

Jordan W Smoller

List of Publications by Year in Descending Order

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Version: 2024-04-17

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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.

253
papers

24,147
citations

65
h-index

153
g-index

285
ext. papers

32,954
ext. citations

10
avg, IF

6.75
L-index

#	Paper	IF	Citations
253	Mammalian brain glycoproteins exhibit diminished glycan complexity compared to other tissues.. <i>Nature Communications</i> , 2022 , 13, 275	17.4	2
252	A framework for employing longitudinally collected multicenter electronic health records to stratify heterogeneous patient populations on disease history.. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2022 ,	8.6	1
251	Prediction of Suicide Attempts Using Clinician Assessment, Patient Self-report, and Electronic Health Records.. <i>JAMA Network Open</i> , 2022 , 5, e2144373	10.4	4
250	Predictive structured-unstructured interactions in EHR models: A case study of suicide prediction.. <i>Npj Digital Medicine</i> , 2022 , 5, 15	15.7	0
249	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. <i>Biological Psychiatry</i> , 2022 , 91, 102-117	7.9	11
248	Low depression frequency is associated with decreased risk of cardiometabolic disease 2022 , 1, 125-131		1
247	Implementing Machine Learning Models for Suicide Risk Prediction in Clinical Practice: Focus Group Study With Hospital Providers.. <i>JMIR Formative Research</i> , 2022 , 6, e30946	2.5	1
246	Characterizing mood disorders in the AFFECT study: a large, longitudinal, and phenotypically rich genetic cohort in the US.. <i>Translational Psychiatry</i> , 2022 , 12, 121	8.6	0
245	Mapping genomic loci implicates genes and synaptic biology in schizophrenia.. <i>Nature</i> , 2022 ,	50.4	35
244	Rare coding variants in ten genes confer substantial risk for schizophrenia.. <i>Nature</i> , 2022 ,	50.4	16
243	Large-scale real-world data analysis identifies comorbidity patterns in schizophrenia.. <i>Translational Psychiatry</i> , 2022 , 12, 154	8.6	
242	Genetic architecture of 11 major psychiatric disorders at biobehavioral, functional genomic and molecular genetic levels of analysis.. <i>Nature Genetics</i> , 2022 , 54, 548-559	36.3	4
241	Psychiatric manifestations of rare variation in medically actionable genes: a PheWAS approach.. <i>BMC Genomics</i> , 2022 , 23, 385	4.5	
240	Examining the impact of the COVID-19 pandemic through the lens of the network approach to psychopathology: Analysis of the Brazilian Longitudinal Study of Health (ELSA-Brasil) cohort over a 12-year timespan.. <i>Journal of Anxiety Disorders</i> , 2021 , 85, 102512	10.9	2
239	Use of the PsycheMERGE Network to Investigate the Association Between Depression Polygenic Scores and White Blood Cell Count. <i>JAMA Psychiatry</i> , 2021 , 78, 1365-1374	14.5	3
238	Returning actionable genomic results in a research biobank: Analytic validity, clinical implementation, and resource utilization. <i>American Journal of Human Genetics</i> , 2021 , 108, 2224-2237	11	3
237	Temporally informed random forests for suicide risk prediction. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2021 ,	8.6	1

