

A mega-analysis of genome-wide association studies for

Molecular Psychiatry

18, 497-511

DOI: [10.1038/mp.2012.21](https://doi.org/10.1038/mp.2012.21)

Citation Report

#	ARTICLE	IF	CITATIONS
1	Mental Illness and the Criminal Justice System. , 2009, , 478-498.		11
2	A genome-wide association study on common SNPs and rare CNVs in anorexia nervosa. <i>Molecular Psychiatry</i> , 2011, 16, 949-959.	4.1	186
3	Revolution Stalled. <i>Science Translational Medicine</i> , 2012, 4, 155cm11.	5.8	207
4	Multi-locus genome-wide association analysis supports the role of glutamatergic synaptic transmission in the etiology of major depressive disorder. <i>Translational Psychiatry</i> , 2012, 2, e184-e184.	2.4	77
5	The Genetic Basis of Depression. <i>Current Topics in Behavioral Neurosciences</i> , 2012, 14, 81-99.	0.8	8
6	Bringing a developmental perspective to anxiety genetics. <i>Development and Psychopathology</i> , 2012, 24, 1179-1193.	1.4	40
7	Individual Differences in Amygdala-Medial Prefrontal Anatomy Link Negative Affect, Impaired Social Functioning, and Polygenic Depression Risk. <i>Journal of Neuroscience</i> , 2012, 32, 18087-18100.	1.7	250
8	Systems Biology, Bioinformatics, and Biomarkers in Neuropsychiatry. <i>Frontiers in Neuroscience</i> , 2012, 6, 187.	1.4	41
10	The genetic basis of mood and anxiety disorders – changing paradigms. <i>Biology of Mood & Anxiety Disorders</i> , 2012, 2, 17.	4.7	17
11	Data-driven subtypes of major depressive disorder: a systematic review. <i>BMC Medicine</i> , 2012, 10, 156.	2.3	229
12	Using summary data from the Danish National Registers to estimate heritabilities for schizophrenia, bipolar disorder, and major depressive disorder. <i>Frontiers in Genetics</i> , 2012, 3, 118.	1.1	176
13	Genetic architectures of psychiatric disorders: the emerging picture and its implications. <i>Nature Reviews Genetics</i> , 2012, 13, 537-551.	7.7	1,025
14	Gene expression profiles associated with depression in patients with chronic hepatitis C (CH-C). <i>Brain and Behavior</i> , 2012, 2, 525-531.	1.0	12
15	Considering trauma exposure in the context of genetics studies of posttraumatic stress disorder: a systematic review. <i>Biology of Mood & Anxiety Disorders</i> , 2013, 3, 2.	4.7	30
16	Cardiovascular Disease, Psychosocial Factors, and Genetics: The Case of Depression. <i>Progress in Cardiovascular Diseases</i> , 2013, 55, 557-562.	1.6	42
17	Understanding the somatic consequences of depression: biological mechanisms and the role of depression symptom profile. <i>BMC Medicine</i> , 2013, 11, 129.	2.3	550
18	Genetic association studies between SNPs and suicidal behavior: A meta-analytical field synopsis. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2013, 46, 36-42.	2.5	34
19	Genome-wide association analysis identifies 13 new risk loci for schizophrenia. <i>Nature Genetics</i> , 2013, 45, 1150-1159.	9.4	1,395

