Joel Gelernter

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

565	29,119	92	145
papers	citations	h-index	g-index
635	34,613 ext. citations	6.7	6.93
ext. papers		avg, IF	L-index

#	Paper	IF	Citations
565	Post-treatment effects of topiramate on alcohol-related outcomes: A combined analysis of two placebo-controlled trials <i>Addiction Biology</i> , 2022 , 27, e13130	4.6	1
564	Epigenome-wide association study of posttraumatic stress disorder identifies novel loci in U.S. military veterans <i>Translational Psychiatry</i> , 2022 , 12, 65	8.6	2
563	Using phenotype risk scores to enhance gene discovery for generalized anxiety disorder and posttraumatic stress disorder <i>Molecular Psychiatry</i> , 2022 ,	15.1	2
562	Polygenic scores for empathy associate with posttraumatic stress severity in response to certain traumatic events <i>Neurobiology of Stress</i> , 2022 , 17, 100439	7.6	O
561	Genetically regulated multi-omics study for symptom clusters of posttraumatic stress disorder highlights pleiotropy with hematologic and cardio-metabolic traits <i>Molecular Psychiatry</i> , 2022 ,	15.1	2
560	Diagnostic Reliability and Validity of the Semi-Structured Assessment for Drug Dependence and Alcoholism (SSADDA) Chinese Version. <i>Complex Psychiatry</i> , 2021 , 6, 62-67	2.3	1
559	The addiction risk factor: A unitary genetic vulnerability characterizes substance use disorders and their associations with common correlates. <i>Neuropsychopharmacology</i> , 2021 ,	8.7	4
558	Drinking and smoking polygenic risk is associated with childhood and early-adulthood psychiatric and behavioral traits independently of substance use and psychiatric genetic risk. <i>Translational Psychiatry</i> , 2021 , 11, 586	8.6	1
557	Attachment Style Moderates Polygenic Risk for Incident Posttraumatic Stress in U.S. Military Veterans: A 7-Year, Nationally Representative, Prospective Cohort Study <i>Biological Psychiatry</i> , 2021 ,	7.9	2
556	Ancestry may confound genetic machine learning: Candidate-gene prediction of opioid use disorder as an example. <i>Drug and Alcohol Dependence</i> , 2021 , 229, 109115	4.9	O
555	Prevalence of predicted gene-drug interactions for antidepressants in the treatment of major depressive disorder in the Precision Medicine in Mental Health Care Study. <i>Journal of Affective Disorders</i> , 2021 , 282, 1272-1277	6.6	O
554	Pleiotropic effects of telomere length loci with brain morphology and brain tissue expression. <i>Human Molecular Genetics</i> , 2021 , 30, 1360-1370	5.6	1
553	Dissecting the heterogeneity of posttraumatic stress disorder: differences in polygenic risk, stress exposures, and course of PTSD subtypes. <i>Psychological Medicine</i> , 2021 , 1-9	6.9	2
552	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. <i>Nature Genetics</i> , 2021 , 53, 817-829	36.3	83
551	The impact of removing former drinkers from genome-wide association studies of AUDIT-C. <i>Addiction</i> , 2021 , 116, 3044-3054	4.6	1
550	Polygenic risk for traumatic loss-related PTSD in US military veterans: Protective effect of secure attachment style. <i>World Journal of Biological Psychiatry</i> , 2021 , 22, 792-799	3.8	2
549	Bi-ancestral depression GWAS in the Million Veteran Program and meta-analysis in >1.2 million individuals highlight new therapeutic directions. <i>Nature Neuroscience</i> , 2021 , 24, 954-963	25.5	32

(2021-2021)

