, 1992, 13, 208-210.	2.9	120
88	Corpus callosum dimensions measured by magnetic resonance imaging in bipolar affective disorder and schizophrenia. Biological Psychiatry, 1989, 26, 659-668.	1.3	119
89	Genomewide linkage scan for cocaine dependence and related traits: Significant linkages for a cocaine-related trait and cocaine-induced paranoia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2005, 136B, 45-52.	1.7	119
90	Association of haplotypic variants in DRD2, ANKK1, TTC12 and NCAM1 to alcohol dependence in independent case–control and family samples. Human Molecular Genetics, 2007, 16, 2844-2853.	2.9	118

#	Article	IF	Citations
91	Serotonin transporter gene associations with psychopathic traits in youth vary as a function of socioeconomic resources Journal of Abnormal Psychology, 2010, 119, 604-609.	1.9	118
92	Genotyping Array Design and Data Quality Control in the Million Veteran Program. American Journal of Human Genetics, 2020, 106, 535-548.	6.2	118
93	Tryptophan hydroxylase genotype is associated with impulsive-aggression measures: a preliminary study. American Journal of Medical Genetics Part A, 1998, 81, 13-17.	2.4	114
94	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. Biological Psychiatry, 2022, 91, 313-327.	1.3	114
95	No Linkage Between D2 Dopamine Receptor Gene Region and Schizophrenia. Archives of General Psychiatry, 1991, 48, 643.	12.3	113
96	Genetic variation in 5HTTLPR is associated with emotional resilience. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2009, 150B, 900-906.	1.7	112
97	Diplotype Trend Regression Analysis of the ADH Gene Cluster and the ALDH2 Gene: Multiple Significant Associations with Alcohol Dependence. American Journal of Human Genetics, 2006, 78, 973-987.	6.2	110
98	No association between D2 dopamine receptor (DRD2) "A―system alleles, or DRD2 haplotypes, and posttraumatic stress disorder. Biological Psychiatry, 1999, 45, 620-625.	1.3	109
99	Genetics of alcohol dependence. Human Genetics, 2009, 126, 91-99.	3.8	109
100	Evidence of CNIH3 involvement in opioid dependence. Molecular Psychiatry, 2016, 21, 608-614.	7.9	109
101	Genetic–epigenetic interactions in cis: a major focus in the post-GWAS era. Genome Biology, 2017, 18, 120.	8.8	109
102	Allelic Variation in the D4 Dopamine Receptor (DRD4) Gene Does Not Predict Response to Clozapine. Archives of General Psychiatry, 1994, 51, 912.	12.3	107
103	Practical population group assignment with selected informative markers: Characteristics and properties of Bayesian clustering via STRUCTURE. Genetic Epidemiology, 2005, 28, 302-312.	1.3	106
104	Genetic influences on eight psychiatric disorders based on family data of 4 408 646 full and half-siblings, and genetic data of 333 748 cases and controls. Psychological Medicine, 2019, 49, 1166-1173.	4.5	106
105	Transcriptomic organization of the human brain in post-traumatic stress disorder. Nature Neuroscience, 2021, 24, 24-33.	14.8	106
106	CNR1 Variation Modulates Risk for Drug and Alcohol Dependence. Biological Psychiatry, 2007, 62, 616-626.	1.3	105
107	Population genetics of a functional variant of the dopamine $\hat{l}^2$ -hydroxylase gene (DBH). American Journal of Medical Genetics Part A, 1997, 74, 374-379.	2.4	104
108	Genome-Wide Association Study of Alcohol Dependence Implicates KIAA0040 on Chromosome 1q. Neuropsychopharmacology, 2012, 37, 557-566.	5.4	104

#	Article	IF	Citations
109	Temporal lobe measurement in primary affective disorder by magnetic resonance imaging. Journal of Neuropsychiatry and Clinical Neurosciences, 1989, 1, 128-134.	1.8	103
110	Variant in RGS2 moderates posttraumatic stress symptoms following potentially traumatic event exposure. Journal of Anxiety Disorders, 2009, 23, 369-373.	3.2	103
111	Variation in Nicotinic Acetylcholine Receptor Genes is Associated with Multiple Substance Dependence Phenotypes. Neuropsychopharmacology, 2010, 35, 1921-1931.	5.4	103
112	X-Chromosome Markers and Manic-Depressive Illness. Archives of General Psychiatry, 1990, 47, 366.	12.3	102
113	ADH4 Gene Variation is Associated with Alcohol Dependence and Drug Dependence in European Americans: Results from HWD Tests and Case–Control Association Studies. Neuropsychopharmacology, 2006, 31, 1085-1095.	5.4	102
114	Linkage disequilibrium between an allele at the dopamine D4 receptor locus and Tourette syndrome, by the transmission-disequilibrium test. American Journal of Human Genetics, 1996, 59, 644-52.	6.2	102
115	Effects of the Brain-Derived Neurotrophic Growth Factor Val66Met Variation on Hippocampus Morphology in Bipolar Disorder. Neuropsychopharmacology, 2009, 34, 944-951.	5.4	101
116	<i>ALDH2</i> is associated to alcohol dependence and is the major genetic determinant of "daily maximum drinks―in a GWAS study of an isolated rural chinese sample. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 103-110.	1.7	101
117	Genetic Variants of Nogo-66 Receptor with Possible Association to Schizophrenia Block Myelin Inhibition of Axon Growth. Journal of Neuroscience, 2008, 28, 13161-13172.	3.6	98
118	Association of the OPRM1 Variant rs1799971 (A118G) with Non-Specific Liability to Substance Dependence in a Collaborative de novo Meta-Analysis of European-Ancestry Cohorts. Behavior Genetics, 2016, 46, 151-169.	2.1	98
119	Haplotypes at the OPRM1 locus are associated with susceptibility to substance dependence in European-Americans. American Journal of Medical Genetics Part A, 2003, 120B, 97-108.	2.4	97
120	Significant association of the neurexin-1 gene (NRXN1) with nicotine dependence in European- and African-American smokers. Human Molecular Genetics, 2008, 17, 1569-1577.	2.9	95
121	Haplotypic Variants in <i>DRD2</i> , <i>ANKK1</i> , <i>TTC12</i> , and <i>NCAM1</i> are Associated With Comorbid Alcohol and Drug Dependence. Alcoholism: Clinical and Experimental Research, 2008, 32, 2117-2127.	2.4	93
122	Prevalence of DSM-IV and DSM-5 alcohol, cocaine, opioid, and cannabis use disorders in a largely substance dependent sample. Drug and Alcohol Dependence, 2013, 127, 215-219.	3.2	93
123	Association of Gamma-Aminobutyric Acid A Receptor $\hat{l}\pm 2$ Gene (GABRA2) with Alcohol Use Disorder. Neuropsychopharmacology, 2014, 39, 907-918.	5.4	93
124	Linkage genome scan for loci predisposing to panic disorder or agoraphobia. American Journal of Medical Genetics Part A, 2001, 105, 548-557.	2.4	91
125	Genetic polymorphism at theCLOCK gene locus and major depression. American Journal of Medical Genetics Part A, 2000, 96, 418-421.	2.4	89
126	ADH4 gene variation is associated with alcohol and drug dependence: results from family controlled and population-structured association studies. Pharmacogenetics and Genomics, 2005, 15, 755-768.	1.5	87

#	Article	IF	CITATIONS
127	Genome-wide meta-analysis reveals common splice site acceptor variant in CHRNA4 associated with nicotine dependence. Translational Psychiatry, 2015, 5, e651-e651.	4.8	86
128	Suicide, impulsive aggression, and HTR1B genotype. Biological Psychiatry, 2001, 50, 62-65.	1.3	85
129	Components of Cross-Frequency Modulation in Health and Disease. Frontiers in Systems Neuroscience, 2011, 5, 59.	2.5	85
130	Genome-wide association study of therapeutic opioid dosing identifies a novel locus upstream of OPRM1. Molecular Psychiatry, 2017, 22, 346-352.	7.9	85
131	Markers in the 5′-Region of GABRG1 Associate to Alcohol Dependence and are in Linkage Disequilibrium with Markers in the Adjacent GABRA2 Gene. Neuropsychopharmacology, 2008, 33, 837-848.	<b>5.</b> 4	84
132	Genetic associations with suicide attempt severity and genetic overlap with major depression. Translational Psychiatry, 2019, 9, 22.	4.8	84
133	Arrayâ€Based Profiling of <scp>DNA</scp> Methylation Changes Associated with Alcohol Dependence. Alcoholism: Clinical and Experimental Research, 2013, 37, E108-15.	2.4	83
134	Genome-wide Association Study of Maximum Habitual Alcohol Intake in >140,000 U.S. European and African American Veterans Yields Novel Risk Loci. Biological Psychiatry, 2019, 86, 365-376.	1.3	82
135	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
136	Family-based association study of serotonin transporter promoter in suicidal adolescents: No association with suicidality but possible role in violence traits. American Journal of Medical Genetics Part A, 2001, 105, 239-245.	2.4	81
137	Association between a serotonin transporter promoter region polymorphism and mood response during tryptophan depletion. Molecular Psychiatry, 2002, 7, 213-216.	7.9	81
138	The Structure of Linkage Disequilibrium at the DBH Locus Strongly Influences the Magnitude of Association between Diallelic Markers and Plasma Dopamine β-Hydroxylase Activity. American Journal of Human Genetics, 2003, 72, 1389-1400.	6.2	81
139	Variation in <i>GABRA2</i> Predicts Drinking Behavior in Project MATCH Subjects. Alcoholism: Clinical and Experimental Research, 2007, 31, 1780-1787.	2.4	81
140	Genome-wide association study across European and African American ancestries identifies a SNP in DNMT3B contributing to nicotine dependence. Molecular Psychiatry, 2018, 23, 1911-1919.	7.9	80
141	Expanding the genetic architecture of nicotine dependence and its shared genetics with multiple traits. Nature Communications, 2020, $11$ , $5562$ .	12.8	80
142	Linkage mapping of serotonin transporter protein gene SLC6A4 on chromosome 17. Human Genetics, 1995, 95, 677-80.	3.8	79
143	D2 Dopamine Receptor Gene (DRD2) Allele and Haplotype Frequencies in Alcohol Dependent and Control Subjects No Association with Phenotype or Severity of Phenotype. Neuropsychopharmacology, 1999, 20, 640-649.	5.4	79
144	Results of a genomewide linkage scan: Support for chromosomes 9 and 11 loci increasing risk for cigarette smoking. American Journal of Medical Genetics Part A, 2004, 128B, 94-101.	2.4	79

