Jordan W Smoller

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65 24,147 153 253 h-index g-index citations papers 6.75 285 10 32,954 L-index ext. citations ext. papers avg, IF

#	Paper	IF	Citations
253	The organization of the human cerebral cortex estimated by intrinsic functional connectivity. Journal of Neurophysiology, 2011 , 106, 1125-65	3.2	3997
252	Molecular genetics of attention-deficit/hyperactivity disorder. <i>Biological Psychiatry</i> , 2005 , 57, 1313-23	7.9	1676
251	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. <i>Nature Genetics</i> , 2018 , 50, 668-681	36.3	1301
250	Collaborative genome-wide association analysis supports a role for ANK3 and CACNA1C in bipolar disorder. <i>Nature Genetics</i> , 2008 , 40, 1056-8	36.3	949
249	A mega-analysis of genome-wide association studies for major depressive disorder. <i>Molecular Psychiatry</i> , 2013 , 18, 497-511	15.1	853
248	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , 2018 , 360,	33.3	666
247	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019 , 51, 793-803	36.3	662
246	Pleiotropy in complex traits: challenges and strategies. <i>Nature Reviews Genetics</i> , 2013 , 14, 483-95	30.1	649
245	Common genetic variants influence human subcortical brain structures. <i>Nature</i> , 2015 , 520, 224-9	50.4	601
244	The ENIGMA Consortium: large-scale collaborative analyses of neuroimaging and genetic data. Brain Imaging and Behavior, 2014 , 8, 153-82	4.1	539
243	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. <i>Nature Genetics</i> , 2017 , 49, 27-35	36.3	530
242	Identification of common variants associated with human hippocampal and intracranial volumes. <i>Nature Genetics</i> , 2012 , 44, 552-61	36.3	498
241	Family, twin, and adoption studies of bipolar disorder 2003 , 123C, 48-58		443
240	Psychiatric Genomics: An Update and an Agenda. American Journal of Psychiatry, 2018, 175, 15-27	11.9	328
239	The "All of Us" Research Program. New England Journal of Medicine, 2019, 381, 668-676	59.2	325
238	Polygenic dissection of diagnosis and clinical dimensions of bipolar disorder and schizophrenia. <i>Molecular Psychiatry</i> , 2014 , 19, 1017-1024	15.1	258
237	Individual differences in amygdala-medial prefrontal anatomy link negative affect, impaired social functioning, and polygenic depression risk. <i>Journal of Neuroscience</i> , 2012 , 32, 18087-100	6.6	223

(2016-2016)

236	The Genetics of Stress-Related Disorders: PTSD, Depression, and Anxiety Disorders. Neuropsychopharmacology, 2016 , 41, 297-319	8.7	220
235	QT interval and antidepressant use: a cross sectional study of electronic health records. <i>BMJ, The</i> , 2013 , 346, f288	5.9	205
234	Brain Genomics Superstruct Project initial data release with structural, functional, and behavioral measures. <i>Scientific Data</i> , 2015 , 2, 150031	8.2	204
233	Combined analysis from eleven linkage studies of bipolar disorder provides strong evidence of susceptibility loci on chromosomes 6q and 8q. <i>American Journal of Human Genetics</i> , 2005 , 77, 582-95	11	192
232	Polygenic prediction via Bayesian regression and continuous shrinkage priors. <i>Nature Communications</i> , 2019 , 10, 1776	17.4	191
231	Antidepressant use and risk of incident cardiovascular morbidity and mortality among postmenopausal women in the Womenß Health Initiative study. <i>Archives of Internal Medicine</i> , 2009 , 169, 2128-39		182
230	Novel genetic loci associated with hippocampal volume. <i>Nature Communications</i> , 2017 , 8, 13624	17.4	173
229	Assessment of Bidirectional Relationships Between Physical Activity and Depression Among Adults: A 2-Sample Mendelian Randomization Study. <i>JAMA Psychiatry</i> , 2019 , 76, 399-408	14.5	165
228	Genetic influences on schizophrenia and subcortical brain volumes: large-scale proof of concept. <i>Nature Neuroscience</i> , 2016 , 19, 420-431	25.5	163
227	Joint analysis of psychiatric disorders increases accuracy of risk prediction for schizophrenia, bipolar disorder, and major depressive disorder. <i>American Journal of Human Genetics</i> , 2015 , 96, 283-94	11	161
226	Cross-disorder genomewide analysis of schizophrenia, bipolar disorder, and depression. <i>American Journal of Psychiatry</i> , 2010 , 167, 1254-63	11.9	157
225	The genetic architecture of the human cerebral cortex. <i>Science</i> , 2020 , 367,	33.3	156
224	Predicting Suicidal Behavior From Longitudinal Electronic Health Records. <i>American Journal of Psychiatry</i> , 2017 , 174, 154-162	11.9	156
223	International meta-analysis of PTSD genome-wide association studies identifies sex- and ancestry-specific genetic risk loci. <i>Nature Communications</i> , 2019 , 10, 4558	17.4	151
222	Developmental timing of child maltreatment and symptoms of depression and suicidal ideation in young adulthood: results from the National Longitudinal Study of Adolescent Health. <i>Depression and Anxiety</i> , 2013 , 30, 955-64	8.4	149
221	Psychiatric genetics and the structure of psychopathology. <i>Molecular Psychiatry</i> , 2019 , 24, 409-420	15.1	149
220	Panic attacks and risk of incident cardiovascular events among postmenopausal women in the Womenß Health Initiative Observational Study. <i>Archives of General Psychiatry</i> , 2007 , 64, 1153-60		147
219	Novel genetic loci underlying human intracranial volume identified through genome-wide association. <i>Nature Neuroscience</i> , 2016 , 19, 1569-1582	25.5	147