548	Sex-stratified gene-by-environment genome-wide interaction study of trauma, posttraumatic-stress, and suicidality. <i>Neurobiology of Stress</i> , 2021 , 14, 100309	7.6	7
547	The relationship between cannabis and schizophrenia: a genetically informed perspective. <i>Addiction</i> , 2021 , 116, 3227-3234	4.6	6
546	Integration of evidence across human and model organism studies: A meeting report. <i>Genes, Brain and Behavior</i> , 2021 , 20, e12738	3.6	1
545	Psychological trauma and the genetic overlap between posttraumatic stress disorder and major depressive disorder. <i>Psychological Medicine</i> , 2021 , 1-10	6.9	3
544	Genome-wide association study of stimulant dependence. <i>Translational Psychiatry</i> , 2021 , 11, 363	8.6	0
543	Investigation of convergent and divergent genetic influences underlying schizophrenia and alcohol use disorder. <i>Psychological Medicine</i> , 2021 , 1-9	6.9	O
542	Genetics of substance use disorders in the era of big data. <i>Nature Reviews Genetics</i> , 2021 , 22, 712-729	30.1	7
541	Dissecting the genetic association of C-reactive protein with PTSD, traumatic events, and social support. <i>Neuropsychopharmacology</i> , 2021 , 46, 1071-1077	8.7	10
540	Attachment Style Moderates Polygenic Risk for Posttraumatic Stress in United States Military Veterans: Results From the National Health and Resilience in Veterans Study. <i>Biological Psychiatry</i> , 2021 , 89, 878-887	7.9	9
539	GABAergic polygenic risk for cocaine use disorder is negatively correlated with precuneus activity during cognitive control in African American individuals. <i>Addictive Behaviors</i> , 2021 , 114, 106695	4.2	1
538	Characterizing the effect of background selection on the polygenicity of brain-related traits. <i>Genomics</i> , 2021 , 113, 111-119	4.3	5
537	Study design and implementation of the PRecision Medicine In MEntal health Care (PRIME Care) Trial. <i>Contemporary Clinical Trials</i> , 2021 , 101, 106247	2.3	2
536	Transcriptomic organization of the human brain in post-traumatic stress disorder. <i>Nature Neuroscience</i> , 2021 , 24, 24-33	25.5	44
535	Multivariate genome-wide analysis of education, socioeconomic status and brain phenome. <i>Nature Human Behaviour</i> , 2021 , 5, 482-496	12.8	6
534	Multi-environment gene interactions linked to the interplay between polysubstance dependence and suicidality. <i>Translational Psychiatry</i> , 2021 , 11, 34	8.6	3
533	Genome-wide association study of phenotypes measuring progression from first cocaine or opioid use to dependence reveals novel risk genes. <i>Exploration of Medicine</i> , 2021 , 2, 60-73	1.1	1
532	Epigenomic Profiles of African-American Val122Ile Carriers Reveals Putatively Dysregulated Amyloid Mechanisms. <i>Circulation Genomic and Precision Medicine</i> , 2021 , 14, e003011	5.2	3
531	Multivariate analysis of 1.5 million people identifies genetic associations with traits related to self-regulation and addiction. <i>Nature Neuroscience</i> , 2021 , 24, 1367-1376	25.5	10