#	ARTICLE	IF	CITATIONS
20	p11 and its role in depression and therapeutic responses to antidepressants. <i>Nature Reviews Neuroscience</i> , 2013, 14, 673-680.	4.9	144
21	On schizophrenia as a "disease of humanity". <i>Schizophrenia Research</i> , 2013, 143, 223-224.	1.1	6
22	Interactive effects of corticotropin-releasing hormone receptor 1 gene and childhood adversity on depressive symptoms in young adults: Findings from a longitudinal study. <i>European Neuropsychopharmacology</i> , 2013, 23, 358-367.	0.3	43
23	Genome-wide scan of job-related exhaustion with three replication studies implicate a susceptibility variant at the UST gene locus. <i>Human Molecular Genetics</i> , 2013, 22, 3363-3372.	1.4	13
24	Circadian clocks, brain function, and development. <i>Annals of the New York Academy of Sciences</i> , 2013, 1306, 43-67.	1.8	36
26	Using Phenotypic Heterogeneity to Increase the Power of Genome-Wide Association Studies: Application to Age at Onset of Ischaemic Stroke Subphenotypes. <i>Genetic Epidemiology</i> , 2013, 37, 495-503.	0.6	10
27	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. <i>Nature Genetics</i> , 2013, 45, 984-994.	9.4	2,067
28	Protocol for a collaborative meta-analysis of 5-HTTLPR, stress, and depression. <i>BMC Psychiatry</i> , 2013, 13, 304.	1.1	35
29	Association Between Autozygosity and Major Depression: Stratification Due to Religious Assortment. <i>Behavior Genetics</i> , 2013, 43, 455-467.	1.4	34
30	Questions about DISC1 as a genetic risk factor for schizophrenia. <i>Molecular Psychiatry</i> , 2013, 18, 1050-1052.	4.1	86
31	A network medicine approach to psychiatric genetics. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2013, 162, 579-586.	1.1	2
32	Cross-species genetics converge to <i>TLL2</i> for mouse avoidance behavior and human bipolar disorder. <i>Genes, Brain and Behavior</i> , 2013, 12, 653-657.	1.1	9
33	Genome-wide association analysis accounting for environmental factors through propensity score matching: Application to stressful life events in major depressive disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2013, 162, 521-529.	1.1	16
35	Recurrent deletions of <i>ULK4</i> in schizophrenia: a novel gene crucial for neuritogenesis and neuronal motility. <i>Journal of Cell Science</i> , 2014, 127, 630-40.	1.2	78
36	Biological substrates underpinning diagnosis of major depression. <i>International Journal of Neuropsychopharmacology</i> , 2013, 16, 1893-1909.	1.0	33
37	The genetic overlap between schizophrenia and height. <i>Schizophrenia Research</i> , 2013, 151, 226-228.	1.1	12
38	Contextualizing experience. <i>Developmental Review</i> , 2013, 33, 273-278.	2.6	3
39	Identification of risk loci with shared effects on five major psychiatric disorders: a genome-wide analysis. <i>Lancet, The</i> , 2013, 381, 1371-1379.	6.3	2,643

#	ARTICLE	IF	CITATIONS
40	A genome-wide association study of a sustained pattern of antidepressant response. <i>Journal of Psychiatric Research</i> , 2013, 47, 1157-1165.	1.5	52
41	Research Review: The role of cytokines in depression in adolescents: a systematic review. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2013, 54, 816-835.	3.1	73
42	Genome-wide association studies in psychiatry: what have we learned?. <i>British Journal of Psychiatry</i> , 2013, 202, 1-4.	1.7	75
43	Gene × environment interactions in the prediction of response to antidepressant treatment. <i>International Journal of Neuropsychopharmacology</i> , 2013, 16, 701-711.	1.0	27
44	Discussion of "The Psychodynamic Psychotherapist's Guide to the Interaction among Sex, Genes, and Environmental Adversity in the Etiology of Depression for Women," by Simone N. Vigod and Valerie H. Taylor. <i>Psychodynamic Psychiatry</i> , 2013, 41, 553-561.	0.1	0
45	Secondary Use of Health Information. <i>JAMA Internal Medicine</i> , 2013, 173, 1806.	2.6	10
46	The future of genomics for developmentalists. <i>Development and Psychopathology</i> , 2013, 25, 1263-1278.	1.4	41
47	Integrative mouse and human mRNA studies using WGCNA nominates novel candidate genes involved in the pathogenesis of major depressive disorder. <i>Pharmacogenomics</i> , 2013, 14, 1979-1990.	0.6	55
48	USING MENDELIAN RANDOMISATION TO INFER CAUSALITY IN DEPRESSION AND ANXIETY RESEARCH. <i>Depression and Anxiety</i> , 2013, 30, 1185-1193.	2.0	27
49	Monozygotic twins affected with major depressive disorder have greater variance in methylation than their unaffected co-twin. <i>Translational Psychiatry</i> , 2013, 3, e269-e269.	2.4	89
50	Assessment of Genetic and Nongenetic Interactions for the Prediction of Depressive Symptomatology: An Analysis of the Wisconsin Longitudinal Study Using Machine Learning Algorithms. <i>American Journal of Public Health</i> , 2013, 103, S136-S144.	1.5	27
52	Microarray Profiling and Co-Expression Network Analysis of Circulating lncRNAs and mRNAs Associated with Major Depressive Disorder. <i>PLoS ONE</i> , 2014, 9, e93388.	1.1	103
53	A Conserved BDNF, Glutamate- and GABA-Enriched Gene Module Related to Human Depression Identified by Coexpression Meta-Analysis and DNA Variant Genome-Wide Association Studies. <i>PLoS ONE</i> , 2014, 9, e90980.	1.1	75
54	Analyzing Genome-Wide Association Studies with an FDR Controlling Modification of the Bayesian Information Criterion. <i>PLoS ONE</i> , 2014, 9, e103322.	1.1	18
55	A Genetic Variant in 12q13, a Possible Risk Factor for Bipolar Disorder, Is Associated with Depressive State, Accounting for Stressful Life Events. <i>PLoS ONE</i> , 2014, 9, e115135.	1.1	13
56	Mental health: Depression needs large human-genetics studies. <i>Nature</i> , 2014, 515, 189-191.	13.7	40
57	Medical research: If depression were cancer. <i>Nature</i> , 2014, 515, 182-184.	13.7	84
58	Jumping on the Train of Personalized Medicine: A Primer for Non-Geneticist Clinicians: Part 2. Fundamental Concepts in Genetic Epidemiology. <i>Current Psychiatry Reviews</i> , 2014, 10, 101-117.	0.9	10