#	Article	IF	CITATIONS
145	Genome-Wide Linkage Scan for Loci Predisposing to Social Phobia: Evidence for a Chromosome 16 Risk Locus. American Journal of Psychiatry, 2004, 161, 59-66.	7.2	78
146	<i>RGS2</i> and generalized anxiety disorder in an epidemiologic sample of hurricane-exposed adults. Depression and Anxiety, 2009, 26, 309-315.	4.1	78
147	Early-life stress, corpus callosum development, hippocampal volumetrics, and anxious behavior in male nonhuman primates. Psychiatry Research - Neuroimaging, 2011, 192, 37-44.	1.8	78
148	Genome-Wide Association Study of Nicotine Dependence in American Populations: Identification of Novel Risk Loci in Both African-Americans and European-Americans. Biological Psychiatry, 2015, 77, 493-503.	1.3	78
149	Genome Scan for Loci Predisposing to Anxiety Disorders Using a Novel Multivariate Approach: Strong Evidence for a Chromosome 4 Risk Locus. American Journal of Human Genetics, 2006, 78, 543-553.	6.2	76
150	A Common Genetic Variant in the Neurexin Superfamily Member CNTNAP2 Is Associated with Increased Risk for Selective Mutism and Social Anxiety-Related Traits. Biological Psychiatry, 2011, 69, 825-831.	1.3	76
151	The association of genetic variation in <i>CACNA1C</i> with structure and function of a frontotemporal system. Bipolar Disorders, 2011, 13, 696-700.	1.9	75
152	Gilles de la Tourette Syndrome Is Not Linked to D2-Dopamine Receptor. Archives of General Psychiatry, 1990, 47, 1073.	12.3	74
153	Genetic Risk Variants Associated With Comorbid Alcohol Dependence and Major Depression. JAMA Psychiatry, 2017, 74, 1234.	11.0	74
154	Evidence of causal effect of major depression on alcohol dependence: findings from the psychiatric genomics consortium. Psychological Medicine, 2019, 49, 1218-1226.	4.5	74
155	Polygenic Scores for Major Depressive Disorder and Risk of Alcohol Dependence. JAMA Psychiatry, 2017, 74, 1153.	11.0	73
156	Widespread signatures of positive selection in common risk alleles associated to autism spectrum disorder. PLoS Genetics, 2017, 13, e1006618.	3.5	73
157	Population Studies of Polymorphisms at Loci of Neuropsychiatric Interest (Tryptophan Hydroxylase) Tj ETQq1 1 0.	784314 rg 2.9	gBT /Overloo 72
158	Genomewide Linkage Scan for Nicotine Dependence: Identification of a Chromosome 5 Risk Locus. Biological Psychiatry, 2007, 61, 119-126.	1.3	72
159	Integrating GWASs and Human Protein Interaction Networks Identifies a Gene Subnetwork Underlying Alcohol Dependence. American Journal of Human Genetics, 2013, 93, 1027-1034.	6.2	72
160	Pervasive pleiotropy between psychiatric disorders and immune disorders revealed by integrative analysis of multiple GWAS. Human Genetics, 2015, 134, 1195-1209.	3.8	72
161	Association study of the CNR1 gene exon 3 alternative promoter region polymorphisms and substance dependence. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2006, 141B, 499-503.	1.7	71
162	Increased Genetic Vulnerability to Smoking at CHRNA5 in Early-Onset Smokers. Archives of General Psychiatry, 2012, 69, 854.	12.3	71

#	Article	IF	Citations
163	Improving genetic risk prediction by leveraging pleiotropy. Human Genetics, 2014, 133, 639-650.	3.8	71
164	Genome-wide analysis of insomnia disorder. Molecular Psychiatry, 2018, 23, 2238-2250.	7.9	71
165	Effects of Opioid Receptor Gene Variation on Targeted Nalmefene Treatment in Heavy Drinkers. Alcoholism: Clinical and Experimental Research, 2008, 32, 1159-1166.	2.4	68
166	DNA co-methylation modules in postmortem prefrontal cortex tissues of European Australians with alcohol use disorders. Scientific Reports, 2016, 6, 19430.	3.3	68
167	Apolipoprotein E ε4 Allele Is Unrelated to Cognitive or Functional Decline in Alzheimer's Disease: Retrospective and Prospective Analysis. Dementia and Geriatric Cognitive Disorders, 2006, 22, 73-82.	1.5	67
168	Genetic Relationship between Schizophrenia and Nicotine Dependence. Scientific Reports, 2016, 6, 25671.	3.3	67
169	Association of psychiatric and substance use disorder comorbidity with cocaine dependence severity and treatment utilization in cocaine-dependent individuals. Drug and Alcohol Dependence, 2009, 99, 193-203.	3.2	66
170	FKBP5 polymorphisms, childhood abuse, and PTSD symptoms: Results from the National Health and Resilience in Veterans Study. Psychoneuroendocrinology, 2016, 69, 98-105.	2.7	66
171	It Is Time to Take a Stand for Medical Research and Against Terrorism Targeting Medical Scientists. Biological Psychiatry, 2008, 63, 725-727.	1.3	65
172	The role of early life stress in development of the anterior limb of the internal capsule in nonhuman primates. Neuroscience Letters, 2010, 480, 93-96.	2.1	65
173	C957T polymorphism of the dopamine D2 receptor gene modulates the effect of nicotine on working memory performance and cortical processing efficiency. Psychopharmacology, 2006, 188, 530-540.	3.1	64
174	A CHRNA5 Smoking Risk Variant Decreases the Aversive Effects of Nicotine in Humans. Neuropsychopharmacology, 2015, 40, 2813-2821.	5.4	64
175	Genome-wide Association Study Identifies a Regulatory Variant of RGMA Associated With Opioid Dependence in European Americans. Biological Psychiatry, 2018, 84, 762-770.	1.3	64
176	Association Study of Alcoholism Subtypes with a Functional Promoter Polymorphism in the Serotonin Transporter Protein Gene. Alcoholism: Clinical and Experimental Research, 2002, 26, 1330-1335.	2.4	63
177	Childhood Adversity Increases Risk for Nicotine Dependence and Interacts with $\hat{l}\pm 5$ Nicotinic Acetylcholine Receptor Genotype Specifically in Males. Neuropsychopharmacology, 2012, 37, 669-676.	5.4	63
178	A polymorphism of the $\hat{l}^21$ -adrenergic receptor is associated with low extraversion. Biological Psychiatry, 2004, 56, 217-224.	1.3	62
179	Association of Markers in the 3′ Region of the GluR5 Kainate Receptor Subunit Gene to Alcohol Dependence. Alcoholism: Clinical and Experimental Research, 2009, 33, 925-930.	2.4	62
180	Rare Nonsynonymous Variants in Alpha-4 Nicotinic Acetylcholine Receptor Gene Protect Against Nicotine Dependence. Biological Psychiatry, 2011, 70, 528-536.	1.3	62

#	Article	IF	Citations
181	Serotonin transporter ⟨i⟩5â€HTTLPR⟨/i⟩ genotype moderates the effects of childhood adversity on posttraumatic stress disorder risk: A replication study. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2012, 159B, 644-652.	1.7	62
182	Meta-Analysis of 15 Genome-Wide Linkage Scans of Smoking Behavior. Biological Psychiatry, 2010, 67, 12-19.	1.3	61
183	<i>ADH1B</i> : From alcoholism, natural selection, and cancer to the human phenome. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2018, 177, 113-125.	1.7	61
184	Genome-wide meta-analysis of insomnia prioritizes genes associated with metabolic and psychiatric pathways. Nature Genetics, 2022, 54, 1125-1132.	21.4	61
185	Genetics of substance use disorders in the era of big data. Nature Reviews Genetics, 2021, 22, 712-729.	16.3	60
186	Genome-wide association study of smoking trajectory and meta-analysis of smoking status in 842,000 individuals. Nature Communications, 2020, 11, 5302.	12.8	59
187	Genotype-controlled analysis of plasma dopamine $\hat{l}^2$ -hydroxylase activity in psychotic unipolar major depression. Biological Psychiatry, 2002, 51, 358-364.	1.3	58
188	Dopamine $\hat{I}^2$ -Hydroxylase Gene (D $\hat{I}^2$ H) -1021C $\hat{a}^{\dagger}$ T Influences Self-Reported Paranoia during Cocaine Self-Administration. Biological Psychiatry, 2007, 61, 1310-1313.	1.3	58
189	Genetic risk prediction and neurobiological understanding of alcoholism. Translational Psychiatry, 2014, 4, e391-e391.	4.8	58
190	Genomewide Association Study for Maximum Number of Alcoholic Drinks in European Americans and African Americans. Alcoholism: Clinical and Experimental Research, 2015, 39, 1137-1147.	2.4	58
191	No association between D2 dopamine receptor (DRD2) alleles or haplotypes and cocaine dependence or severity of cocaine dependence in European- and African-Americans. Biological Psychiatry, 1999, 45, 340-345.	1.3	57
192	Measurement of admixture proportions and description of admixture structure in different U.S. populations. Human Mutation, 2009, 30, 1299-1309.	2.5	57
193	Hypermethylation of OPRM1 promoter region in European Americans with alcohol dependence. Journal of Human Genetics, 2012, 57, 670-675.	2.3	57
194	Typologies of drug dependence: comparative validity of a multivariate and four univariate models. Drug and Alcohol Dependence, 2004, 73, 289-300.	3.2	56
195	Population-specific effects of the Asn40Asp polymorphism at the $\hat{l}^{1}\!\!/\!\!4$ -opioid receptor gene (OPRM1) on HPA-axis activation. Pharmacogenetics and Genomics, 2007, 17, 1031-1038.	1.5	56
196	Interaction between Two Independent CNR1 Variants Increases Risk for Cocaine Dependence in European Americans: A Replication Study in Family-Based Sample and Population-Based Sample. Neuropsychopharmacology, 2009, 34, 1504-1513.	5.4	56
197	The genetics of alcohol dependence: Twin and SNPâ€based heritability, and genomeâ€wide association study based on AUDIT scores. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 739-748.	1.7	56
198	A chromosome 14 risk locus for simple phobia: results from a genomewide linkage scan. Molecular Psychiatry, 2003, 8, 71-82.	7.9	55