530	Combined analysis of the moderating effect of a GRIK1 polymorphism on the effects of topiramate for treating alcohol use disorder. <i>Drug and Alcohol Dependence</i> , 2021 , 225, 108762	4.9	2
529	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). <i>Drug and Alcohol Dependence</i> , 2021 , 229, 109047	4.9	1
528	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. <i>Biological Psychiatry</i> , 2021 ,	7.9	11
527	Enhancing Discovery of Genetic Variants for Posttraumatic Stress Disorder Through Integration of Quantitative Phenotypes and Trauma Exposure Information. <i>Biological Psychiatry</i> , 2021 ,	7.9	3
526	Disentangling sex differences in the shared genetic architecture of posttraumatic stress disorder, traumatic experiences, and social support with body size and composition. <i>Neurobiology of Stress</i> , 2021 , 15, 100400	7.6	
525	Genome-wide association analyses of post-traumatic stress disorder and its symptom subdomains in the Million Veteran Program. <i>Nature Genetics</i> , 2021 , 53, 174-184	36.3	40
524	Expanding the genetic architecture of nicotine dependence and its shared genetics with multiple traits. <i>Nature Communications</i> , 2020 , 11, 5562	17.4	25
523	Association of OPRM1 Functional Coding Variant With Opioid Use Disorder: A Genome-Wide Association Study. <i>JAMA Psychiatry</i> , 2020 , 77, 1072-1080	14.5	46
522	Tubulin Polymerization Promoting Protein (TPPP) gene methylation and corpus callosum measures in maltreated children. <i>Psychiatry Research - Neuroimaging</i> , 2020 , 298, 111058	2.9	3
521	Genomic influences on self-reported childhood maltreatment. <i>Translational Psychiatry</i> , 2020 , 10, 38	8.6	20
520	Leveraging genome-wide data to investigate differences between opioid use vs. opioid dependence in 41,176 individuals from the Psychiatric Genomics Consortium. <i>Molecular Psychiatry</i> , 2020 , 25, 1673-1687	15.1	30
519	Genome-Wide Association Study of Opioid Cessation. <i>Journal of Clinical Medicine</i> , 2020 , 9,	5.1	10
0	Systematic review and meta-analysis of the moderating effect of rs1799971 in OPRM1, the		
518	mu-opioid receptor gene, on response to naltrexone treatment of alcohol use disorder. <i>Addiction</i> , 2020 , 115, 1426-1437	4.6	11
518	mu-opioid receptor gene, on response to naltrexone treatment of alcohol use disorder. Addiction,	4.6 4·5	8
	mu-opioid receptor gene, on response to naltrexone treatment of alcohol use disorder. <i>Addiction</i> , 2020 , 115, 1426-1437 A genome-wide association study of cocaine use disorder accounting for phenotypic heterogeneity		
517	mu-opioid receptor gene, on response to naltrexone treatment of alcohol use disorder. <i>Addiction</i> , 2020 , 115, 1426-1437 A genome-wide association study of cocaine use disorder accounting for phenotypic heterogeneity and gene#nvironment interaction. <i>Journal of Psychiatry and Neuroscience</i> , 2020 , 45, 34-44 Identifying factors associated with opioid cessation in a biracial sample using machine learning.	4.5	8
517 516	mu-opioid receptor gene, on response to naltrexone treatment of alcohol use disorder. <i>Addiction</i> , 2020 , 115, 1426-1437 A genome-wide association study of cocaine use disorder accounting for phenotypic heterogeneity and gene®nvironment interaction. <i>Journal of Psychiatry and Neuroscience</i> , 2020 , 45, 34-44 Identifying factors associated with opioid cessation in a biracial sample using machine learning. <i>Exploration of Medicine</i> , 2020 , 1, 27-41 Genome-wide meta-analysis of problematic alcohol use in 435,563 individuals yields insights into	4.5	8