#	ARTICLE	IF	CITATIONS
61	Genome-wide scans of genetic variants for psychophysiological endophenotypes: A methodological overview. <i>Psychophysiology</i> , 2014, 51, 1207-1224.	1.2	28
62	Gene-environment interaction research in psychiatric epidemiology: a framework and implications for study design. <i>Social Psychiatry and Psychiatric Epidemiology</i> , 2014, 49, 1525-1529.	1.6	8
63	Epigenetics of Depression. <i>Progress in Molecular Biology and Translational Science</i> , 2014, 128, 103-137.	0.9	28
65	Association of HTR2A T102C and A-1438G polymorphisms with susceptibility to major depressive disorder: a meta-analysis. <i>Neurological Sciences</i> , 2014, 35, 1857-1866.	0.9	39
66	Type I interferon signaling genes in recurrent major depression: increased expression detected by whole-blood RNA sequencing. <i>Molecular Psychiatry</i> , 2014, 19, 1267-1274.	4.1	151
67	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, 1872-1879.	2.8	49
68	Functional SNPs are enriched for schizophrenia association signals. <i>Molecular Psychiatry</i> , 2014, 19, 276-277.	4.1	23
69	Association between serotonin transporter genotype, brain structure and adolescent-onset major depressive disorder: a longitudinal prospective study. <i>Translational Psychiatry</i> , 2014, 4, e445-e445.	2.4	22
70	MZ twin pairs or MZ singletons in population family-based GWAS? More power in pairs. <i>Molecular Psychiatry</i> , 2014, 19, 1154-1155.	4.1	20
71	Investigating the possible causal association of smoking with depression and anxiety using Mendelian randomisation meta-analysis: the CARTA consortium. <i>BMJ Open</i> , 2014, 4, e006141.	0.8	150
72	Premorbid risk factors for major depressive disorder: Are they associated with early onset and recurrent course?. <i>Development and Psychopathology</i> , 2014, 26, 1477-1493.	1.4	54
74	A Polygenic Risk Score Associated with Measures of Depressive Symptoms Among Older Adults. <i>Biodemography and Social Biology</i> , 2014, 60, 199-211.	0.4	51
75	Depression and BMI influences the serum vascular endothelial growth factor level. <i>International Journal of Neuropsychopharmacology</i> , 2014, 17, 1409-1417.	1.0	27
76	Temperamental Contributions to the Development of Psychological Profiles: I. Basic Issues. , 2014, , 377-418.		1
77	Circadian rhythms and mood: Opportunities for multi-level analyses in genomics and neuroscience. <i>BioEssays</i> , 2014, 36, 305-315.	1.2	10
78	Applying polygenic risk scores to postpartum depression. <i>Archives of Women's Mental Health</i> , 2014, 17, 519-528.	1.2	62
79	Social neuroscience and its potential contribution to psychiatry. <i>World Psychiatry</i> , 2014, 13, 131-139.	4.8	56
80	A recessive genetic model and runs of homozygosity in major depressive disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2014, 165, 157-166.	1.1	20