#	Article	IF	Citations
199	Rating the severity and character of transient cocaine-induced delusions and hallucinations with a new instrument, the Scale for Assessment of Positive Symptoms for Cocaine-Induced Psychosis (SAPS-CIP). Drug and Alcohol Dependence, 2005, 80, 23-33.	3.2	55
200	Support for association of RORA variant and post traumatic stress symptoms in a population-based study of hurricane exposed adults. Molecular Psychiatry, 2013, 18, 1148-1149.	7.9	55
201	Effect of Pharmacogenomic Testing for Drug-Gene Interactions on Medication Selection and Remission of Symptoms in Major Depressive Disorder. JAMA - Journal of the American Medical Association, 2022, 328, 151.	7.4	55
202	Role of Variation in the Serotonin Transporter Protein Gene (SLC6A4) in Trait Disturbances in the Ventral Anterior Cingulate in Bipolar Disorder. Neuropsychopharmacology, 2009, 34, 1301-1310.	5.4	54
203	A <i>CRHR1</i> haplotype moderates the effect of adverse childhood experiences on lifetime risk of major depressive episode in Africanâ€American women. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 960-968.	1.7	54
204	No association between ADCYAP1R1 and post-traumatic stress disorder in two independent samples. Molecular Psychiatry, 2012, 17, 239-241.	7.9	54
205	Novel gene identified in an exomeâ€wide association study of tanning dependence. Experimental Dermatology, 2014, 23, 757-759.	2.9	54
206	Rate of progression from first use to dependence on cocaine or opioids: A cross-substance examination of associated demographic, psychiatric, and childhood risk factors. Addictive Behaviors, 2014, 39, 473-479.	3.0	54
207	Population studies of polymorphisms of the serotonin transporter protein gene. American Journal of Medical Genetics Part A, 1999, 88, 61-6.	2.4	54
208	Assignment of the 5HT7 receptor gene (HTR7) to chromosome 10q and exclusion of genetic linkage with tourette syndrome. Genomics, 1995, 26, 207-209.	2.9	53
209	DRD2 Allele Frequencies and Linkage Disequilibria, Including the -141CIns/DelPromoter Polymorphism, in European-American, African-American, and Japanese Subjects. Genomics, 1998, 51, 21-26.	2.9	53
210	Influence of Vascular Endothelial Growth Factor Variation on Human Hippocampus Morphology. Biological Psychiatry, 2008, 64, 901-903.	1.3	52
211	Genetic associations with performance on a behavioral measure of distress intolerance. Journal of Psychiatric Research, 2012, 46, 87-94.	3.1	52
212	Genomewide association studies of suicide attempts in US soldiers. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2017, 174, 786-797.	1.7	52
213	Polymorphism of DCDC2 Reveals Differences in Cortical Morphology of Healthy Individuals—A Preliminary Voxel Based Morphometry Study. Brain Imaging and Behavior, 2008, 2, 21-26.	2.1	51
214	A pilot multivariate parallel ICA study to investigate differential linkage between neural networks and genetic profiles in schizophrenia. NeuroImage, 2010, 53, 1007-1015.	4.2	50
215	5-HTTLPR as a potential moderator of the effects of adverse childhood experiences on risk of antisocial personality disorder. Psychiatric Genetics, 2011, 21, 240-248.	1.1	50
216	The addiction risk factor: A unitary genetic vulnerability characterizes substance use disorders and their associations with common correlates. Neuropsychopharmacology, 2022, 47, 1739-1745.	5.4	50

#	Article	IF	CITATIONS
217	Genetic Associations of Brain Structural Networks in Schizophrenia: A Preliminary Study. Biological Psychiatry, 2010, 68, 657-666.	1.3	49
218	Genetic risk variants for social anxiety. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2017, 174, 120-131.	1.7	49
219	Review: DNA methylation and alcohol use disorders: Progress and challenges. American Journal on Addictions, 2017, 26, 502-515.	1.4	49
220	Association of Alcohol or Other Drug Dependence with Alleles of the mu Opioid Receptor Gene (OPRM1). Alcoholism: Clinical and Experimental Research, 1998, 22, 1359-1362.	2.4	48
221	Intronic variants in the dopa decarboxylase (DDC) gene are associated with smoking behavior in European-Americans and African-Americans. Human Molecular Genetics, 2006, 15, 2192-2199.	2.9	48
222	Manic depressive illness not linked to factor IX region in an independent series of pedigrees. Genomics, 1990, 8, 648-655.	2.9	47
223	Assignment of the norepinephrine transporter protein (NET1) locusto chromosome 16. Genomics, 1993, 18, 690-692.	2.9	47
224	<i>NPY</i> moderates the relation between hurricane exposure and generalized anxiety disorder in an epidemiologic sample of hurricane-exposed adults. Depression and Anxiety, 2010, 27, 270-275.	4.1	47
225	Genetics of Complex Traits in Psychiatry. Biological Psychiatry, 2015, 77, 36-42.	1.3	47
226	A genome-wide gene-by-trauma interaction study of alcohol misuse in two independent cohorts identifies PRKG1 as a risk locus. Molecular Psychiatry, 2018, 23, 154-160.	7.9	47
227	Genomic influences on self-reported childhood maltreatment. Translational Psychiatry, 2020, 10, 38.	4.8	47
228	Association study of cannabinoid receptor gene (CNR1) alleles and drug dependence. Molecular Psychiatry, 2001, 6, 501-502.	7.9	46
229	Comorbid Psychiatric Diagnoses and Their Association with Cocaine-Induced Psychosis in Cocaine-Dependent Subjects. American Journal on Addictions, 2007, 16, 343-351.	1.4	46
230	Genetic influences of cortical gray matter in language-related regions in healthy controls and schizophrenia. Schizophrenia Research, 2011, 129, 141-148.	2.0	46
231	Linkage Analysis Followed by Association Show NRG1 Associated with Cannabis Dependence in African Americans. Biological Psychiatry, 2012, 72, 637-644.	1.3	46
232	Oxytocin receptor gene polymorphisms, attachment, and PTSD: Results from the National Health and Resilience in Veterans Study. Journal of Psychiatric Research, 2017, 94, 139-147.	3.1	46
233	Progress in a genome scan for linkage in schizophrenia in a large Swedish kindred. American Journal of Medical Genetics Part A, 1994, 54, 51-58.	2.4	45
234	Case control and family-based studies of tryptophan hydroxylase gene A218C polymorphism and suicidality in adolescents. American Journal of Medical Genetics Part A, 2001, 105, 451-457.	2.4	45

#	Article	IF	Citations
235	Polymorphism of the 5-HT1B Receptor Gene (HTR1B) Strong Within-Locus Linkage Disequilibrium without Association to Antisocial Substance Dependence. Neuropsychopharmacology, 2002, 26, 115-122.	5.4	45
236	Variation at APOE and STH loci and Alzheimer's disease. Behavioral and Brain Functions, 2006, 2, 13.	3.3	45
237	Association of CRHR1 variants and posttraumatic stress symptoms in hurricane exposed adults. Journal of Anxiety Disorders, 2013, 27, 678-683.	3.2	44
238	Dopamine Beta-Hydroxylase (DBH) gene and schizophrenia phenotypic variability: A genetic association study. American Journal of Medical Genetics Part A, 2003, 117B, 33-38.	2.4	43
239	Genomewide Association Study of Alcohol Dependence Identifies Risk Loci Altering Ethanolâ€Response Behaviors in Model Organisms. Alcoholism: Clinical and Experimental Research, 2017, 41, 911-928.	2.4	43
240	Genetics of Alcoholism. Current Psychiatry Reports, 2019, 21, 26.	4.5	43
241	No association between an allele at the D2 dopamine receptor gene (DRD2) and alcoholism. JAMA - Journal of the American Medical Association, 1991, 266, 1801-7.	7.4	43
242	Further clarification of the contribution of the ADH1C gene to vulnerability of alcoholism and selected liver diseases. Human Genetics, 2012, 131, 1361-1374.	3.8	42
243	Exclusion of close linkage of Tourette's syndrome to D1 dopamine receptor. American Journal of Psychiatry, 1993, 150, 449-453.	7.2	41
244	Risk factors for cocaine-induced paranoia in cocaine-dependent sibling pairs. Drug and Alcohol Dependence, 2006, 84, 77-84.	3.2	41
245	Profiling of Childhood Adversity-Associated DNA Methylation Changes in Alcoholic Patients and Healthy Controls. PLoS ONE, 2013, 8, e65648.	2.5	41
246	Clinical features of methamphetamine-induced paranoia and preliminary genetic association with <i>DBH </i> -1021 Câ†'T in a Thai treatment cohort. Addiction, 2014, 109, 965-976.	3.3	41
247	D2 Dopamine Receptor Alleles Do Not Influence Severity of Tourette's Syndrome. Archives of Neurology, 1994, 51, 397.	4.5	40
248	Apolipoprotein E epsilon4 is associated with atrophy of the amygdala in Alzheimer's disease. Neurobiology of Aging, 2006, 27, 1416-1424.	3.1	40
249	Apolipoprotein E É>4 Allele Increases Risk for Psychotic Symptoms in Alzheimer's Disease. Neuropsychopharmacology, 2007, 32, 171-179.	5.4	40
250	Multiple ADH genes modulate risk for drug dependence in both African- and European-Americans. Human Molecular Genetics, 2007, 16, 380-390.	2.9	40
251	The Interplay Between Risky Sexual Behaviors and Alcohol Dependence: Genome-Wide Association and Neuroimaging Support for LHPP as a Risk Gene. Neuropsychopharmacology, 2017, 42, 598-605.	5.4	40
252	Association of Economic Status and Educational Attainment With Posttraumatic Stress Disorder. JAMA Network Open, 2019, 2, e193447.	5.9	40