512	Post-GWAS analysis of six substance use traits improves the identification and functional interpretation of genetic risk loci. <i>Drug and Alcohol Dependence</i> , 2020 , 206, 107703	4.9	8
511	Molecular genetic overlap between posttraumatic stress disorder and sleep phenotypes. <i>Sleep</i> , 2020 , 43,	1.1	9
510	Investigating Causality Between Blood Metabolites and Emotional and Behavioral Responses to Traumatic Stress: a Mendelian Randomization Study. <i>Molecular Neurobiology</i> , 2020 , 57, 1542-1552	6.2	3
509	Genome-wide association study of smoking trajectory and meta-analysis of smoking status in 842,000 individuals. <i>Nature Communications</i> , 2020 , 11, 5302	17.4	15
508	A large-scale genome-wide association study meta-analysis of cannabis use disorder. <i>Lancet Psychiatry,the</i> , 2020 , 7, 1032-1045	23.3	43
507	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. <i>PLoS Genetics</i> , 2020 , 16, e1009036	6	2
506	Differentiating Types of Self-Reported Alcohol Abstinence. AIDS and Behavior, 2020, 24, 655-665	4.3	4
505	Genome-Wide Meta-Analyses of FTND and TTFC Phenotypes. <i>Nicotine and Tobacco Research</i> , 2020 , 22, 900-909	4.9	9
504	Genome-wide scan identifies opioid overdose risk locus close to MCOLN1. <i>Addiction Biology</i> , 2020 , 25, e12811	4.6	7
503	Genotyping Array Design and Data Quality Control in the Million Veteran Program. <i>American Journal of Human Genetics</i> , 2020 , 106, 535-548	11	22
502	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 2020 , 16, e1009036		
501	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 2020 , 16, e1009036		
500	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 2020 , 16, e1009036		
499	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 2020 , 16, e1009036		
498	Genetic associations with suicide attempt severity and genetic overlap with major depression. <i>Translational Psychiatry</i> , 2019 , 9, 22	8.6	48
497	Salivary microRNAs identified by small RNA sequencing and machine learning as potential biomarkers of alcohol dependence. <i>Epigenomics</i> , 2019 , 11, 739-749	4.4	7
496	Genome-wide analyses of psychological resilience in U.S. Army soldiers. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019 , 180, 310-319	3.5	22
495	Inviting in the Exome for Alcohol and Smoking Traits. <i>Biological Psychiatry</i> , 2019 , 85, 889-890	7.9	1