#	ARTICLE	IF	CITATIONS
81	PSYCHIATRIC GENETICS AND THE FUTURE OF PERSONALIZED TREATMENT. <i>Depression and Anxiety</i> , 2014, 31, 893-898.	2.0	16
82	Genome-wide polygenic scoring for a 14-year long-term average depression phenotype. <i>Brain and Behavior</i> , 2014, 4, 298-311.	1.0	19
83	Further confirmation of the association between anxiety and <i>CTNND2</i> : replication in humans. <i>Genes, Brain and Behavior</i> , 2014, 13, 195-201.	1.1	43
84	Neuroplasticity and memory formation in major depressive disorder: An imaging genetics perspective on serotonin and BDNF. <i>Restorative Neurology and Neuroscience</i> , 2014, 32, 25-49.	0.4	22
85	Resistance to antidepressant drugs. <i>Behavioural Pharmacology</i> , 2014, 25, 352-371.	0.8	29
86	Autism Spectrum Disorder Genetics. <i>Harvard Review of Psychiatry</i> , 2014, 22, 65-75.	0.9	59
87	Prenatal risk factors for depression: a critical review of the evidence and potential mechanisms. <i>Journal of Developmental Origins of Health and Disease</i> , 2014, 5, 339-350.	0.7	21
88	Clarifying the causal relationship in women between childhood sexual abuse and lifetime major depression. <i>Psychological Medicine</i> , 2014, 44, 1213-1221.	2.7	31
89	The dynamic nature of depression: a new micro-level perspective of mental disorder that meets current challenges. <i>Psychological Medicine</i> , 2014, 44, 1349-1360.	2.7	213
90	Hypermethylation in the ZBTB20 gene is associated with major depressive disorder. <i>Genome Biology</i> , 2014, 15, R56.	13.9	87
92	The ceramide system as a novel antidepressant target. <i>Trends in Pharmacological Sciences</i> , 2014, 35, 293-304.	4.0	96
93	Brief, unidimensional melancholia rating scales are highly sensitive to the effect of citalopram and may have biological validity: Implications for the Research Domain Criteria (RDoC). <i>Journal of Affective Disorders</i> , 2014, 163, 18-24.	2.0	36
94	Genetic influences on alcohol use across stages of development: <i>GABRA2</i> and longitudinal trajectories of drunkenness from adolescence to young adulthood. <i>Addiction Biology</i> , 2014, 19, 1055-1064.	1.4	41
95	Explaining additional genetic variation in complex traits. <i>Trends in Genetics</i> , 2014, 30, 124-132.	2.9	128
96	The Genetics of Major Depression. <i>Neuron</i> , 2014, 81, 484-503.	3.8	559
97	DNA mismatch repair MSH2 gene-based SNP associated with different populations. <i>Molecular Genetics and Genomics</i> , 2014, 289, 469-487.	1.0	1
98	Testing the role of circadian genes in conferring risk for psychiatric disorders. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2014, 165, 254-260.	1.1	39
99	The gender-specific association of EHD3 polymorphisms with major depressive disorder. <i>Neuroscience Letters</i> , 2014, 567, 11-14.	1.0	9