236	An independently validated, portable algorithm for the rapid identification of COPD patients using electronic health records. <i>Scientific Reports</i> , 2021 , 11, 19959	4.9	1
235	Testing the Stability and Validity of an Executive Dysfunction Classification Using Task-Based Assessment in Children and Adolescents. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2021 , 60, 1501-1512	7.2	0
234	Conditional GWAS analysis to identify disorder-specific SNPs for psychiatric disorders. <i>Molecular Psychiatry</i> , 2021 , 26, 2070-2081	15.1	19
233	Socio-demographic and trauma-related predictors of PTSD within 8 weeks of a motor vehicle collision in the AURORA study. <i>Molecular Psychiatry</i> , 2021 , 26, 3108-3121	15.1	6
232	Increase in Suicidal Thinking During COVID-19. <i>Clinical Psychological Science</i> , 2021 , 9, 482-488	6	12
231	Overlap in genetic risk for cross-disorder vulnerability to mental disorders and genetic risk for altered subcortical brain volumes. <i>Journal of Affective Disorders</i> , 2021 , 282, 740-756	6.6	2
230	Prevalence and risk factors of psychiatric symptoms and diagnoses before and during the COVID-19 pandemic: findings from the ELSA-Brasil COVID-19 mental health cohort. <i>Psychological Medicine</i> , 2021 , 1-12	6.9	12
229	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
228	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
227	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
226	Reconciling Statistical and Clinicians' Predictions of Suicide Risk. <i>Psychiatric Services</i> , 2021 , 72, 555-562	3.3	6
225	The association between genetically determined ABO blood types and major depressive disorder. <i>Psychiatry Research</i> , 2021 , 299, 113837	9.9	
224	Examining Sex-Differentiated Genetic Effects Across Neuropsychiatric and Behavioral Traits. <i>Biological Psychiatry</i> , 2021 , 89, 1127-1137	7.9	12
223	Accuracy Requirements for Cost-effective Suicide Risk Prediction Among Primary Care Patients in the US. <i>JAMA Psychiatry</i> , 2021 , 78, 642-650	14.5	10
222	Hematopoietic mosaic chromosomal alterations increase the risk for diverse types of infection. <i>Nature Medicine</i> , 2021 , 27, 1012-1024	50.5	16
221	Hypertension prevalence in the All of Us Research Program among groups traditionally underrepresented in medical research. <i>Scientific Reports</i> , 2021 , 11, 12849	4.9	1
220	Genome-wide association study identifies 48 common genetic variants associated with handedness. <i>Nature Human Behaviour</i> , 2021 , 5, 59-70	12.8	33
219	Pleiotropy and Cross-Disorder Genetics Among Psychiatric Disorders. <i>Biological Psychiatry</i> , 2021 , 89, 20-31	7.9	14

218	Evidence in the UK Biobank for the underdiagnosis of erythropoietic protoporphyria. <i>Genetics in Medicine</i> , 2021 , 23, 140-148	8.1	5
217	Lossless integration of multiple electronic health records for identifying pleiotropy using summary statistics. <i>Nature Communications</i> , 2021 , 12, 168	17.4	1
216	Clinical laboratory test-wide association scan of polygenic scores identifies biomarkers of complex disease. <i>Genome Medicine</i> , 2021 , 13, 6	14.4	11
215	Cortical thickness across the lifespan: Data from 17,075 healthy individuals aged 3-90 years. <i>Human Brain Mapping</i> , 2021 ,	5.9	26
214	Subcortical volumes across the lifespan: Data from 18,605 healthy individuals aged 3-90 years. <i>Human Brain Mapping</i> , 2021 ,	5.9	13
213	Characterisation of age and polarity at onset in bipolar disorder.. <i>British Journal of Psychiatry</i> , 2021 , 219, 659-669	5.4	2
212	Generalizability of Polygenic Risk Scores for Breast Cancer Among Women With European, African, and Latinx Ancestry. <i>JAMA Network Open</i> , 2021 , 4, e2119084	10.4	5
211	Investigating Shared Genetic Basis Across Tourette Syndrome and Comorbid Neurodevelopmental Disorders Along the Impulsivity-Compulsivity Spectrum. <i>Biological Psychiatry</i> , 2021 , 90, 317-327	7.9	12
210	Transcriptome-based polygenic score links depression-related corticolimbic gene expression changes to sex-specific brain morphology and depression risk. <i>Neuropsychopharmacology</i> , 2021 , 46, 2304-2311 ¹	8.7	11
209	The Genetic Architecture of Depression in Individuals of East Asian Ancestry: A Genome-Wide Association Study. <i>JAMA Psychiatry</i> , 2021 , 78, 1258-1269	14.5	7
208	Cross-Disorder Genomics Data Analysis Elucidates a Shared Genetic Basis Between Major Depression and Osteoarthritis Pain. <i>Frontiers in Genetics</i> , 2021 , 12, 687687	4.5	3
207	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. <i>Biological Psychiatry</i> , 2021 ,	7.9	11
206	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
205	Cortical and subcortical brain structure in generalized anxiety disorder: findings from 28 research sites in the ENIGMA-Anxiety Working Group. <i>Translational Psychiatry</i> , 2021 , 11, 502	8.6	4
204	Plasma metabolomic profiles associated with chronic distress in women. <i>Psychoneuroendocrinology</i> , 2021 , 133, 105420	5	0
203	Geographic Variation in Obesity at the State Level in the All of Us Research Program.. <i>Preventing Chronic Disease</i> , 2021 , 18, E104	3.7	2
202	Glycobiology and schizophrenia: a biological hypothesis emerging from genomic research. <i>Molecular Psychiatry</i> , 2020 , 25, 3129-3139	15.1	18
201	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