494	Association of Economic Status and Educational Attainment With Posttraumatic Stress Disorder: A Mendelian Randomization Study. <i>JAMA Network Open</i> , 2019 , 2, e193447	10.4	18
493	Validation of an Electronic Medical Record-Based Algorithm for Identifying Posttraumatic Stress Disorder in U.S. Veterans. <i>Journal of Traumatic Stress</i> , 2019 , 32, 226-237	3.8	11
492	Multivariate Analyses Reveal Biological Components Related to Neuronal Signaling and Immunity Mediating Electroencephalograms Abnormalities in Alcohol-Dependent Individuals from the Collaborative Study on the Genetics of Alcoholism Cohort. <i>Alcoholism: Clinical and Experimental</i>	3.7	2
491	Research, 2019, 43, 1462-1477 Increased Risk of Multiple Outpatient Surgeries in African-American Carriers of Val122Ile Mutation Is Modulated by Non-Coding Variants. <i>Journal of Clinical Medicine</i> , 2019, 8,	5.1	9
490	Genetics of Alcoholism. <i>Current Psychiatry Reports</i> , 2019 , 21, 26	9.1	23
489	Alcohol-responsive genes identified in human iPSC-derived neural cultures. <i>Translational Psychiatry</i> , 2019 , 9, 96	8.6	9
488	Genomewide Study of Epigenetic Biomarkers of Opioid Dependence in European- American Women. <i>Scientific Reports</i> , 2019 , 9, 4660	4.9	14
487	Genome-wide Association Study of Maximum Habitual Alcohol Intake in >140,000 U.S. European and African American Veterans Yields Novel Risk Loci. <i>Biological Psychiatry</i> , 2019 , 86, 365-376	7.9	37
486	Evidence of causal effect of major depression on alcohol dependence: findings from the psychiatric genomics consortium. <i>Psychological Medicine</i> , 2019 , 49, 1218-1226	6.9	33
485	Genome-wide association study of alcohol consumption and use disorder in 274,424 individuals from multiple populations. <i>Nature Communications</i> , 2019 , 10, 1499	17.4	164
485 484		17.4	164
	from multiple populations. <i>Nature Communications</i> , 2019 , 10, 1499 Genomics of posttraumatic stress disorder in veterans: Methods and rationale for Veterans Affairs	, ' 	
484	Genomics of posttraumatic stress disorder in veterans: Methods and rationale for Veterans Affairs Cooperative Study #575B. <i>International Journal of Methods in Psychiatric Research</i> , 2019 , 28, e1767 Apolipoprotein E gene polymorphism, posttraumatic stress disorder, and cognitive function in older U.S. veterans: Results from the National Health and Resilience in Veterans Study. <i>Depression</i>	4-3	5
484	Genomics of posttraumatic stress disorder in veterans: Methods and rationale for Veterans Affairs Cooperative Study #575B. <i>International Journal of Methods in Psychiatric Research</i> , 2019 , 28, e1767 Apolipoprotein E gene polymorphism, posttraumatic stress disorder, and cognitive function in older U.S. veterans: Results from the National Health and Resilience in Veterans Study. <i>Depression and Anxiety</i> , 2019 , 36, 834-845 Epigenome-Wide DNA Methylation Association Analysis Identified Novel Loci in Peripheral Cells for Alcohol Consumption Among European American Male Veterans. <i>Alcoholism: Clinical and</i>	4-3	5
484 483 482	Genomics of posttraumatic stress disorder in veterans: Methods and rationale for Veterans Affairs Cooperative Study #575B. International Journal of Methods in Psychiatric Research, 2019, 28, e1767 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 Epigenome-Wide DNA 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 Patterns and Correlates of Prescription Opioid Receipt Among US Veterans: A National, 18-Year	4·3 8·4 3·7	5 10 16
484 483 482 481	Genomics of posttraumatic stress disorder in veterans: Methods and rationale for Veterans Affairs Cooperative Study #575B. International Journal of Methods in Psychiatric Research, 2019, 28, e1767 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 Epigenome-Wide DNA 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 Patterns and Correlates of Prescription Opioid Receipt Among US Veterans: A National, 18-Year Observational Cohort Study. AIDS and Behavior, 2019, 23, 3340-3349 Genome-wide association study of post-traumatic stress disorder reexperiencing symptoms in	4·3 8·4 3·7 4·3	5 10 16 6
484 483 482 481 480	Genomics of posttraumatic stress disorder in veterans: Methods and rationale for Veterans Affairs Cooperative Study #575B. International Journal of Methods in Psychiatric Research, 2019, 28, e1767 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 Epigenome-Wide DNA 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 Patterns and Correlates of Prescription Opioid Receipt Among US Veterans: A National, 18-Year Observational Cohort Study. AIDS and Behavior, 2019, 23, 3340-3349 Genome-wide association study of post-traumatic stress disorder reexperiencing symptoms in >165,000 US veterans. Nature Neuroscience, 2019, 22, 1394-1401 International meta-analysis of PTSD genome-wide association studies identifies sex- and	4·3 8·4 3·7 4·3 25·5	5 10 16 6

(2018-2019)