#	ARTICLE	IF	CITATIONS
100	Brain galanin system genes interact with life stresses in depression-related phenotypes. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E1666-73.	3.3	83
101	Large-scale genomics unveils the genetic architecture of psychiatric disorders. Nature Neuroscience, 2014, 17, 782-790.	7.1	321
102	Revitalizing Psychiatric Therapeutics. Neuropsychopharmacology, 2014, 39, 220-229.	2.8	76
103	Stratified medicine for mental disorders. European Neuropsychopharmacology, 2014, 24, 5-50.	0.3	152
104	Epigenetics and the regulation of stress vulnerability and resilience. Neuroscience, 2014, 264, 157-170.	1.1	165
105	Gene-Environment Interaction. Annual Review of Psychology, 2014, 65, 41-70.	9.9	224
106	Gene-environment interactions in common mental disorders: an update and strategy for a genome-wide search. Social Psychiatry and Psychiatric Epidemiology, 2014, 49, 3-14.	1.6	74
107	Genetic, Molecular and Clinical Determinants for the Involvement of Aldosterone and Its Receptors in Major Depression. Nephron Physiology, 2014, 128, 17-25.	1.5	27
108	Interaction among childhood trauma and functional polymorphisms in the serotonin pathway moderate the risk of depressive disorders. European Archives of Psychiatry and Clinical Neuroscience, 2014, 264, 45-54.	1.8	29
109	Commentary: G × E in child psychiatry and psychology: a broadening of the scope of enquiry as prompted by Munaf et al. (2014). Journal of Child Psychology and Psychiatry and Allied Disciplines, 2014, 55, 1102-1104.	3.1	4
110	Chipping away at major depressive disorder. Genome Biology, 2014, 15, 421.	3.8	4
111	Practitioner Review: A critical perspective on gene-environment interaction models – what impact should they have on clinical perceptions and practice?. Journal of Child Psychology and Psychiatry and Allied Disciplines, 2014, 55, 1092-1101.	3.1	33
112	Advancements of Mass Spectrometry in Biomedical Research. Advances in Experimental Medicine and Biology, 2014, , .	0.8	6
113	Decreased peripheral brain-derived neurotrophic factor levels are a biomarker of disease activity in major psychiatric disorders: a comparative meta-analysis. Molecular Psychiatry, 2014, 19, 750-751.	4.1	135
114	Introduction to Epigenetics in Psychiatry. , 2014, , 3-25.		6
115	Genetic Studies of Major Depressive Disorder: Why Are There No Genome-wide Association Study Findings and What Can We Do About It?. Biological Psychiatry, 2014, 76, 510-512.	0.7	161
116	Metamoodics: meta-analysis and bioinformatics resource for mood disorders. Molecular Psychiatry, 2014, 19, 748-749.	4.1	16
117	A genome-wide association study of anorexia nervosa. Molecular Psychiatry, 2014, 19, 1085-1094.	4.1	282

#	ARTICLE	IF	CITATIONS
118	Biomarkers in Major Depressive Disorder: The Role of Mass Spectrometry. <i>Advances in Experimental Medicine and Biology</i> , 2014, 806, 545-560.	0.8	13
119	Chasing Mendel: five questions for personalized medicine. <i>Journal of Physiology</i> , 2014, 592, 2381-2388.	1.3	30
120	The endogenous and reactive depression subtypes revisited: integrative animal and human studies implicate multiple distinct molecular mechanisms underlying major depressive disorder. <i>BMC Medicine</i> , 2014, 12, 73.	2.3	52
121	Genetic relationships between suicide attempts, suicidal ideation and major psychiatric disorders: A genome-wide association and polygenic scoring study. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2014, 165, 428-437.	1.1	99
122	Arguments for the sake of endophenotypes: Examining common misconceptions about the use of endophenotypes in psychiatric genetics. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2014, 165, 122-130.	1.1	135
123	A New Piece to Understanding the Intimate Partner Violence Puzzle. <i>Violence Against Women</i> , 2014, 20, 414-419.	1.1	2
124	An Excess of Risk-Increasing Low-Frequency Variants Can Be a Signal of Polygenic Inheritance in Complex Diseases. <i>American Journal of Human Genetics</i> , 2014, 94, 437-452.	2.6	55
125	Investigating the genetic variation underlying episodicity in major depressive disorder: Suggestive evidence for a bipolar contribution. <i>Journal of Affective Disorders</i> , 2014, 155, 81-89.	2.0	15
126	Letter to editor: Failure to replicate the association of glucocorticoid and type 1 corticotropin-releasing hormone receptors gene variants with risk of depression during pregnancy and post-partum reported by Engineer et Al. (2013). <i>Journal of Psychiatric Research</i> , 2014, 56, 168-170.	1.5	14
127	ITIH3 polymorphism may confer susceptibility to psychiatric disorders by altering the expression levels of GLT8D1. <i>Journal of Psychiatric Research</i> , 2014, 50, 79-83.	1.5	24
128	Effect of polygenic risk scores on depression in childhood trauma. <i>British Journal of Psychiatry</i> , 2014, 205, 113-119.	1.7	167
129	708 Common and 2010 rare DISC1 locus variants identified in 1542 subjects: analysis for association with psychiatric disorder and cognitive traits. <i>Molecular Psychiatry</i> , 2014, 19, 668-675.	4.1	59
130	Allelic differences between Europeans and Chinese for CREB1 SNPs and their implications in gene expression regulation, hippocampal structure and function, and bipolar disorder susceptibility. <i>Molecular Psychiatry</i> , 2014, 19, 452-461.	4.1	61
131	The genetic interacting landscape of 63 candidate genes in Major Depressive Disorder: an explorative study. <i>BioData Mining</i> , 2014, 7, 19.	2.2	7
133	Familiality and SNP heritability of age at onset and episodicity in major depressive disorder. <i>Psychological Medicine</i> , 2015, 45, 2215-2225.	2.7	21
134	Epistatic and gene wide effects in YWHA and aromatic amino hydroxylase genes across ADHD and other common neuropsychiatric disorders: Association with YWHA. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2015, 168, 423-432.	1.1	21
135	The interaction between stress and genetic factors in the etiopathogenesis of depression. <i>World Psychiatry</i> , 2015, 14, 161-163.	4.8	51
136	The relationship between schizophrenia and rheumatoid arthritis revisited: Genetic and epidemiological analyses. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2015, 168, 81-88.	1.1	29