#	Article	IF	CITATIONS
253	Neuropeptide Y-like immunoreactivity in schizophrenia. Schizophrenia Research, 1990, 3, 287-294.	2.0	39
254	<i>GABRG1</i> and <i>GABRA2</i> Variation Associated with Alcohol Dependence in African Americans. Alcoholism: Clinical and Experimental Research, 2012, 36, 588-593.	2.4	39
255	Identification of methylation quantitative trait loci (mQTLs) influencing promoter DNA methylation of alcohol dependence risk genes. Human Genetics, 2014, 133, 1093-1104.	3.8	39
256	Genome-Wide Association Study of Copy Number Variations (CNVs) with Opioid Dependence. Neuropsychopharmacology, 2015, 40, 1016-1026.	5.4	39
257	Genome-wide association study identifies a novel locus for cannabis dependence. Molecular Psychiatry, 2018, 23, 1293-1302.	7.9	39
258	Family-Based study of DRD2 alleles in alcohol and drug dependence. American Journal of Medical Genetics Part A, 2000, 96, 659-664.	2.4	38
259	Methylation in OTX2 and related genes, maltreatment, and depression in children. Neuropsychopharmacology, 2018, 43, 2204-2211.	5.4	38
260	Alleles of a Functional Serotonin Transporter Promoter Polymorphism Are Associated With Major Depression in Alcoholics. Alcoholism: Clinical and Experimental Research, 2003, 27, 1402-1408.	2.4	37
261	Early-life stress, corticotropin-releasing factor, and serotonin transporter gene: A pilot study. Psychoneuroendocrinology, 2011, 36, 289-293.	2.7	37
262	Improved methods to identify stable, highly heritable subtypes of opioid use and related behaviors. Addictive Behaviors, 2012, 37, 1138-1144.	3.0	37
263	Adverse Childhood Experiences, Epigenetic Measures, and Obesity in Youth. Journal of Pediatrics, 2018, 202, 150-156.e3.	1.8	37
264	The A1 allele at the D2 dopamine receptor gene and alcoholism. A reappraisal. JAMA - Journal of the American Medical Association, 1993, 269, 1673-7.	7.4	37
265	Association study of personality factors and the Asn40Asp polymorphism at the $\hat{l}^{1}\!\!/_{4}$ -opioid receptor gene (OPRM1). Psychiatric Genetics, 2004, 14, 89-92.	1.1	36
266	Examining the relation between the serotonin transporter 5-HTTPLR genotype x trauma exposure interaction on a contemporary phenotypic model of posttraumatic stress symptomatology: A pilot study. Journal of Affective Disorders, 2013, 148, 123-128.	4.1	36
267	Differentially co-expressed genes in postmortem prefrontal cortex of individuals with alcohol use disorders: influence on alcohol metabolism-related pathways. Human Genetics, 2014, 133, 1383-1394.	3.8	36
268	DSM-5 gambling disorder: Prevalence and characteristics in a substance use disorder sample Experimental and Clinical Psychopharmacology, 2014, 22, 50-56.	1.8	36
269	Dense Genomewide Linkage Scan for Alcohol Dependence in African Americans: Significant Linkage on Chromosome 10. Biological Psychiatry, 2009, 65, 111-115.	1.3	35
270	Allelic Variation of Calsyntenin 2 (CLSTN2) Modulates the Impact of Developmental Tobacco Smoke Exposure on Mnemonic Processing in Adolescents. Biological Psychiatry, 2009, 65, 671-679.	1.3	35

#	Article	IF	CITATIONS
271	The Dopamine Transporter Protein Gene (SLC6A3): Primary Linkage Mapping and Linkage Studies in Tourette Syndrome. Genomics, 1995, 30, 459-463.	2.9	34
272	GRIK1 Genotype moderates topiramate's effects on daily drinking level, expectations of alcohol's positive effects and desire to drink. International Journal of Neuropsychopharmacology, 2014, 17, 1549-1556.	2.1	34
273	Pharmacogenetics of naltrexone and disulfiram in alcohol dependent, dually diagnosed veterans. American Journal on Addictions, 2014, 23, 288-293.	1.4	34
274	Genomeâ€wide analyses of psychological resilience in U.S. Army soldiers. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, 310-319.	1.7	34
275	Sensory gating and psychosis vulnerability in cocaine-dependent individuals: preliminary data. Biological Psychiatry, 2002, 51, 683-686.	1.3	33
276	The validity of cocaine dependence subtypes. Addictive Behaviors, 2008, 33, 41-53.	3.0	33
277	Variation in Genes Encoding the Neuroactive Steroid Synthetic Enzymes 5α-Reductase Type 1 and 3α-Reductase Type 2 Is Associated With Alcohol Dependence. Alcoholism: Clinical and Experimental Research, 2011, 35, 946-952.	2.4	33
278	The Research Domain Criteria (RDoC) Project and Studies of Risk and Resilience in Maltreated Children. Journal of the American Academy of Child and Adolescent Psychiatry, 2015, 54, 617-625.	0.5	33
279	Phenome-Wide Association Study for Alcohol and Nicotine Risk Alleles in 26394 Women. Neuropsychopharmacology, 2016, 41, 2688-2696.	5.4	33
280	Ethanol Upregulates NMDA Receptor Subunit Gene Expression in Human Embryonic Stem Cell-Derived Cortical Neurons. PLoS ONE, 2015, 10, e0134907.	2.5	33
281	Variant detection at the $\hat{l}$ opioid receptor (OPRD1) locus and population genetics of a novel variant affecting protein sequence. Human Genetics, 2000, 107, 86-88.	3.8	32
282	The relationship between risk-taking propensity and the COMT Val158Met polymorphism among early adolescents as a function of sex. Journal of Psychiatric Research, 2012, 46, 940-945.	3.1	32
283	Distinct Loci in the <i>CHRNA5</i> / <i>CHRNA3</i> / <i>CHRNB4</i> Gene Cluster Are Associated With Onset of Regular Smoking. Genetic Epidemiology, 2013, 37, 846-859.	1.3	32
284	Alcohol Misuse and Coâ€Occurring Mental Disorders Among New Soldiers in the U.S. Army. Alcoholism: Clinical and Experimental Research, 2017, 41, 139-148.	2.4	32
285	Genomewide Association Study of Alcohol Dependence and Related Traits in a Thai Population. Alcoholism: Clinical and Experimental Research, 2018, 42, 861-868.	2.4	32
286	Molecular genetic overlap between posttraumatic stress disorder and sleep phenotypes. Sleep, 2020, 43, .	1.1	32
287	Dissecting the genetic association of C-reactive protein with PTSD, traumatic events, and social support. Neuropsychopharmacology, 2021, 46, 1071-1077.	5.4	32
288	Sex-stratified gene-by-environment genome-wide interaction study of trauma, posttraumatic-stress, and suicidality. Neurobiology of Stress, 2021, 14, 100309.	4.0	32

#	Article	IF	CITATIONS
289	Pro-Opiomelanocortin Gene Variation Related to Alcohol or Drug Dependence: Evidence and Replications Across Family- and Population-based Studies. Biological Psychiatry, 2009, 66, 128-136.	1.3	31
290	Connecting the dots, genome-wide association studies in substance use. Molecular Psychiatry, 2016, 21, 733-735.	7.9	31
291	A putative causal relationship between genetically determined female body shape and posttraumatic stress disorder. Genome Medicine, 2017, 9, 99.	8.2	31
292	The relationship between cannabis and schizophrenia: a genetically informed perspective. Addiction, 2021, 116, 3227-3234.	3.3	31
293	Absence of an Apolipoprotein E ϵ4 Allele Is Associated With Increased Parietal Regional Cerebral Blood Flow Asymmetry in Alzheimer Disease. Archives of Neurology, 1998, 55, 1460.	4.5	30
294	Association between polymorphisms in catecholâ€ <i>O</i> àâ€methyltransferase ( <i>COMT</i> ) and cocaineâ€induced paranoia in Europeanâ€American and Africanâ€American populations. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 651-660.	1.7	30
295	Validation of an Electronic Medical Record–Based Algorithm for Identifying Posttraumatic Stress Disorder in U.S. Veterans. Journal of Traumatic Stress, 2019, 32, 226-237.	1.8	30
296	Genomewide Study of Epigenetic Biomarkers of Opioid Dependence in European-American Women. Scientific Reports, 2019, 9, 4660.	3.3	30
297	Multivariate genome-wide analysis of education, socioeconomic status and brain phenome. Nature Human Behaviour, 2021, 5, 482-496.	12.0	30
298	DOPA decarboxylase gene is associated with nicotine dependence. Pharmacogenomics, 2006, 7, 1159-1166.	1.3	29
299	Meta-Analyses of Genome-Wide Association Data Hold New Promise for Addiction Genetics. Journal of Studies on Alcohol and Drugs, 2016, 77, 676-680.	1.0	29
300	No association between D3 dopamine receptor (DRD3) alleles and cocaine dependence. Addiction Biology, 1996, 1, 281-287.	2.6	28
301	Cognitive Flexibility is Associated with KIBRA Variant and Modulated by Recent Tobacco Use. Neuropsychopharmacology, 2009, 34, 2508-2516.	5.4	28
302	A Genomewide Linkage Scan of Cocaine Dependence and Major Depressive Episode in Two Populations. Neuropsychopharmacology, 2011, 36, 2422-2430.	5.4	28
303	M3: an improved SNP calling algorithm for Illumina BeadArray data. Bioinformatics, 2012, 28, 358-365.	4.1	28
304	Meta-analyses of genome-wide linkage scans of anxiety-related phenotypes. European Journal of Human Genetics, 2012, 20, 1078-1084.	2.8	28
305	The Joint Effects of <i><scp>ADH</scp>1B</i> Variants and Childhood Adversity on Alcohol Related Phenotypes in Africanâ€American and Europeanâ€American Women and Men. Alcoholism: Clinical and Experimental Research, 2014, 38, 2907-2914.	2.4	28
306	Childhood adversity moderates the effect of <i> ADH1B &lt; /i &gt; on risk for alcohol-related phenotypes in Jewish Israeli drinkers. Addiction Biology, 2015, 20, 205-214.</i>	2.6	28