476	A regulatory variant of CHRM3 is associated with cannabis-induced hallucinations in European Americans. <i>Translational Psychiatry</i> , 2019 , 9, 309	8.6	O
475	Deciphering the Biological Mechanisms Underlying the Genome-Wide Associations between Computerized Device Use and Psychiatric Disorders. <i>Journal of Clinical Medicine</i> , 2019 , 8,	5.1	5
474	Genomewide Gene-by-Sex Interaction Scans Identify ADGRV1 for Sex Differences in Opioid Dependent African Americans. <i>Scientific Reports</i> , 2019 , 9, 18070	4.9	6
473	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. <i>Psychological Medicine</i> , 2019 , 49, 1166-11	7 5 .9	52
472	Discovery of the first genome-wide significant risk loci for attention deficit/hyperactivity disorder. <i>Nature Genetics</i> , 2019 , 51, 63-75	36.3	826
471	GWAS and network analysis of co-occurring nicotine and alcohol dependence identifies significantly associated alleles and network. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019 , 180, 3-11	3.5	1
470	Using DNA methylation to validate an electronic medical record phenotype for smoking. <i>Addiction Biology</i> , 2019 , 24, 1056-1065	4.6	7
469	Attachment style moderates effects of polymorphisms and childhood abuse on post-traumatic stress symptoms: Results from the National Health and Resilience in Veterans Study. <i>World Journal of Biological Psychiatry</i> , 2019 , 20, 289-300	3.8	17
468	Genomewide Association Study of Alcohol Dependence and Related Traits in a Thai Population. <i>Alcoholism: Clinical and Experimental Research</i> , 2018 , 42, 861-868	3.7	16
467	Genome-wide analysis of insomnia disorder. <i>Molecular Psychiatry</i> , 2018 , 23, 2238-2250	15.1	39
466	Genome-wide Association Study Identifies a Regulatory Variant of RGMA Associated With Opioid Dependence in European Americans. <i>Biological Psychiatry</i> , 2018 , 84, 762-770	7.9	41
465	Local adaptation in European populations affected the genetics of psychiatric disorders and behavioral traits. <i>Genome Medicine</i> , 2018 , 10, 24	14.4	2
464	Multivariate Pattern Analysis of Genotype-Phenotype Relationships in Schizophrenia. <i>Schizophrenia Bulletin</i> , 2018 , 44, 1045-1052	1.3	8
463	A genome-wide gene-by-trauma interaction study of alcohol misuse in two independent cohorts identifies PRKG1 as a risk locus. <i>Molecular Psychiatry</i> , 2018 , 23, 154-160	15.1	32
462	ADH1B: From alcoholism, natural selection, and cancer to the human phenome. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018 , 177, 113-125	3.5	34
461	Psychiatric Genomics: An Update and an Agenda. <i>American Journal of Psychiatry</i> , 2018 , 175, 15-27	11.9	328
460	Genome-wide association study across European and African American ancestries identifies a SNP in DNMT3B contributing to nicotine dependence. <i>Molecular Psychiatry</i> , 2018 , 23, 1911-1919	15.1	52
459	Cortical Emyloid burden, gray matter, and memory in adults at varying APOE prisk for Alzheimerk disease. <i>Neurobiology of Aging</i> , 2018 , 61, 207-214	5.6	20

458	Genome-wide association meta-analysis of age at first cannabis use. Addiction, 2018, 113, 2073-2086	4.6	16
457	AUDIT-C and ICD codes as phenotypes for harmful alcohol use: association with ADH1B polymorphisms in two US populations. <i>Addiction</i> , 2018 , 113, 2214-2224	4.6	13
456	Biochemical, demographic, and self-reported tobacco-related predictors of the acute heart rate response to nicotine in smokers. <i>Pharmacology Biochemistry and Behavior</i> , 2018 , 173, 36-43	3.9	2
455	Genome-wide association study of cognitive flexibility assessed by the Wisconsin Card Sorting Test. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018 , 177, 511-519	3.5	O
454	GWAS of lifetime cannabis use reveals new risk loci, genetic overlap with psychiatric traits, and a causal influence of schizophrenia. <i>Nature Neuroscience</i> , 2018 , 21, 1161-1170	25.5	270
453	Methylation in OTX2 and related genes, maltreatment, and depression in children. <i>Neuropsychopharmacology</i> , 2018 , 43, 2204-2211	8.7	31
452	Translational studies support a role for serotonin 2B receptor (HTR2B) gene in aggression-related cannabis response. <i>Molecular Psychiatry</i> , 2018 , 23, 2277-2286	15.1	14
451	Genetically determined schizophrenia is not associated with impaired glucose homeostasis. <i>Schizophrenia Research</i> , 2018 , 195, 286-289	3.6	16
450	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. <i>Depression and Anxiety</i> , 2018 , 35, 168-177	8.4	11
449	Trauma exposure interacts with the genetic risk of bipolar disorder in alcohol misuse of US soldiers. <i>Acta Psychiatrica Scandinavica</i> , 2018 , 137, 148-156	6.5	8
448	A Genetic Investigation of Sex Bias in the Prevalence of Attention-Deficit/Hyperactivity Disorder. <i>Biological Psychiatry</i> , 2018 , 83, 1044-1053	7.9	93
447	Genome-wide association study identifies a novel locus for cannabis dependence. <i>Molecular Psychiatry</i> , 2018 , 23, 1293-1302	15.1	28
446	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. <i>Nature Neuroscience</i> , 2018 , 21, 1656-1669	25.5	257
445	Inferring phenotypes from substance use via collaborative matrix completion. <i>BMC Systems Biology</i> , 2018 , 12, 104	3.5	
444	Genome-wide association study identifies glutamate ionotropic receptor GRIA4 as a risk gene for comorbid nicotine dependence and major depression. <i>Translational Psychiatry</i> , 2018 , 8, 208	8.6	8
443	Phenomic Impact of Genetically-Determined Euthyroid Function and Molecular Differences between Thyroid Disorders. <i>Journal of Clinical Medicine</i> , 2018 , 7,	5.1	10
442	Risk Locus Identification Ties Alcohol Withdrawal Symptoms to SORCS2. <i>Alcoholism: Clinical and Experimental Research</i> , 2018 , 42, 2337-2348	3.7	8
441	Adverse Childhood Experiences, Epigenetic Measures, and Obesity in Youth. <i>Journal of Pediatrics</i> , 2018 , 202, 150-156.e3	3.6	26