218	Genome-wide Association Studies in Ancestrally Diverse Populations: Opportunities, Methods, Pitfalls, and Recommendations. <i>Cell</i> , 2019 , 179, 589-603	56.2	145
217	ENIGMA and the individual: Predicting factors that affect the brain in 35 countries worldwide. <i>NeuroImage</i> , 2017 , 145, 389-408	7.9	142
216	Association of the SLC1A1 glutamate transporter gene and obsessive-compulsive disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2007, 144B, 1027-33	3.5	142
215	Panic and phobic anxiety: defining phenotypes for genetic studies. <i>American Journal of Psychiatry</i> , 1998 , 155, 1152-62	11.9	138
214	The corticotropin-releasing hormone gene and behavioral inhibition in children at risk for panic disorder. <i>Biological Psychiatry</i> , 2005 , 57, 1485-92	7.9	137
213	Phenome-wide heritability analysis of the UK Biobank. <i>PLoS Genetics</i> , 2017 , 13, e1006711	6	131
212	Influence of RGS2 on anxiety-related temperament, personality, and brain function. <i>Archives of General Psychiatry</i> , 2008 , 65, 298-308		120
211	Association of a genetic marker at the corticotropin-releasing hormone locus with behavioral inhibition. <i>Biological Psychiatry</i> , 2003 , 54, 1376-81	7.9	120
210	Genome-wide Association Studies of Posttraumatic Stress Disorder in 2 Cohorts of US Army Soldiers. <i>JAMA Psychiatry</i> , 2016 , 73, 695-704	14.5	114
209	GWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. <i>American Journal of Psychiatry</i> , 2019 , 176, 651-660	11.9	103
208	Genetic determinants of depression: recent findings and future directions. <i>Harvard Review of Psychiatry</i> , 2015 , 23, 1-18	4.1	102
207	Mind the gap: why many geneticists and psychological scientists have discrepant views about gene-environment interaction (GE) research. <i>American Psychologist</i> , 2014 , 69, 249-68	9.5	101
206	Microstructural abnormalities in subcortical reward circuitry of subjects with major depressive disorder. <i>PLoS ONE</i> , 2010 , 5, e13945	3.7	100
205	Interrogating the Genetic Determinants of Touretteß Syndrome and Other Tic Disorders Through Genome-Wide Association Studies. <i>American Journal of Psychiatry</i> , 2019 , 176, 217-227	11.9	95
204	The Psychiatric Genomics Consortium Posttraumatic Stress Disorder Workgroup: Posttraumatic Stress Disorder Enters the Age of Large-Scale Genomic Collaboration. <i>Neuropsychopharmacology</i> , 2015 , 40, 2287-97	8.7	94
203	Individual Differences in Cognitive Control Circuit Anatomy Link Sensation Seeking, Impulsivity, and Substance Use. <i>Journal of Neuroscience</i> , 2016 , 36, 4038-49	6.6	90
202	Polygenic risk of Alzheimer disease is associated with early- and late-life processes. <i>Neurology</i> , 2016 , 87, 481-8	6.5	86
201	An electronic health records study of long-term weight gain following antidepressant use. <i>JAMA Psychiatry</i> , 2014 , 71, 889-96	14.5	84