#	Article	IF	CITATIONS
307	The role of genes involved in stress, neural plasticity, and brain circuitry in depressive phenotypes: Convergent findings in a mouse model of neglect. Behavioural Brain Research, 2016, 315, 71-74.	2.2	28
308	Cortical $\hat{l}^2$ -amyloid burden, gray matter, and memory in adults at varying APOE $\hat{l}\mu 4$ risk for Alzheimer's disease. Neurobiology of Aging, 2018, 61, 207-214.	3.1	28
309	Associations between moderate alcohol consumption, brain iron, and cognition in UK Biobank participants: Observational and mendelian randomization analyses. PLoS Medicine, 2022, 19, e1004039.	8.4	28
310	NOTCH4 gene haplotype is associated with schizophrenia in African Americans. Biological Psychiatry, 2004, 55, 112-117.	1.3	27
311	Human clock, PER1 and PER2 polymorphisms: lack of association with cocaine dependence susceptibility and cocaine-induced paranoia. Psychiatric Genetics, 2006, 16, 245-249.	1.1	27
312	The efficacies of clozapine and haloperidol in refractory schizophrenia are related to DTNBP1 variation. Pharmacogenetics and Genomics, 2009, 19, 437-446.	1.5	27
313	Functional impact of a single-nucleotide polymorphism in the OPRD1 promoter region. Journal of Human Genetics, 2010, 55, 278-284.	2.3	27
314	Association of <i>CHRNA4</i> polymorphisms with smoking behavior in two populations. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 421-429.	1.7	27
315	Comparing the utility of homogeneous subtypes of cocaine use and related behaviors with DSMâ€Ⅳ cocaine dependence as traits for genetic association analysis. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 148-156.	1.7	27
316	Exome sequencing and genome-wide copy number variant mapping reveal novel associations with sensorineural hereditary hearing loss. BMC Genomics, 2014, 15, 1155.	2.8	27
317	Polygenic risk for alcohol dependence associates with alcohol consumption, cognitive function and social deprivation in a populationâ€based cohort. Addiction Biology, 2016, 21, 469-480.	2.6	27
318	RDoC and translational perspectives on the genetics of traumaâ€related psychiatric disorders. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2016, 171, 81-91.	1.7	27
319	Sex Differences in Methamphetamine Use and Dependence in a Thai Treatment Center. Journal of Addiction Medicine, 2017, 11, 19-27.	2.6	27
320	Epigenomeâ€Wide <scp>DNA</scp> Methylation Association Analysis Identified Novel Loci in Peripheral Cells for Alcohol Consumption Among European American Male Veterans. Alcoholism: Clinical and Experimental Research, 2019, 43, 2111-2121.	2.4	27
321	Systematic review and metaâ€analysis of the moderating effect of rs1799971 in <i>OPRM1</i> , the muâ€opioid receptor gene, on response to naltrexone treatment of alcohol use disorder. Addiction, 2020, 115, 1426-1437.	3.3	27
322	A genome-wide association study of cocaine use disorder accounting for phenotypic heterogeneity and gene–environment interaction. Journal of Psychiatry and Neuroscience, 2020, 45, 34-44.	2.4	27
323	Variation in the Nicotinic Acetylcholine Receptor Gene Cluster CHRNA5–CHRNA3–CHRNB4 and Its Interaction with Recent Tobacco Use Influence Cognitive Flexibility. Neuropsychopharmacology, 2010, 35, 2211-2224.	5.4	26
324	Arguable Assumptions, Debatable Conclusions. Biological Psychiatry, 2010, 67, e19-e20.	1.3	26

#	Article	IF	CITATIONS
325	Inter-rater reliability and concurrent validity of DSM-IV opioid dependence in a Hmong isolate using the Thai version of the Semi-Structured Assessment for Drug Dependence and Alcoholism (SSADDA). Addictive Behaviors, 2011, 36, 156-160.	3.0	26
326	Empirically derived subtypes of opioid use and related behaviors. Addiction, 2011, 106, 1146-1154.	3.3	26
327	Genetic variation of ANK3 is associated with lower white matter structural integrity in bipolar disorder. Molecular Psychiatry, 2017, 22, 1225-1225.	7.9	26
328	Genomic regions controlling corticosterone levels in rats. Biological Psychiatry, 2004, 55, 634-641.	1.3	25
329	Multiple <i>OPR</i> genes influence personality traits in substance dependent and healthy subjects in two American populations. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2008, 147B, 1028-1039.	1.7	25
330	Association Between <i>COMT </i> , PTSD, and Increased Smoking Following Hurricane Exposure in an Epidemiologic Sample. Psychiatry (New York), 2009, 72, 360-369.	0.7	25
331	Variation in the gene encoding the serotonin transporter is associated with a measure of sociopathy in alcoholics. Addiction Biology, 2011, 16, 124-132.	2.6	25
332	Genetics of drug dependence. Dialogues in Clinical Neuroscience, 2010, 12, 77-84.	3.7	25
333	Ciliary neurotrophic factor null allele frequencies in schizophrenia, affective disorders, and Alzheimer's disease. American Journal of Medical Genetics Part A, 1997, 74, 497-500.	2.4	24
334	Characterization of a likelihood based method and effects of markers informativeness in evaluation of admixture and population group assignment. BMC Genetics, 2005, 6, 50.	2.7	24
335	Personality Traits of Agreeableness and Extraversion are Associated with ADH4 Variation. Biological Psychiatry, 2007, 61, 599-608.	1.3	24
336	Twenty-one-base-pair insertion polymorphism creates an enhancer element and potentiates SLC6A1 GABA transporter promoter activity. Pharmacogenetics and Genomics, 2009, 19, 53-65.	1.5	24
337	Adolescent cannabis use increases risk for cocaine-induced paranoia. Drug and Alcohol Dependence, 2010, 107, 196-201.	3.2	24
338	Biases and Errors on Allele Frequency Estimation and Disease Association Tests of Nextâ€Generation Sequencing of Pooled Samples. Genetic Epidemiology, 2012, 36, 549-560.	1.3	24
339	The effects of a <i>MAP2K5</i> microRNA target site SNP on risk for anxiety and depressive disorders. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 175-183.	1.7	24
340	Inter-observer reliability of DSM-5 substance use disorders. Drug and Alcohol Dependence, 2015, 153, 229-235.	3.2	24
341	Genomeâ€wide association metaâ€analysis of age at first cannabis use. Addiction, 2018, 113, 2073-2086.	3.3	24
342	AUDIT  and ICD codes as phenotypes for harmful alcohol use: association with <i>ADH1B</i> polymorphisms in two US populations. Addiction, 2018, 113, 2214-2224.	3.3	24

#	Article	IF	Citations
343	Characterizing the effect of background selection on the polygenicity of brain-related traits. Genomics, 2021, 113, 111-119.	2.9	24
344	Association of alcohol or other drug dependence with alleles of the mu opioid receptor gene (OPRM1). Alcoholism: Clinical and Experimental Research, 1998, 22, 1359-62.	2.4	24
345	CHRM2 variation predisposes to personality traits of agreeableness and conscientiousness. Human Molecular Genetics, 2007, 16, 1557-1568.	2.9	23
346	Mutation screen of the GAD2 gene and association study of alcoholism in three populations. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2007, 144B, 183-192.	1.7	23
347	Demographic changes and marker properties affect detection of human population differentiation. BMC Genetics, 2007, 8, 21.	2.7	23
348	Substance dependence low-density whole genome association study in two distinct American populations. Human Genetics, 2008, 123, 495-506.	3.8	23
349	Interleukin 1B gene ( <i>IL1B</i> ) variation and internalizing symptoms in maltreated preschoolers. Development and Psychopathology, 2014, 26, 1277-1287.	2.3	23
350	Effects of ANK3 variation on gray and white matter in bipolar disorder. Molecular Psychiatry, 2017, 22, 1345-1351.	7.9	23
351	Transient Cocaine-Associated Behavioral Symptoms Rated with a New Instrument, the Scale for Assessment of Positive Symptoms for Cocaine-Induced Psychosis (SAPS-CIP). American Journal on Addictions, 2009, 18, 339-345.	1.4	23
352	Schizophrenia: Instability in Norepinephrine, Serotonin, and $\hat{I}^3$ -Aminobutyric Acid Systems. International Review of Neurobiology, 1988, 29, 309-347.	2.0	22
353	Confirmation and fine mapping of the chromosome 1 alcohol dependence risk locus. Molecular Psychiatry, 2004, 9, 312-319.	7.9	22
354	Variation in <i>RGS2</i> is Associated with Suicidal Ideation in an Epidemiological Study of Adults Exposed to the 2004 Florida Hurricanes. Archives of Suicide Research, 2009, 13, 349-357.	2.3	22
355	Association of Variants in MANEA With Cocaine-Related Behaviors. Archives of General Psychiatry, 2009, 66, 267.	12.3	22
356	Posttreatment Effects of Topiramate Treatment for Heavy Drinking. Alcoholism: Clinical and Experimental Research, 2014, 38, 3017-3023.	2.4	22
357	Deep resequencing of 17 glutamate system genes identifies rare variants in <scp><i>Disc1</i></scp> and <scp><i>GRIN2B</i></scp> affecting risk of opioid dependence. Addiction Biology, 2014, 19, 955-964.	2.6	22
358	Dissecting ancestry genomic background in substance dependence genome-wide association studies. Pharmacogenomics, 2015, 16, 1487-1498.	1.3	22
359	Using phenotype risk scores to enhance gene discovery for generalized anxiety disorder and posttraumatic stress disorder. Molecular Psychiatry, 2022, 27, 2206-2215.	7.9	22
360	An in-frame deletion in the α2C adrenergic receptor is common in African–Americans. Molecular Psychiatry, 2001, 6, 168-172.	7.9	21

#	Article	IF	CITATIONS
361	Genetic influences of resting state fMRI activity in language-related brain regions in healthy controls and schizophrenia patients: a pilot study. Brain Imaging and Behavior, 2013, 7, 15-27.	2.1	21
362	COMT Val158Met modulates subjective responses to intravenous nicotine and cognitive performance in abstinent smokers. Pharmacogenomics Journal, 2013, 13, 490-497.	2.0	21
363	Genetic association of impulsivity in young adults: a multivariate study. Translational Psychiatry, 2014, 4, e451-e451.	4.8	21
364	Selfâ€efficacy mediates the effects of topiramate and <scp><i>GRIK1</i></scp> genotype on drinking. Addiction Biology, 2016, 21, 450-459.	2.6	21
365	Genomeâ€wide association study of body mass index in subjects with alcohol dependence. Addiction Biology, 2017, 22, 535-549.	2.6	21
366	Cross-Phenotype Polygenic Risk Score Analysis of Persistent Post-Concussive Symptoms in U.S. Army Soldiers with Deployment-Acquired Traumatic Brain Injury. Journal of Neurotrauma, 2017, 34, 781-789.	3.4	21
367	Genetically determined schizophrenia is not associated with impaired glucose homeostasis. Schizophrenia Research, 2018, 195, 286-289.	2.0	21
368	Genome-wide association study of alcohol dependence in male Han Chinese and cross-ethnic polygenic risk score comparison. Translational Psychiatry, 2019, 9, 249.	4.8	21
369	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. World Journal of Biological Psychiatry, 2019, 20, 289-300.	2.6	21
370	Enhancing Discovery of Genetic Variants for Posttraumatic Stress Disorder Through Integration of Quantitative Phenotypes and Trauma Exposure Information. Biological Psychiatry, 2022, 91, 626-636.	1.3	21
371	Human GABAB receptor 1 gene: Eight novel sequence variants. Human Mutation, 2001, 17, 349-350.	2.5	20
372	Subtypes of major depression in substance dependence. Addiction, 2009, 104, 1700-1709.	3.3	20
373	SLC6A4 polymorphism, population genetics, and psychiatric traits. Human Genetics, 2014, 133, 459-461.	3.8	20
374	Schizophrenia and substance use comorbidity: a genome-wide perspective. Genome Medicine, 2017, 9, 25.	8.2	20
375	Apolipoprotein E gene polymorphism, trauma burden, and posttraumatic stress symptoms in U.S. military veterans: Results from the National Health and Resilience in Veterans Study. Depression and Anxiety, 2018, 35, 168-177.	4.1	20
376	Translational studies support a role for serotonin 2B receptor (HTR2B) gene in aggression-related cannabis response. Molecular Psychiatry, 2018, 23, 2277-2286.	7.9	20
377	Multi-environment gene interactions linked to the interplay between polysubstance dependence and suicidality. Translational Psychiatry, 2021, 11, 34.	4.8	20
378	Setting Priorities for Genomic Research. Science, 2004, 304, 1445-1447.	12.6	19