200	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
199	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
198	Resilience to mental disorders in a low-income, non-Westernized setting. <i>Psychological Medicine</i> , 2020 , 1-10	6.9	6
197	The genetic architecture of the human cerebral cortex. <i>Science</i> , 2020 , 367,	33.3	156
196	Minimal phenotyping yields genome-wide association signals of low specificity for major depression. <i>Nature Genetics</i> , 2020 , 52, 437-447	36.3	80
195	Mega-analysis methods in ENIGMA: The experience of the generalized anxiety disorder working group. <i>Human Brain Mapping</i> , 2020 ,	5.9	19
194	Making the right moves to prevent depression in young people. <i>Lancet Psychiatry</i> , 2020 , 7, 221-222	23.3	1
193	Genomic influences on self-reported childhood maltreatment. <i>Translational Psychiatry</i> , 2020 , 10, 38	8.6	20
192	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
191	Maternal vitamin D status during pregnancy and offspring risk of childhood/adolescent depression: Results from the Avon Longitudinal Study of Parents and Children (ALSPAC). <i>Journal of Affective Disorders</i> , 2020 , 265, 255-262	6.6	1
190	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
189	Hematopoietic mosaic chromosomal alterations and risk for infection among 767,891 individuals without blood cancer 2020 ,		5
188	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
187	Genome-wide association study of shared liability to anxiety disorders in Army STARRS. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2020 , 183, 197-207	3.5	5
186	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
185	Molecular genetic overlap between posttraumatic stress disorder and sleep phenotypes. <i>Sleep</i> , 2020 , 43,	1.1	9
184	Greater male than female variability in regional brain structure across the lifespan. <i>Human Brain Mapping</i> , 2020 ,	5.9	31
183	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

182	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
181	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
180	The schizophrenia risk locus in SLC39A8 alters brain metal transport and plasma glycosylation. <i>Scientific Reports</i> , 2020 , 10, 13162	4.9	18
179	N-glycome analysis detects dysglycosylation missed by conventional methods in SLC39A8 deficiency. <i>Journal of Inherited Metabolic Disease</i> , 2020 , 43, 1370-1381	5.4	4
178	Prospective study of polygenic risk, protective factors, and incident depression following combat deployment in US Army soldiers. <i>Psychological Medicine</i> , 2020 , 50, 737-745	6.9	11
177	Using life history calendars to improve measurement of lifetime experience with mental disorders. <i>Psychological Medicine</i> , 2020 , 50, 515-522	6.9	18
176	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
175	Translating Discoveries in Attention-Deficit/Hyperactivity Disorder Genomics to an Outpatient Child and Adolescent Psychiatric Cohort. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2020 , 59, 964-977	7.2	11
174	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. <i>Biological Psychiatry</i> , 2020 , 87, 419-430	7.9	9
173	Harmonizing Clinical Sequencing and Interpretation for the eMERGE III Network. <i>American Journal of Human Genetics</i> , 2019 , 105, 588-605	11	63
172	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
171	Genetic associations with suicide attempt severity and genetic overlap with major depression. <i>Translational Psychiatry</i> , 2019 , 9, 22	8.6	48
170	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
169	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
168	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
167	Polygenic prediction via Bayesian regression and continuous shrinkage priors. <i>Nature Communications</i> , 2019 , 10, 1776	17.4	191
166	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019 , 51, 793-803	36.3	662
165	Interrogating the Genetic Determinants of Tourette's Syndrome and Other Tic Disorders Through Genome-Wide Association Studies. <i>American Journal of Psychiatry</i> , 2019 , 176, 217-227	11.9	95