200	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	
199	Targeted genome screen of panic disorder and anxiety disorder proneness using homology to murine QTL regions. <i>American Journal of Medical Genetics Part A</i> , 2001 , 105, 195-206		82	
198	Genetic architecture of subcortical brain structures in 38,851 individuals. <i>Nature Genetics</i> , 2019 , 51, 16	24 3 6630	6 81	
197	Minimal phenotyping yields genome-wide association signals of low specificity for major depression. <i>Nature Genetics</i> , 2020 , 52, 437-447	36.3	80	
196	Validation of electronic health record phenotyping of bipolar disorder cases and controls. <i>American Journal of Psychiatry</i> , 2015 , 172, 363-72	11.9	79	
195	Heritability analysis with repeat measurements and its application to resting-state functional connectivity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017 , 114, 5521-5526	11.5	76	
194	GENOME-WIDE ASSOCIATION STUDY (GWAS) AND GENOME-WIDE BY ENVIRONMENT INTERACTION STUDY (GWEIS) OF DEPRESSIVE SYMPTOMS IN AFRICAN AMERICAN AND HISPANIC/LATINA WOMEN. <i>Depression and Anxiety</i> , 2016 , 33, 265-80	8.4	76	
193	Association between the 5HT1B receptor gene (HTR1B) and the inattentive subtype of ADHD. <i>Biological Psychiatry</i> , 2006 , 59, 460-7	7.9	74	
192	Penetrance and Pleiotropy of Polygenic Risk Scores for Schizophrenia in 106,160 Patients Across Four Health Care Systems. <i>American Journal of Psychiatry</i> , 2019 , 176, 846-855	11.9	73	
191	Cohort of birth modifies the association between FTO genotype and BMI. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015 , 112, 354-9	11.5	73	
190	Corticotrophin-Releasing Hormone Type 1 Receptor Gene (CRHR1) Variants Predict Posttraumatic Stress Disorder Onset and Course in Pediatric Injury Patients. <i>Disease Markers</i> , 2011 , 30, 89-99	3.2	73	
189	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. <i>Biological Psychiatry</i> , 2017 , 82, 322-329	7.9	68	
188	Childhood Adversity Is Associated with Adult Theory of Mind and Social Affiliation, but Not Face Processing. <i>PLoS ONE</i> , 2015 , 10, e0129612	3.7	65	
187	Genetics of anxiety disorders: the complex road from DSM to DNA. <i>Depression and Anxiety</i> , 2009 , 26, 965-75	8.4	65	
186	Harmonizing Clinical Sequencing and Interpretation for the eMERGE III Network. <i>American Journal of Human Genetics</i> , 2019 , 105, 588-605	11	63	
185	The genetic basis of panic and phobic anxiety disorders. <i>American Journal of Medical Genetics, Part C: Seminars in Medical Genetics</i> , 2008 , 148C, 118-26	3.1	61	
184	Individual Aesthetic Preferences for Faces Are Shaped Mostly by Environments, Not Genes. <i>Current Biology</i> , 2015 , 25, 2684-9	6.3	60	
183	Antidepressant response in patients with major depression exposed to NSAIDs: a pharmacovigilance study. <i>American Journal of Psychiatry</i> , 2012 , 169, 1065-72	11.9	59	