#	Article	IF	CITATIONS
379	Association of COL25A1 with Comorbid Antisocial Personality Disorder and Substance Dependence. Biological Psychiatry, 2012, 71, 733-740.	1.3	19
380	Indoor Tanning and Tanning Dependence in Young People After a Diagnosis of Basal Cell Carcinoma. JAMA Dermatology, 2013, 149, 1110.	4.1	19
381	Salivary microRNAs identified by small RNA sequencing and machine learning as potential biomarkers of alcohol dependence. Epigenomics, 2019, 11, 739-749.	2.1	19
382	Post-GWAS analysis of six substance use traits improves the identification and functional interpretation of genetic risk loci. Drug and Alcohol Dependence, 2020, 206, 107703.	3.2	19
383	Attachment Style Moderates Polygenic Risk for Posttraumatic Stress in United States Military Veterans: Results From the National Health and Resilience in Veterans Study. Biological Psychiatry, 2021, 89, 878-887.	1.3	19
384	An Atlas of Genetic Correlations and Genetically Informed Associations Linking Psychiatric and Immune-Related Phenotypes. JAMA Psychiatry, 2022, 79, 667.	11.0	19
385	Analysis of variations in the tryptophan hydroxylase-2 (TPH2) gene in cocaine dependence. Addiction Biology, 2006, 11, 76-83.	2.6	18
386	<i>ADH7</i> variation modulates extraversion and conscientiousness in substanceâ€dependent subjects. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2008, 147B, 179-186.	1.7	18
387	Interpopulation linkage disequilibrium patterns of GABRA2 and GABRG1 genes at the GABA cluster locus on human chromosome 4. Genomics, 2008, 91, 61-69.	2.9	18
388	Rare Human Nicotinic Acetylcholine Receptor $\hat{l}\pm 4$ Subunit (CHRNA4) Variants Affect Expression and Function of High-Affinity Nicotinic Acetylcholine Receptors. Journal of Pharmacology and Experimental Therapeutics, 2014, 348, 410-420.	2.5	18
389	Multivariate Imaging Genetics Study of MRI Gray Matter Volume and SNPs Reveals Biological Pathways Correlated with Brain Structural Differences in Attention Deficit Hyperactivity Disorder. Frontiers in Psychiatry, 2016, 7, 128.	2.6	18
390	A preliminary examination of the relationship between the 5-HTTLPR and childhood emotional abuse on depressive symptoms in 10-12-year-old youth Psychological Trauma: Theory, Research, Practice, and Policy, 2014, 6, 1-7.	2.1	17
391	Ancestryâ€specific and sexâ€specific risk alleles identified in a genomeâ€wide geneâ€byâ€alcohol dependence interaction study of risky sexual behaviors. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2017, 174, 846-853.	1.7	17
392	Genetic factor common to schizophrenia and HIV infection is associated with risky sexual behavior: antagonistic vs. synergistic pleiotropic SNPs enriched for distinctly different biological functions. Human Genetics, 2017, 136, 75-83.	3.8	17
393	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. Depression and Anxiety, 2019, 36, 834-845.	4.1	17
394	Genome-Wide Meta-Analyses of FTND and TTFC Phenotypes. Nicotine and Tobacco Research, 2020, 22, 900-909.	2.6	17
395	Genome-Wide Association Study of Opioid Cessation. Journal of Clinical Medicine, 2020, 9, 180.	2.4	17
396	Variants in the ?2A AR adrenergic receptor gene in psychiatric patients. , 1998, 81, 405-410.		16

#	Article	IF	CITATIONS
397	Functional variants at CYP2A6: New genotyping methods, population genetics, and relevance to studies of tobacco dependence. American Journal of Medical Genetics Part A, 2000, 96, 638-645.	2.4	16
398	Southeast Asian origins of five Hill Tribe populations and correlation of genetic to linguistic relationships inferred with genomeâ€wide SNP data. American Journal of Physical Anthropology, 2011, 144, 300-308.	2.1	16
399	Ordered subset linkage analysis based on admixture proportion identifies new linkage evidence for alcohol dependence in African-Americans. Human Genetics, 2013, 132, 397-403.	3.8	16
400	Alcohol-Metabolizing Genes and Alcohol Phenotypes in an Israeli Household Sample. Alcoholism: Clinical and Experimental Research, 2013, 37, 1872-1881.	2.4	16
401	FKBP5 variation is associated with the acute and chronic effects of nicotine. Pharmacogenomics Journal, 2015, 15, 340-346.	2.0	16
402	Pstl RFLP at the SERT locus. Human Molecular Genetics, 1994, 3, 383-383.	2.9	15
403	Exploring the genetic architecture of alcohol dependence in African-Americans via analysis of a genomewide set of common variants. Human Genetics, 2014, 133, 617-624.	3.8	15
404	Ancestry informative markers for distinguishing between Thai populations based on genome-wide association datasets. Legal Medicine, 2015, 17, 245-250.	1.3	15
405	Alcohol and nicotine codependence-associated DNA methylation changes in promoter regions of addiction-related genes. Scientific Reports, 2017, 7, 41816.	3.3	15
406	Validating Harmful Alcohol Use as a Phenotype for Genetic Discovery Using Phosphatidylethanol and a Polymorphism in <i><scp>ADH</scp>1B</i> . Alcoholism: Clinical and Experimental Research, 2017, 41, 998-1003.	2.4	15
407	Multivariate Pattern Analysis of Genotype–Phenotype Relationships in Schizophrenia. Schizophrenia Bulletin, 2018, 44, 1045-1052.	4.3	15
408	Increased Risk of Multiple Outpatient Surgeries in African-American Carriers of Transthyretin Val122lle Mutation Is Modulated by Non-Coding Variants. Journal of Clinical Medicine, 2019, 8, 269.	2.4	15
409	Detection of marker associations with a dominant disease gene in genetically complex and heterogeneous diseases. American Journal of Human Genetics, 1989, 45, 578-85.	6.2	15
410	Genetically regulated multi-omics study for symptom clusters of posttraumatic stress disorder highlights pleiotropy with hematologic and cardio-metabolic traits. Molecular Psychiatry, 2022, 27, 1394-1404.	7.9	15
411	Association of Kidney Comorbidities and Acute Kidney Failure With Unfavorable Outcomes After COVID-19 in Individuals With the Sickle Cell Trait. JAMA Internal Medicine, 0, , .	5.1	15
412	Variant detection at the ? opioid receptor (OPRD1) locus and population genetics of a novel variant affecting protein sequence. Human Genetics, 2000, 107, 86-88.	3.8	14
413	Association study of DTNBP1 with schizophrenia in a US sample. Psychiatric Genetics, 2009, 19, 292-304.	1.1	14
414	Autosomal linkage scan for loci predisposing to comorbid dependence on multiple substances. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2012, 159B, 361-369.	1.7	14

#	Article	IF	CITATIONS
415	The α-endomannosidase gene (MANEA) is associated with panic disorder and social anxiety disorder. Translational Psychiatry, 2014, 4, e353-e353.	4.8	14
416	Nf1 Regulates Alcohol Dependence-Associated Excessive Drinking and Gamma-Aminobutyric Acid Release in the Central Amygdala in Mice and Is Associated with Alcohol Dependence in Humans. Biological Psychiatry, 2015, 77, 870-879.	1.3	14
417	Trauma exposure interacts with the genetic risk of bipolar disorder in alcohol misuse of <scp>US</scp> soldiers. Acta Psychiatrica Scandinavica, 2018, 137, 148-156.	4.5	14
418	Genome-wide association study identifies glutamate ionotropic receptor GRIA4 as a risk gene for comorbid nicotine dependence and major depression. Translational Psychiatry, 2018, 8, 208.	4.8	14
419	Risk Locus Identification Ties Alcohol Withdrawal Symptoms to <i><scp>SORCS</scp>2</i> . Alcoholism: Clinical and Experimental Research, 2018, 42, 2337-2348.	2.4	14
420	Alcohol-responsive genes identified in human iPSC-derived neural cultures. Translational Psychiatry, 2019, 9, 96.	4.8	14
421	Deciphering the Biological Mechanisms Underlying the Genome-Wide Associations between Computerized Device Use and Psychiatric Disorders. Journal of Clinical Medicine, 2019, 8, 2040.	2.4	14
422	CALCYON gene variation, schizophrenia, and cocaine dependence. American Journal of Medical Genetics Part A, 2004, 125B, 25-30.	2.4	13
423	BDNF Variants, Premorbid Educational Attainment, and Disease Characteristics in Alzheimer's Disease: An Exploratory Study. Journal of Alzheimer's Disease, 2009, 17, 887-898.	2.6	13
424	Variation in regulator of G-protein signaling 17 gene (RGS17) is associated with multiple substance dependence diagnoses. Behavioral and Brain Functions, 2012, 8, 23.	3.3	13
425	Patterns and Correlates of Prescription Opioid Receipt Among US Veterans: A National, 18-Year Observational Cohort Study. AIDS and Behavior, 2019, 23, 3340-3349.	2.7	13
426	Retrospective Association Analysis of Longitudinal Binary Traits Identifies Important Loci and Pathways in Cocaine Use. Genetics, 2019, 213, 1225-1236.	2.9	13
427	Genomeâ€wide association study of shared liability to anxiety disorders in Army STARRS. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2020, 183, 197-207.	1.7	13
428	Identification of POMC Exonic Variants Associated with Substance Dependence and Body Mass Index. PLoS ONE, 2012, 7, e45300.	2.5	13
429	Differential Expression of miR-130a in Postmortem Prefrontal Cortex of Subjects with Alcohol Use Disorders. Journal of Addiction Research & Therapy, 2013, 04, .	0.2	13
430	Integrating human brain proteomes with genome-wide association data implicates novel proteins in post-traumatic stress disorder. Molecular Psychiatry, 2022, 27, 3075-3084.	7.9	13
431	<i>ADH1A</i> variation predisposes to personality traits and substance dependence. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 376-386.	1.7	12
432	Identification of population substructure among Jews using STR markers and dependence on reference populations included. BMC Genetics, 2010, 11, 48.	2.7	12