164	Associations of antepartum suicidal behaviour with adverse infant and obstetric outcomes. <i>Paediatric and Perinatal Epidemiology</i> , 2019 , 33, 137-144	2.7	1
163	The "All of Us" Research Program. <i>New England Journal of Medicine</i> , 2019 , 381, 668-676	59.2	325
162	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
161	Psychiatric Genetics Begins to Find Its Footing. <i>American Journal of Psychiatry</i> , 2019 , 176, 609-614	11.9	22
160	Genomics and psychological resilience: a research agenda. <i>Molecular Psychiatry</i> , 2019 , 24, 1770-1778	15.1	28
159	Genome-wide Association Studies in Ancestrally Diverse Populations: Opportunities, Methods, Pitfalls, and Recommendations. <i>Cell</i> , 2019 , 179, 589-603	56.2	145
158	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
157	Detecting potential pleiotropy across cardiovascular and neurological diseases using univariate, bivariate, and multivariate methods on 43,870 individuals from the eMERGE network. <i>Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing</i> , 2019 , 24, 272-283	1.3	6
156	Genetic architecture of subcortical brain structures in 38,851 individuals. <i>Nature Genetics</i> , 2019 , 51, 1624-1636	36.36	81
155	Genetic overlap between vascular pathologies and Alzheimer's dementia and potential causal mechanisms. <i>Alzheimer's and Dementia</i> , 2019 , 15, 65-75	1.2	24
154	GWAS Identifies Risk Locus for Erectile Dysfunction and Implicates Hypothalamic Neurobiology and Diabetes in Etiology. <i>American Journal of Human Genetics</i> , 2019 , 104, 157-163	11	12
153	Joint Contributions of Rare Copy Number Variants and Common SNPs to Risk for Schizophrenia. <i>American Journal of Psychiatry</i> , 2019 , 176, 29-35	11.9	59
152	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
151	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
150	The Shared Genetic Basis of Educational Attainment and Cerebral Cortical Morphology. <i>Cerebral Cortex</i> , 2019 , 29, 3471-3481	5.1	11
149	Psychiatric genetics and the structure of psychopathology. <i>Molecular Psychiatry</i> , 2019 , 24, 409-420	15.1	149
148	Genome-wide analysis of insomnia disorder. <i>Molecular Psychiatry</i> , 2018 , 23, 2238-2250	15.1	39
147	Genetic validation of bipolar disorder identified by automated phenotyping using electronic health records. <i>Translational Psychiatry</i> , 2018 , 8, 86	8.6	14

146	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
145	Reply to Risk and Zhu: Mixed-effects modeling as a principled approach to heritability analysis with repeat measurements. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018 , 115, E123	11.5	
144	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
143	Dissociable influences of β and polygenic risk of AD dementia on amyloid and cognition. <i>Neurology</i> , 2018 , 90, e1605-e1612	6.5	43
142	The human cortex possesses a reconfigurable dynamic network architecture that is disrupted in psychosis. <i>Nature Communications</i> , 2018 , 9, 1157	17.4	42
141	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
140	Psychiatric Genomics: An Update and an Agenda. <i>American Journal of Psychiatry</i> , 2018 , 175, 15-27	11.9	328
139	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , 2018 , 360,	33.3	666
138	Precision Psychiatry-Will Genomic Medicine Lead the Way?. <i>JAMA Psychiatry</i> , 2018 , 75, 663-664	14.5	24
137	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
136	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
135	Adverse obstetric outcomes during delivery hospitalizations complicated by suicidal behavior among US pregnant women. <i>PLoS ONE</i> , 2018 , 13, e0192943	3.7	19
134	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
133	Genome-wide association study of depressive symptoms in the Hispanic Community Health Study/Study of Latinos. <i>Journal of Psychiatric Research</i> , 2018 , 99, 167-176	5.2	9
132	Precision Psychiatry-Yes, But Precisely What?-Reply. <i>JAMA Psychiatry</i> , 2018 , 75, 1303	14.5	2
131	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
130	ENIGMA and the individual: Predicting factors that affect the brain in 35 countries worldwide. <i>NeuroImage</i> , 2017 , 145, 389-408	7.9	142
129	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

128	Novel genetic loci associated with hippocampal volume. <i>Nature Communications</i> , 2017 , 8, 13624	17.4	173
127	Genetic risk variants for social anxiety. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017 , 174, 120-131	3.5	30
126	Anxiety genetics: Dispatches from the frontier. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017 , 174, 117-119	3.5	
125	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
124	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
123	The interplay of childhood behavior problems and IQ in the development of later schizophrenia and affective psychoses. <i>Schizophrenia Research</i> , 2017 , 184, 45-51	3.6	9
122	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
121	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, 781-789	5.4	20
120	Predicting Suicidal Behavior From Longitudinal Electronic Health Records. <i>American Journal of Psychiatry</i> , 2017 , 174, 154-162	11.9	156
119	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
118	A Quarter Century of Progress in Psychiatric Genetics. <i>Harvard Review of Psychiatry</i> , 2017 , 25, 256-258	4.1	7
117	Phenome-wide heritability analysis of the UK Biobank. <i>PLoS Genetics</i> , 2017 , 13, e1006711	6	131
116	Stratifying Risk for Renal Insufficiency Among Lithium-Treated Patients: An Electronic Health Record Study. <i>Neuropsychopharmacology</i> , 2016 , 41, 1138-43	8.7	49
115	Multidimensional heritability analysis of neuroanatomical shape. <i>Nature Communications</i> , 2016 , 7, 13291	17.4	38
114	Polygenic risk of Alzheimer disease is associated with early- and late-life processes. <i>Neurology</i> , 2016 , 87, 481-8	6.5	86
113	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
112	Genetic influences on schizophrenia and subcortical brain volumes: large-scale proof of concept. <i>Nature Neuroscience</i> , 2016 , 19, 420-431	25.5	163
111	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