182	Joint Contributions of Rare Copy Number Variants and Common SNPs to Risk for Schizophrenia. American Journal of Psychiatry, 2019 , 176, 29-35	11.9	59
181	Research review: gene-environment interaction research in youth depression - a systematic review with recommendations for future research. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2011 , 52, 1223-38	7.9	58
180	The human ortholog of acid-sensing ion channel gene ASIC1a is associated with panic disorder and amygdala structure and function. <i>Biological Psychiatry</i> , 2014 , 76, 902-10	7.9	57
179	Genetic association analysis of behavioral inhibition using candidate loci from mouse models. <i>American Journal of Medical Genetics Part A</i> , 2001 , 105, 226-35		56
178	Rare copy number variation in treatment-resistant major depressive disorder. <i>Biological Psychiatry</i> , 2014 , 76, 536-41	7.9	54
177	Dopamine genetic risk score predicts depressive symptoms in healthy adults and adults with depression. <i>PLoS ONE</i> , 2014 , 9, e93772	3.7	53
176	Massively expedited genome-wide heritability analysis (MEGHA). <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015 , 112, 2479-84	11.5	52
175	Folate, vitamin B-6, and vitamin B-12 intake and mild cognitive impairment and probable dementia in the Womenß Health Initiative Memory Study. <i>Journal of the Academy of Nutrition and Dietetics</i> , 2015 , 115, 231-241	3.9	51
174	A new role for endophenotypes in the GWAS era: functional characterization of risk variants. <i>Harvard Review of Psychiatry</i> , 2010 , 18, 67-74	4.1	50
173	Stratifying Risk for Renal Insufficiency Among Lithium-Treated Patients: An Electronic Health Record Study. <i>Neuropsychopharmacology</i> , 2016 , 41, 1138-43	8.7	49
172	Genetic modifiers and subtypes in schizophrenia: investigations of age at onset, severity, sex and family history. <i>Schizophrenia Research</i> , 2014 , 154, 48-53	3.6	49
171	Genetic associations with suicide attempt severity and genetic overlap with major depression. <i>Translational Psychiatry</i> , 2019 , 9, 22	8.6	48
170	The use of electronic health records for psychiatric phenotyping and genomics. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018 , 177, 601-612	3.5	48
169	Genome-wide gene-environment analyses of major depressive disorder and reported lifetime traumatic experiences in UK Biobank. <i>Molecular Psychiatry</i> , 2020 , 25, 1430-1446	15.1	47
168	Socioeconomic disadvantage and neural development from infancy through early childhood. <i>International Journal of Epidemiology</i> , 2015 , 44, 1889-99	7.8	47
167	Pathway analyses implicate glial cells in schizophrenia. <i>PLoS ONE</i> , 2014 , 9, e89441	3.7	45
166	Dissociable influences of A and polygenic risk of AD dementia on amyloid and cognition. <i>Neurology</i> , 2018 , 90, e1605-e1612	6.5	43
165	The human cortex possesses a reconfigurable dynamic network architecture that is disrupted in psychosis. <i>Nature Communications</i> , 2018 , 9, 1157	17.4	42