#	Article	IF	CITATIONS
433	Agitated depression in substance dependence. Drug and Alcohol Dependence, 2011, 116, 163-169.	3.2	12
434	Variation inâ€, <i>OPRM1</i> à€,and Risk of Suicidal Behavior in Drugâ€Dependent Individuals. American Journal on Addictions, 2012, 21, 5-10.	1.4	12
435	Which alcohol use disorder criteria contribute to the association of ADH1B with alcohol dependence?. Addiction Biology, 2016, 21, 924-938.	2.6	12
436	<i>CHRNA4</i> and <i>ANKK1</i> Polymorphisms Influence Smoking-Induced Nicotinic Acetylcholine Receptor Upregulation. Nicotine and Tobacco Research, 2016, 18, 1845-1852.	2.6	12
437	Phenomic Impact of Genetically-Determined Euthyroid Function and Molecular Differences between Thyroid Disorders. Journal of Clinical Medicine, 2018, 7, 296.	2.4	12
438	Integration of evidence across human and model organism studies: A meeting report. Genes, Brain and Behavior, 2021, 20, e12738.	2.2	12
439	Drinking and smoking polygenic risk is associated with childhood and early-adulthood psychiatric and behavioral traits independently of substance use and psychiatric genetic risk. Translational Psychiatry, 2021, 11, 586.	4.8	12
440	Mutational analysis of candidate genes in psychiatric disorders. American Journal of Medical Genetics Part A, 1993, 48, 184-191.	2.4	11
441	A revised allele frequency estimate and haplotype analysis of the DBH deficiency mutation IVS1+2T → C African- and European-Americans. , 2003, 123A, 190-192.	C in	11
442	ACSL6 Is Associated with the Number of Cigarettes Smoked and Its Expression Is Altered by Chronic Nicotine Exposure. PLoS ONE, 2011, 6, e28790.	2.5	11
443	GABRA2 and KIBRA genotypes predict early relapse to substance use. Drug and Alcohol Dependence, 2012, 123, 154-159.	3.2	11
444	Genetic risk variants for social anxiety. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2017, 174, 470-482.	1.7	11
445	Using DNA methylation to validate an electronic medical record phenotype for smoking. Addiction Biology, 2019, 24, 1056-1065.	2.6	11
446	Differentiating Types of Self-Reported Alcohol Abstinence. AIDS and Behavior, 2020, 24, 655-665.	2.7	11
447	The impact of removing former drinkers from genomeâ€wide association studies of AUDIT . Addiction, 2021, 116, 3044-3054.	3.3	11
448	Psychological trauma and the genetic overlap between posttraumatic stress disorder and major depressive disorder. Psychological Medicine, 2022, 52, 3975-3984.	4.5	11
449	Evidence of Polygenic Adaptation in the Systems Genetics of Anthropometric Traits. PLoS ONE, 2016, 11, e0160654.	2.5	11
450	ANKRD7 and CYTL1 are novel risk genes for alcohol drinking behavior. Chinese Medical Journal, 2012, 125, 1127-34.	2.3	11

#	Article	IF	CITATIONS
451	<i>GABRA2</i> Genotype, Impulsivity, and Body Mass. American Journal on Addictions, 2012, 21, 404-410.	1.4	10
452	A protocadherin gene cluster regulatory variant is associated with nicotine withdrawal and the urge to smoke. Molecular Psychiatry, 2017, 22, 242-249.	7.9	10
453	Sexâ€specific linkage scans in opioid dependence. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2017, 174, 261-268.	1.7	10
454	Genomeâ€wide scan identifies opioid overdose risk locus close to MCOLN1. Addiction Biology, 2020, 25, e12811.	2.6	10
455	Genetic Factors in Alcoholism: Evidence and Implications. Handbook of Experimental Pharmacology, 1995, , 297-313.	1.8	10
456	Genome-wide meta-analysis of alcohol use disorder in East Asians. Neuropsychopharmacology, 2022, 47, 1791-1797.	5.4	10
457	Epigenome-wide association study of posttraumatic stress disorder identifies novel loci in U.S. military veterans. Translational Psychiatry, 2022, 12, 65.	4.8	10
458	Sequence tagged site (STS) Taql RFLP at dopamine $\hat{l}^2$ -hydroxylase (DBH). Nucleic Acids Research, 1991, 19, 1957-1957.	14.5	9
459	Recessive genetic mode of an ADH4 variant in substance dependence in African-Americans: A model of utility of the HWD test. Behavioral and Brain Functions, 2008, 4, 42.	3.3	9
460	Sources of Unreliability in the Diagnosis of Substance Dependence. Journal of Studies on Alcohol and Drugs, 2009, 70, 475-481.	1.0	9
461	Confirmation and Generalization of an Alcohol-Dependence Locus on Chromosome 10q. Neuropsychopharmacology, 2010, 35, 1325-1332.	5.4	9
462	Alcohol Consumption Mediates the Relationship Between <i>ADH1B</i> and DSM-IV Alcohol Use Disorder and Criteria. Journal of Studies on Alcohol and Drugs, 2014, 75, 635-642.	1.0	9
463	Phenome-wide association study for CYP2A6 alleles: rs113288603 is associated with hearing loss symptoms in elderly smokers. Scientific Reports, 2017, 7, 1034.	3.3	9
464	<i><i><scp>GRIK</scp>1</i> and <i><scp>GABRA</scp>2</i> Variants Have Distinct Effects on the Doseâ€Related Subjective Response to Intravenous Alcohol in Healthy Social Drinkers. Alcoholism: Clinical and Experimental Research, 2017, 41, 2025-2032.</i>	2.4	9
465	Genetic and psychosocial predictors of alcohol use trajectories among disaster-exposed adolescents. American Journal on Addictions, 2017, 26, 623-631.	1.4	9
466	S100A10 identified in a genome-wide gene $\tilde{A}-$ cannabis dependence interaction analysis of risky sexual behaviours. Journal of Psychiatry and Neuroscience, 2017, 42, 252-261.	2.4	9
467	Genomewide Gene-by-Sex Interaction Scans Identify ADGRV1 for Sex Differences in Opioid Dependent African Americans. Scientific Reports, 2019, 9, 18070.	3.3	9
468	Population studies of polymorphisms of the serotonin transporter protein gene. American Journal of Medical Genetics Part A, 1999, 88, 61-66.	2.4	9

#	Article	IF	CITATIONS
469	Psychosocial moderators of polygenic risk for suicidal ideation: Results from a 7-year population-based, prospective cohort study of U.S. veterans. Molecular Psychiatry, 2022, 27, 1068-1074.	7.9	9
470	Taql RFLP at norepinephrine transporter protein (NET) locus. Human Molecular Genetics, 1993, 2, 820-820.	2.9	8
471	Population admixture modulates risk for alcohol dependence. Human Genetics, 2009, 125, 605-613.	3.8	8
472	Variation in <i>NGFB</i> is associated with primary affective disorders in women. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 401-412.	1.7	8
473	Polygenic risk for autism spectrum disorder associates with anger recognition in a neurodevelopment-focused phenome-wide scan of unaffected youths from a population-based cohort. PLoS Genetics, 2020, 16, e1009036.	3.5	8
474	Dissecting the heterogeneity of posttraumatic stress disorder: differences in polygenic risk, stress exposures, and course of PTSD subtypes. Psychological Medicine, 2021, , 1-9.	4.5	8
475	Expert systems and diagnostic monitors in psychiatry. Medical Informatics = Medecine Et Informatique, 1986, 11, 23-28.	0.8	7
476	A detailed physical map of the 6p reading disability locus, including new markers and confirmation of recombination suppression. Human Genetics, 2002, 111, 339-349.	3.8	7
477	Incentive program decreases no-shows in nontreatment substance abuse research Experimental and Clinical Psychopharmacology, 2005, 13, 376-380.	1.8	7
478	Quantitative Trait Locus Analysis Identifies Rat Genomic Regions Related to Amphetamine-Induced Locomotion and Gαi3 Levels in Nucleus Accumbens. Neuropsychopharmacology, 2008, 33, 2735-2746.	5.4	7
479	A Complex Interplay between Personality Domains, Marital Status and a Variant in CHRNA5 on the Risks of Cocaine, Nicotine Dependences and Cocaine-Induced Paranoia. PLoS ONE, 2013, 8, e49368.	2.5	7
480	GRIK1 genotype and daily expectations of alcohol's positive effects moderate the reduction of heavy drinking by topiramate Experimental and Clinical Psychopharmacology, 2014, 22, 494-501.	1.8	7
481	Association Study of Genotype by Depressive Response during Acute Tryptophan Depletion in Subjects Recovered from Major Depression. Molecular Neuropsychiatry, 2015, 1, 165-174.	2.9	7
482	Eye color: A potential indicator of alcohol dependence risk in European Americans. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 347-353.	1.7	7
483	Epigenomic Profiles of African-American Transthyretin Val122lle Carriers Reveals Putatively Dysregulated Amyloid Mechanisms. Circulation Genomic and Precision Medicine, 2021, 14, e003011.	3.6	7
484	Investigation of convergent and divergent genetic influences underlying schizophrenia and alcohol use disorder. Psychological Medicine, 2023, 53, 1196-1204.	4.5	7
485	Combined analysis of the moderating effect of a GRIK1 polymorphism on the effects of topiramate for treating alcohol use disorder. Drug and Alcohol Dependence, 2021, 225, 108762.	3.2	7
486	Critical evaluation of transcription factor Atf2 as a candidate modulator of alcohol preference in mouse and human populations. Genetics and Molecular Research, 2013, 12, 5992-6005.	0.2	7