110	The Genetics of Stress-Related Disorders: PTSD, Depression, and Anxiety Disorders. <i>Neuropsychopharmacology</i> , 2016 , 41, 297-319	8.7	220
109	An eMERGE Clinical Center at Partners Personalized Medicine. <i>Journal of Personalized Medicine</i> , 2016 , 6,	3.6	22
108	Implementation of Electronic Consent at a Biobank: An Opportunity for Precision Medicine Research. <i>Journal of Personalized Medicine</i> , 2016 , 6,	3.6	37
107	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
106	Common Measures for National Institute of Mental Health Funded Research. <i>Biological Psychiatry</i> , 2016 , 79, e91-6	7.9	18
105	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
104	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
103	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
102	Novel genetic loci underlying human intracranial volume identified through genome-wide association. <i>Nature Neuroscience</i> , 2016 , 19, 1569-1582	25.5	147
101	Nicotine dependence and psychosis in Bipolar disorder and Schizoaffective disorder, Bipolar type. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2016 , 171, 521-4	3.5	3
100	Common genetic variants influence human subcortical brain structures. <i>Nature</i> , 2015 , 520, 224-9	50.4	601
99	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
98	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
97	Frontal P3 event-related potential is related to brain glutamine/glutamate ratio measured in vivo. <i>NeuroImage</i> , 2015 , 111, 186-91	7.9	24
96	Individual Aesthetic Preferences for Faces Are Shaped Mostly by Environments, Not Genes. <i>Current Biology</i> , 2015 , 25, 2684-9	6.3	60
95	Folate, vitamin B-6, and vitamin B-12 intake and mild cognitive impairment and probable dementia in the Women's Health Initiative Memory Study. <i>Journal of the Academy of Nutrition and Dietetics</i> , 2015 , 115, 231-241	3.9	51
94	Brain Genomics Superstruct Project initial data release with structural, functional, and behavioral measures. <i>Scientific Data</i> , 2015 , 2, 150031	8.2	204
93	Prevalence of Huntington's disease gene CAG trinucleotide repeat alleles in patients with bipolar disorder. <i>Bipolar Disorders</i> , 2015 , 17, 403-8	3.8	5

92	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
91	Genetic determinants of depression: recent findings and future directions. <i>Harvard Review of Psychiatry</i> , 2015 , 23, 1-18	4.1	102
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16	The restricted nature of protein glycosylation in the mammalian brain		1
15	Findings and insights from the genetic investigation of age of first reported occurrence for complex disorders in the UK Biobank and FinnGen		3
14	Global biobank analyses provide lessons for computing polygenic risk scores across diverse cohorts		3
13	Testing Causal Bidirectional Influences between Physical Activity and Depression using Mendelian Randomization		
12	The schizophrenia-associated variant in SLC39A8 alters N-glycosylation in the mouse brain		1
11	Pleiotropy in complex traits: challenges and strategies		1
10	Examining sex-differentiated genetic effects across neuropsychiatric and behavioral traits		3
9	Cortical Thickness Trajectories across the Lifespan: Data from 17,075 healthy individuals aged 3-90 years		4
8	Genetic Architecture of 11 Major Psychiatric Disorders at Biobehavioral, Functional Genomic, and Molecular Genetic Levels of Analysis		4
7	Minimal phenotyping yields GWAS hits of reduced specificity for major depression		12
6	Genome wide meta-analysis identifies genomic relationships, novel loci, and pleiotropic mechanisms across eight psychiatric disorders		10
5	Analysis of Genetically Regulated Gene Expression identifies a trauma type specific PTSD gene, SNRNP35		1
4	The schizophrenia risk locus in SLC39A8 alters brain metal transport and plasma glycosylation		6
3	A two-stage approach to identifying and validating modifiable factors for the prevention of depression		1

2	Cross-disorder GWAS meta-analysis for Attention Deficit/Hyperactivity Disorder, Autism Spectrum Disorder, Obsessive Compulsive Disorder, and Tourette Syndrome	2
1	Characterization of Age and Polarity at Onset in Bipolar Disorder	1