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164	Genetic association analysis of 300 genes identifies a risk haplotype in SLC18A2 for post-traumatic stress disorder in two independent samples. <i>Neuropsychopharmacology</i> , 2014 , 39, 1872-9	8.7	42	
163	Disorders and borders: psychiatric genetics and nosology. <i>American Journal of Medical Genetics Part</i> B: Neuropsychiatric Genetics, 2013 , 162B, 559-78	3.5	41	
162	Contributions of common genetic variants to risk of schizophrenia among individuals of African and Latino ancestry. <i>Molecular Psychiatry</i> , 2020 , 25, 2455-2467	15.1	41	
161	An Exposure-Wide and Mendelian Randomization Approach to Identifying Modifiable Factors for the Prevention of Depression. <i>American Journal of Psychiatry</i> , 2020 , 177, 944-954	11.9	40	
160	Genome-wide analysis of insomnia disorder. <i>Molecular Psychiatry</i> , 2018 , 23, 2238-2250	15.1	39	
159	Multidimensional heritability analysis of neuroanatomical shape. <i>Nature Communications</i> , 2016 , 7, 1329	117.4	38	
158	The AURORA Study: a longitudinal, multimodal library of brain biology and function after traumatic stress exposure. <i>Molecular Psychiatry</i> , 2020 , 25, 283-296	15.1	38	
157	Cryptic and complex chromosomal aberrations in early-onset neuropsychiatric disorders. <i>American Journal of Human Genetics</i> , 2014 , 95, 454-61	11	37	
156	Serotonin gene polymorphisms and bipolar I disorder: focus on the serotonin transporter. <i>Annals of Medicine</i> , 2005 , 37, 590-602	1.5	37	
155	Implementation of Electronic Consent at a Biobank: An Opportunity for Precision Medicine Research. <i>Journal of Personalized Medicine</i> , 2016 , 6,	3.6	37	
154	Incident user cohort study of risk for gastrointestinal bleed and stroke in individuals with major depressive disorder treated with antidepressants. <i>BMJ Open</i> , 2012 , 2, e000544	3	36	
153	Prevalence and correlates of panic attacks in postmenopausal women: results from an ancillary study to the Womenß Health Initiative. <i>Archives of Internal Medicine</i> , 2003 , 163, 2041-50		36	
152	Mapping genomic loci implicates genes and synaptic biology in schizophrenia Nature, 2022,	50.4	35	
151	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	
150	Bringing a developmental perspective to anxiety genetics. <i>Development and Psychopathology</i> , 2012 , 24, 1179-93	4.3	34	
149	Limitations of Contemporary Guidelines for Managing Patients at High Genetic Risk of Coronary Artery Disease. <i>Journal of the American College of Cardiology</i> , 2020 , 75, 2769-2780	15.1	33	
148	Limbic system white matter microstructure and long-term treatment outcome in major depressive disorder: a diffusion tensor imaging study using legacy data. <i>World Journal of Biological Psychiatry</i> , 2014 , 15, 122-34	3.8	33	
147	Genome-wide association study identifies 48 common genetic variants associated with handedness. Nature Human Behaviour, 2021 , 5, 59-70	12.8	33	

146	Early Childhood IQ Trajectories in Individuals Later Developing Schizophrenia and Affective Psychoses in the New England Family Studies. <i>Schizophrenia Bulletin</i> , 2015 , 41, 817-23	1.3	32
145	Genetic predictors of risk and resilience in psychiatric disorders: a cross-disorder genome-wide association study of functional impairment in major depressive disorder, bipolar disorder, and schizophrenia. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2013 , 162B, 779-8	3.5 88	32
144	Ethical Considerations Related to Return of Results from Genomic Medicine Projects: The eMERGE Network (Phase III) Experience. <i>Journal of Personalized Medicine</i> , 2018 , 8,	3.6	32
143	Panic disorder and serotonergic genes (SLC6A4, HTR1A and HTR2A): Association and interaction with childhood trauma and parenting. <i>Neuroscience Letters</i> , 2010 , 485, 11-5	3.3	31
142	Genetics of anxiety: would the genome recognize the DSM?. Depression and Anxiety, 2008, 25, 368-77	8.4	31
141	Greater male than female variability in regional brain structure across the lifespan. <i>Human Brain Mapping</i> , 2020 ,	5.9	31
140	Genetic risk variants for social anxiety. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017 , 174, 120-131	3.5	30
139	Implications of comorbidity and ascertainment bias for identifying disease genes. <i>American Journal of Medical Genetics Part A</i> , 2000 , 96, 817-22		30
138	Morphometricity as a measure of the neuroanatomical signature of a trait. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, E5749-56	11.5	30
137	Genomics and psychological resilience: a research agenda. <i>Molecular Psychiatry</i> , 2019 , 24, 1770-1778	15.1	28
136	Influence of RGS2 on sertraline treatment for social anxiety disorder. <i>Neuropsychopharmacology</i> , 2014 , 39, 1340-6	8.7	28
135	Genome-wide association study of generalized anxiety symptoms in the Hispanic Community Health Study/Study of Latinos. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017 , 174, 132-143	3.5	27
134	Cortical thickness across the lifespan: Data from 17,075 healthy individuals aged 3-90 years. <i>Human Brain Mapping</i> , 2021 ,	5.9	26
133	Screening pregnant women for suicidal behavior in electronic medical records: diagnostic codes vs. clinical notes processed by natural language processing. <i>BMC Medical Informatics and Decision Making</i> , 2018 , 18, 30	3.6	25
132	Frontal P3 event-related potential is related to brain glutamine/glutamate ratio measured in vivo. NeuroImage, 2015 , 111, 186-91	7.9	24
131	Does developmental timing of exposure to child maltreatment predict memory performance in adulthood? Results from a large, population-based sample. <i>Child Abuse and Neglect</i> , 2016 , 51, 181-91	4.3	24
130	Precision Psychiatry-Will Genomic Medicine Lead the Way?. JAMA Psychiatry, 2018, 75, 663-664	14.5	24
129	Patterns of deficits in brain function in bipolar disorder and schizophrenia: a cluster analytic study. <i>Psychiatry Research</i> , 2012 , 200, 272-80	9.9	24