(2017-2017)

440	Effects of ANK3 variation on gray and white matter in bipolar disorder. <i>Molecular Psychiatry</i> , 2017 , 22, 1345-1351	15.1	15
439	A protocadherin gene cluster regulatory variant is associated with nicotine withdrawal and the urge to smoke. <i>Molecular Psychiatry</i> , 2017 , 22, 242-249	15.1	7
438	Genome-wide association study of body mass index in subjects with alcohol dependence. <i>Addiction Biology</i> , 2017 , 22, 535-549	4.6	18
437	Genome-wide association study of therapeutic opioid dosing identifies a novel locus upstream of OPRM1. <i>Molecular Psychiatry</i> , 2017 , 22, 346-352	15.1	67
436	Genomewide Association Study of Alcohol Dependence Identifies Risk Loci Altering Ethanol-Response Behaviors in Model Organisms. <i>Alcoholism: Clinical and Experimental Research</i> , 2017 , 41, 911-928	3.7	30
435	Genetic risk variants for social anxiety. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017 , 174, 120-131	3.5	30
434	Alcohol and nicotine codependence-associated DNA methylation changes in promoter regions of addiction-related genes. <i>Scientific Reports</i> , 2017 , 7, 41816	4.9	13
433	Phenome-wide association study for CYP2A6 alleles: rs113288603 is associated with hearing loss symptoms in elderly smokers. <i>Scientific Reports</i> , 2017 , 7, 1034	4.9	5
432	Genetic risk variants for social anxiety. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017 , 174, 470-482	3.5	2
431	Genetic-epigenetic interactions in cis: a major focus in the post-GWAS era. <i>Genome Biology</i> , 2017 , 18, 120	18.3	76
430	Validating Harmful Alcohol Use as a Phenotype for Genetic Discovery Using Phosphatidylethanol and a Polymorphism in ADH1B. <i>Alcoholism: Clinical and Experimental Research</i> , 2017 , 41, 998-1003	3.7	12
429	Genetic Risk Variants Associated With Comorbid Alcohol Dependence and Major Depression. <i>JAMA Psychiatry</i> , 2017 , 74, 1234-1241	14.5	47
428	Genome-Wide Association Studies of a Broad Spectrum of Antisocial Behavior. <i>JAMA Psychiatry</i> , 2017 , 74, 1242-1250	14.5	124
427	Ancestry-specific and sex-specific risk alleles identified in a genome-wide gene-by-alcohol dependence interaction study of risky sexual behaviors. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017 , 174, 846-853	3.5	14
426	A putative causal relationship between genetically determined female body shape and posttraumatic stress disorder. <i>Genome Medicine</i> , 2017 , 9, 99	14.4	23
425	identified in a genome-wide gene Itannabis dependence interaction analysis of risky sexual behaviours. <i>Journal of Psychiatry and Neuroscience</i> , 2017 , 42, 252-261	4.5	5
424	Genetic variation of ANK3 is associated with lower white matter structural integrity in bipolar disorder. <i>Molecular Psychiatry</i> , 2017 , 22, 1225	15.1	22
423	Genomewide association studies of suicide attempts in US soldiers. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017 , 174, 786-797	3.5	34