#	Article	IF	CITATIONS
487	Attachment style moderates polygenic risk for incident posttraumatic stress in U.S. military veterans: A 7-year, nationally representative, prospective cohort study. Biological Psychiatry, 2021, , .	1.3	7
488	Characteristics and Course of Dependence in Cocaine-Dependent Individuals Who Never Used Alcohol or Marijuana or Used Cocaine First. Journal of Studies on Alcohol and Drugs, 2014, 75, 423-427.	1.0	6
489	Topiramate's reduction of body mass index in heavy drinkers: Lack of moderation by a GRIK1 polymorphism Experimental and Clinical Psychopharmacology, 2014, 22, 419-423.	1.8	6
490	Biochemical, demographic, and self-reported tobacco-related predictors of the acute heart rate response to nicotine in smokers. Pharmacology Biochemistry and Behavior, 2018, 173, 36-43.	2.9	6
491	Investigating Causality Between Blood Metabolites and Emotional and Behavioral Responses to Traumatic Stress: a Mendelian Randomization Study. Molecular Neurobiology, 2020, 57, 1542-1552.	4.0	6
492	Genome-wide association study of phenotypes measuring progression from first cocaine or opioid use to dependence reveals novel risk genes. Exploration of Medicine, 2021, 2, 60-73.	1.5	6
493	Ancestry may confound genetic machine learning: Candidate-gene prediction of opioid use disorder as an example. Drug and Alcohol Dependence, 2021, 229, 109115.	3.2	6
494	Linkage map of eight human chromosome 11q markers, including DRD2, spanning 60 cM. Cytogenetic and Genome Research, 1992, 60, 26-28.	1.1	5
495	Genetic Basis of Social Anxiety Disorder. , 2010, , 313-322.		5
496	Local adaptation in European populations affected the genetics of psychiatric disorders and behavioral traits. Genome Medicine, 2018, 10, 24.	8.2	5
497	Genomics of posttraumatic stress disorder in veterans: Methods and rationale for <scp>V</scp> eterans <scp>A</scp> ffairs <scp>C</scp> ooperative <scp>S</scp> tudy #575B. International Journal of Methods in Psychiatric Research, 2019, 28, e1767.	2.1	5
498	Study design and implementation of the PRecision Medicine In MEntal health Care (PRIME Care) Trial. Contemporary Clinical Trials, 2021, 101, 106247.	1.8	5
499	Polygenic risk for traumatic loss-related PTSD in US military veterans: Protective effect of secure attachment style. World Journal of Biological Psychiatry, 2021, 22, 792-799.	2.6	5
500	Polygenic risk for major depression is associated with lifetime suicide attempt in <scp>US</scp> soldiers independent of personal and parental history of major depression. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2021, 186, 469-475.	1.7	5
501	Alcoholism and the D2 Dopamine Receptor Gene-Reply. JAMA - Journal of the American Medical Association, 1993, 270, 1547.	7.4	4
502	Sequence variation and linkage disequilibrium in the GABA transporter-1 gene (SLC6A1) in five populations: implications for pharmacogenetic research. BMC Genetics, 2007, 8, 71.	2.7	4
503	Developmental Perspective on the Role of Genes in Smoking Risk. Biological Psychiatry, 2011, 69, 616-617.	1.3	4
504	Evaluating the role of a galanin enhancer genotype on a range of metabolic, depressive and addictive phenotypes. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 654-664.	1.7	4

#	Article	IF	CITATIONS
505	<i>CHRNA5/A3/B4</i> Variant rs3743078 and Nicotine-Related Phenotypes: Indirect Effects Through Nicotine Craving. Journal of Studies on Alcohol and Drugs, 2016, 77, 227-237.	1.0	4
506	Genomeâ€wide association study of cognitive flexibility assessed by the Wisconsin Card Sorting Test. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2018, 177, 511-519.	1.7	4
507	Tubulin Polymerization Promoting Protein (TPPP) gene methylation and corpus callosum measures in maltreated children. Psychiatry Research - Neuroimaging, 2020, 298, 111058.	1.8	4
508	GABAergic polygenic risk for cocaine use disorder is negatively correlated with precuneus activity during cognitive control in African American individuals. Addictive Behaviors, 2021, 114, 106695.	3.0	4
509	Pleiotropic effects of telomere length loci with brain morphology and brain tissue expression. Human Molecular Genetics, 2021, 30, 1360-1370.	2.9	4
510	Genome-wide association study of stimulant dependence. Translational Psychiatry, 2021, 11, 363.	4.8	4
511	Reliability and validity of DSM-IV and DSM-5 methamphetamine use disorder diagnoses using the Chinese Version of the Semi-Structured Assessment for Drug Dependence and Alcoholism (SSADDA). Drug and Alcohol Dependence, 2021, 229, 109047.	3.2	4
512	Postâ€treatment effects of topiramate on alcoholâ€related outcomes: A combined analysis of two placeboâ€controlled trials. Addiction Biology, 2022, 27, e13130.	2.6	4
513	Alcohol withdrawal in pastâ€year drinkers with unhealthy alcohol use: Prevalence, characteristics, and correlates in a national epidemiologic survey. Alcoholism: Clinical and Experimental Research, 2022, 46, 422-433.	2.4	4
514	Response to Dr. Kopke's comments on haplotypes at the OPRM1 locus. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2005, 135B, 102-102.	1.7	3
515	Novel QTL at chromosome 6p22 for alcohol consumption: Implications for the genetic liability of alcohol use disorders. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 294-302.	1.7	3
516	Ttn as a likely causal gene for QTL of alcohol preference on mouse chromosome 2. BMC Bioinformatics, 2014, 15, P12.	2.6	3
517	A regulatory variant of CHRM3 is associated with cannabis-induced hallucinations in European Americans. Translational Psychiatry, 2019, 9, 309.	4.8	3
518	Diagnostic Reliability and Validity of the Semi-Structured Assessment for Drug Dependence and Alcoholism (SSADDA) Chinese Version. Complex Psychiatry, 2020, 6, 62-67.	0.9	3
519	Disentangling sex differences in the shared genetic architecture of posttraumatic stress disorder, traumatic experiences, and social support with body size and composition. Neurobiology of Stress, 2021, 15, 100400.	4.0	3
520	Candidate Genes and Psychiatric Genetics: Tomorrow Never Knows. Psychiatric Annals, 1997, 27, 262-267.	0.1	3
521	Polygenic scores for empathy associate with posttraumatic stress severity in response to certain traumatic events. Neurobiology of Stress, 2022, 17, 100439.	4.0	3
522	Intravenous physostigmine increases cerebrospinal fluid neuropeptide-Y. Biological Psychiatry, 1989, 26, 623-630.	1.3	2

#	Article	IF	CITATIONS
523	Dopamine D2 Receptor and Tourette's Syndrome-Reply. Archives of Neurology, 1995, 52, 442-443.	4.5	2
524	Inviting in the Exome for Alcohol and Smoking Traits. Biological Psychiatry, 2019, 85, 889-890.	1.3	2
525	Multivariate Analyses Reveal Biological Components Related to Neuronal Signaling and Immunity Mediating Electroencephalograms Abnormalities in Alcoholâ€Dependent Individuals from the Collaborative Study on the Genetics of Alcoholism Cohort. Alcoholism: Clinical and Experimental Research, 2019, 43, 1462-1477.	2.4	2
526	Prevalence of predicted gene-drug interactions for antidepressants in the treatment of major depressive disorder in the Precision Medicine in Mental Health Care Study. Journal of Affective Disorders, 2021, 282, 1272-1277.	4.1	2
527	Tryptophan hydroxylase genotype is associated with impulsive-aggression measures: a preliminary study. , 1998, 81, 13.		2
528	Association Study of Alcoholism Subtypes with a Functional Promoter Polymorphism in the Serotonin Transporter Protein Gene. Alcoholism: Clinical and Experimental Research, 2002, 26, 1330-1335.	2.4	2
529	Unique and joint associations of polygenic risk for major depression and opioid use disorder with endogenous opioid system function. Neuropsychopharmacology, 2022, 47, 1784-1790.	5.4	2
530	No allelic association of an exon 13 polymorphism of the Gsl $^\pm$ gene to alcohol and/or drug dependence. Addiction Biology, 1997, 2, 309-316.	2.6	1
531	550. Phenotypic assessment of cocaine-induced paranoia and psychotic symptoms (CIPPS). Biological Psychiatry, 2000, 47, S167-S168.	1.3	1
532	Further analyses support the association between light eye color and alcohol dependence. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 757-760.	1.7	1
533	Collaborative phenotype inference from comorbid substance use disorders and genotypes., 2017, 2017, 392-397.		1
534	GWAS and network analysis of coâ€occurring nicotine and alcohol dependence identifies significantly associated alleles and network. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, 3-11.	1.7	1
535	Ciliary neurotrophic factor null allele frequencies in schizophrenia, affective disorders, and Alzheimer's disease., 1997, 74, 497.		1
536	Identifying factors associated with opioid cessation in a biracial sample using machine learning., 2020, 1, 27-41.		1
537	Medical informatics is a research field. Medical Informatics = Medecine Et Informatique, 1984, 9, 221-221.	0.8	0
538	Report of the DNA committee and catalogs of cloned and mapped genes and DNA polymorphisms pp. 892–921. Cytogenetic and Genome Research, 1989, 51, 892-921.	1.1	0
539	Candidate gene study of the catecholamine system in schizophrenia. European Neuropsychopharmacology, 1994, 4, 373.	0.7	0
540	Bipolar affective disorder is not linked to norepinephrine transporter protein or serotonin transporter protein genes. European Neuropsychopharmacology, 1994, 4, 316.	0.7	0

#	Article	IF	CITATIONS
541	Genetic variation and drug dependence risk factors. , 2002, , 372-388.		О
542	A sequencingâ€based survey of functional <i>APAF1</i> alleles in a large sample of individuals with affective illness and population controls. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 332-335.	1.7	0
543	P3-143: Amyloid burden is associated with decreased gray matter volume but not episodic memory performance in cognitively normal first-degree relatives at varying ApoE4 risk for Alzheimer's disease. , 2015, 11, P680-P681.		O
544	IC-P-101: Amyloid burden is associated with decreased gray matter volume but not episodic memory performance in cognitively normal first-degree relatives at varying ApoE4 risk for Alzheimer's disease. , 2015, 11, P69-P69.		0
545	Introduction to statistical methods in genome-wide association studies. , 0, , 26-52.		O
546	Inferring phenotypes from substance use via collaborative matrix completion. BMC Systems Biology, 2018, 12, 104.	3.0	0
547	Introducing <b><i>Complex Psychiatry</i></b> . Complex Psychiatry, 2020, 6, 2-4.	0.9	0
548	Genome-wide association study of phenotypes measuring progression from first cocaine or opioid use to dependence reveals novel risk genes. Exploration of Medicine, 0, , .	1.5	0
549	Molecular Genetic Analysis of Plasma Dopamine β-Hydroxylase in Depression. Advances in Behavioral Biology, 2002, , 423-426.	0.2	0
550	Genetics of Substance Dependence. , 2007, , 865-881.		0
551	Heritability and Genetics of Anxiety Disorders. , 2008, , .		O
552	The current status of linkage studies in schizophrenia. Research Publications - Association for Research in Nervous and Mental Disease, 1991, 69, 137-52.	0.1	0
553	Title is missing!. , 2020, 16, e1009036.		0
554	Title is missing!. , 2020, 16, e1009036.		0
555	Title is missing!. , 2020, 16, e1009036.		0
556	Title is missing!. , 2020, 16, e1009036.		O