(2007-2019)

128	Genetic overlap between vascular pathologies and Alzheimerß dementia and potential causal mechanisms. <i>Alzheimermand Dementia</i> , 2019 , 15, 65-75	1.2	24	
127	Investigation of parent-of-origin effects in ADHD candidate genes. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2007 , 144B, 776-80	3.5	23	
126	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	
125	Psychiatric Genetics Begins to Find Its Footing. <i>American Journal of Psychiatry</i> , 2019 , 176, 609-614	11.9	22	
124	The genetics of panic disorder. Current Psychiatry Reports, 2001, 3, 131-7	9.1	22	
123	An eMERGE Clinical Center at Partners Personalized Medicine. <i>Journal of Personalized Medicine</i> , 2016 , 6,	3.6	22	
122	Validation of an Electronic Health Record-Based Suicide Risk Prediction Modeling Approach Across Multiple Health Care Systems. <i>JAMA Network Open</i> , 2020 , 3, e201262	10.4	21	
121	Statistical power and utility of meta-analysis methods for cross-phenotype genome-wide association studies. <i>PLoS ONE</i> , 2018 , 13, e0193256	3.7	21	
120	Analysis of Genetically Regulated Gene Expression Identifies a Prefrontal PTSD Gene, SNRNP35, Specific to Military Cohorts. <i>Cell Reports</i> , 2020 , 31, 107716	10.6	21	
119	Genomic influences on self-reported childhood maltreatment. <i>Translational Psychiatry</i> , 2020 , 10, 38	8.6	20	
118	Cross-Phenotype Polygenic Risk Score Analysis of Persistent Post-Concussive Symptoms in U.S. Army Soldiers with Deployment-Acquired Traumatic Brain Injury. <i>Journal of Neurotrauma</i> , 2017 , 34, 781	- 7 8 1 9	20	
117	Dissociable genetic contributions to error processing: a multimodal neuroimaging study. <i>PLoS ONE</i> , 2014 , 9, e101784	3.7	20	
116	Physical activity offsets genetic risk for incident depression assessed via electronic health records in a biobank cohort study. <i>Depression and Anxiety</i> , 2020 , 37, 106-114	8.4	20	
115	Contribution of Rare Copy Number Variants to Bipolar Disorder Risk Is Limited to Schizoaffective Cases. <i>Biological Psychiatry</i> , 2019 , 86, 110-119	7.9	20	
114	Emotion sensitivity and self-reported symptoms of generalized anxiety disorder across the lifespan: A population-based sample approach. <i>Brain and Behavior</i> , 2019 , 9, e01282	3.4	19	
113	Mega-analysis methods in ENIGMA: The experience of the generalized anxiety disorder working group. <i>Human Brain Mapping</i> , 2020 ,	5.9	19	
112	Modifiers and subtype-specific analyses in whole-genome association studies: a likelihood framework. <i>Human Heredity</i> , 2011 , 72, 10-20	1.1	19	
111	Genetics of bipolar disorder. <i>Current Psychiatry Reports</i> , 2007 , 9, 504-11	9.1	19	