422	Oxytocin receptor gene polymorphisms, attachment, and PTSD: Results from the National Health and Resilience in Veterans Study. <i>Journal of Psychiatric Research</i> , 2017 , 94, 139-147	5.2	34
421	Sex Differences in Methamphetamine Use and Dependence in a Thai Treatment Center. <i>Journal of Addiction Medicine</i> , 2017 , 11, 19-27	3.8	18
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(2015-2016)

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(2014-2015)

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(2011-2012)

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26	DRD2 and FOXP2 are implicated in the associations between computerized device use and psychiatric disorders	1
25	Psychiatric Genomics: An Update and an Agenda	3
24	Discovery of the first genome-wide significant risk loci for ADHD	62
23	The effect of the genetic liability to autism spectrum disorder on emotion recognition in young unaffected probands from a population-based cohort	1
22	GWAS including 82,707 subjects identifies functional coding variant in OPRM1 gene associated with opioid use disorder	2
21	Dissecting the association of C-reactive protein levels with PTSD, traumatic events, and social support	1
20	Measuring genetic variation in the multi-ethnic Million Veteran Program (MVP)	4
19	Expanding the Genetic Architecture of Nicotine Dependence and its Shared Genetics with Multiple Traits: Findings from the Nicotine Dependence GenOmics (iNDiGO) Consortium	3
18	Transcriptomic Organization of Human Posttraumatic Stress Disorder	2
17	Natural selection influenced the genetic architecture of brain structure, behavioral and neuropsychiatric traits	1
16	GWAS of Depression Phenotypes in the Million Veteran Program and Meta-analysis in More than 1.2 Million Participants Yields 178 Independent Risk Loci	7
15	Trauma and posttraumatic stress interact with sex-specific risk loci for suicidality and converge on brain extracellular matrix biology and synaptic plasticity	2
14	Ancestry May Confound Genetic Machine Learning: Candidate-Gene Prediction of Opioid Use Disorder as an Example	2
13	Genome-wide association study of over 40,000 bipolar disorder cases provides new insights into the underlying biology	11
12	Multivariate genomic analysis of 1.5 million people identifies genes related to addiction, antisocial behavior, and health	6
11	Trans-ancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders	7
10	Evidence of causal effect of major depression on alcohol dependence: Findings from the Psychiatric Genomics Consortium	4
9	Largest genome-wide association study for PTSD identifies genetic risk loci in European and African ancestries and implicates novel biological pathways	6

LIST OF PUBLICATIONS

8	Economic status mediates the relationship between educational attainment and posttraumatic stress disorder: a multivariable Mendelian randomization study	1
7	Genome wide meta-analysis identifies genomic relationships, novel loci, and pleiotropic mechanisms across eight psychiatric disorders	10
6	Meta-analysis of problematic alcohol use in 435,563 individuals identifies 29 risk variants and yields insights into biology, pleiotropy and causality	6
5	Genomic Characterization of Posttraumatic Stress Disorder in a Large US Military Veteran Sample	7
4	Leveraging genome-wide data to investigate differences between opioid use vs. opioid dependence in 41,176 individuals from the Psychiatric Genomics Consortium	2
3	Introduction to statistical methods in genome-wide association studies26-52	
2	The Addiction Risk Factor: A Unitary Genetic Vulnerability Characterizes Substance Use Disorders and Their Associations with Common Correlates	3
1	Genetic Factors in Social Anxiety Disorder53-66	1