110	Adverse obstetric outcomes during delivery hospitalizations complicated by suicidal behavior among US pregnant women. <i>PLoS ONE</i> , 2018 , 13, e0192943	3.7	19
109	Conditional GWAS analysis to identify disorder-specific SNPs for psychiatric disorders. <i>Molecular Psychiatry</i> , 2021 , 26, 2070-2081	15.1	19
108	Glycobiology and schizophrenia: a biological hypothesis emerging from genomic research. <i>Molecular Psychiatry</i> , 2020 , 25, 3129-3139	15.1	18
107	Aberrant glycosylation in schizophrenia: a review of 25 years of post-mortem brain studies. <i>Molecular Psychiatry</i> , 2020 , 25, 3198-3207	15.1	18
106	Using Administrative Data to Predict Suicide After Psychiatric Hospitalization in the Veterans Health Administration System. <i>Frontiers in Psychiatry</i> , 2020 , 11, 390	5	18
105	A kernel machine method for detecting effects of interaction between multidimensional variable sets: an imaging genetics application. <i>Neurolmage</i> , 2015 , 109, 505-514	7.9	18
104	The schizophrenia risk locus in SLC39A8 alters brain metal transport and plasma glycosylation. <i>Scientific Reports</i> , 2020 , 10, 13162	4.9	18
103	Common Measures for National Institute of Mental Health Funded Research. <i>Biological Psychiatry</i> , 2016 , 79, e91-6	7.9	18
102	Using life history calendars to improve measurement of lifetime experience with mental disorders. <i>Psychological Medicine</i> , 2020 , 50, 515-522	6.9	18
101	A Comparison of Ten Polygenic Score Methods for Psychiatric Disorders Applied Across Multiple Cohorts. <i>Biological Psychiatry</i> , 2021 , 90, 611-620	7.9	17
100	Hematopoietic mosaic chromosomal alterations increase the risk for diverse types of infection. <i>Nature Medicine</i> , 2021 , 27, 1012-1024	50.5	16
99	Use of natural language processing in electronic medical records to identify pregnant women with suicidal behavior: towards a solution to the complex classification problem. <i>European Journal of Epidemiology</i> , 2019 , 34, 153-162	12.1	16
98	Rare coding variants in ten genes confer substantial risk for schizophrenia Nature, 2022,	50.4	16
97	Genetic validation of bipolar disorder identified by automated phenotyping using electronic health records. <i>Translational Psychiatry</i> , 2018 , 8, 86	8.6	14
96	Polygenic Risk for Depression Increases Risk of Ischemic Stroke: From the Stroke Genetics Network Study. <i>Stroke</i> , 2018 , 49, 543-548	6.7	14
95	The Genetics of Endophenotypes of Neurofunction to Understand Schizophrenia (GENUS) consortium: A collaborative cognitive and neuroimaging genetics project. <i>Schizophrenia Research</i> , 2018 , 195, 306-317	3.6	14
94	ENIGMA-anxiety working group: Rationale for and organization of large-scale neuroimaging studies of anxiety disorders. <i>Human Brain Mapping</i> , 2020 ,	5.9	14
93	Pleiotropy and Cross-Disorder Genetics Among Psychiatric Disorders. <i>Biological Psychiatry</i> , 2021 , 89, 20-31	7.9	14

(2008-2016)

92	Attention Bias to Emotional Faces Varies by IQ and Anxiety in Williams Syndrome. <i>Journal of Autism and Developmental Disorders</i> , 2016 , 46, 2174-2185	4.6	13
91	Psychiatric genetics and the future of personalized treatment. <i>Depression and Anxiety</i> , 2014 , 31, 893-8	8.4	13
90	Subcortical volumes across the lifespan: Data from 18,605 healthy individuals aged 3-90 years. <i>Human Brain Mapping</i> , 2021 ,	5.9	13
89	Feasibility of studying brain morphology in major depressive disorder with structural magnetic resonance imaging and clinical data from the electronic medical record: a pilot study. <i>Psychiatry Research - Neuroimaging</i> , 2013 , 211, 202-13	2.9	12
88	Preliminary evidence of association between EFHC2, a gene implicated in fear recognition, and harm avoidance. <i>Neuroscience Letters</i> , 2009 , 452, 84-6	3.3	12
87	Minimal phenotyping yields GWAS hits of reduced specificity for major depression		12
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41	Testing Causal Bidirectional Influences between Physical Activity and Depression using Mendelian Rand	domiza	ntion
40	Examining sex-differentiated genetic effects across neuropsychiatric and behavioral traits		3
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19	Pleiotropy in complex traits: challenges and strategies		1
18	Analysis of Genetically Regulated Gene Expression identifies a trauma type specific PTSD gene, SNRNP3	5	1
17	A two-stage approach to identifying and validating modifiable factors for the prevention of depression		1
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