#	ARTICLE	IF	CITATIONS
138	Personalized Medicine: Review and Perspectives of Promising Baseline EEG Biomarkers in Major Depressive Disorder and Attention Deficit Hyperactivity Disorder. <i>Neuropsychobiology</i> , 2015, 72, 229-240.	0.9	127
141	Early life trauma, depression and the glucocorticoid receptor gene â€“ an epigenetic perspective. <i>Psychological Medicine</i> , 2015, 45, 3393-3410.	2.7	51
143	Epigenome-Wide Association Study of Aggressive Behavior. <i>Twin Research and Human Genetics</i> , 2015, 18, 686-698.	0.3	53
144	Tryptophan hydroxylase 2 and Bcl-xL in the rat raphe nucleus after acute and chronic forced swim stress. <i>Russian Journal of Genetics: Applied Research</i> , 2015, 5, 577-581.	0.4	1
145	Large Scale Genetic Research on Neuropsychiatric Disorders in African Populations is Needed. <i>EBioMedicine</i> , 2015, 2, 1259-1261.	2.7	32
146	Cumulative role of rare and common putative functional genetic variants at <i>NPAS3</i> in schizophrenia susceptibility. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2015, 168, 528-535.	1.1	5
147	Shared additive genetic influences on DSMâ€™ criteria for alcohol dependence in subjects of European ancestry. <i>Addiction</i> , 2015, 110, 1922-1931.	1.7	20
148	Comparative genome-wide association studies of a depressive symptom phenotype in a repeated measures setting by race/ethnicity in the multi-ethnic study of atherosclerosis. <i>BMC Genetics</i> , 2015, 16, 118.	2.7	12
149	Wholeâ€™genome sequencing provides insight into the genetics of major depressive disorder. <i>Clinical Genetics</i> , 2015, 88, 340-342.	1.0	1
150	Genetic Analyses Benefit From Using Less Heterogeneous Phenotypes: An Illustration With the Hospital Anxiety and Depression Scale (HADS). <i>Genetic Epidemiology</i> , 2015, 39, 317-324.	0.6	13
152	Genetic Predictors of Depressive Symptoms in the Look AHEAD Trial. <i>Psychosomatic Medicine</i> , 2015, 77, 982-992.	1.3	7
153	Genome-Wide Scan Informed by Age-Related Disease Identifies Loci for Exceptional Human Longevity. <i>PLoS Genetics</i> , 2015, 11, e1005728.	1.5	128
154	Explore the Features of Brain-Derived Neurotrophic Factor in Mood Disorders. <i>PLoS ONE</i> , 2015, 10, e0128605.	1.1	12
155	Epidemiology and Heritability of Major Depressive Disorder, Stratified by Age of Onset, Sex, and Illness Course in Generation Scotland: Scottish Family Health Study (GS:SFHS). <i>PLoS ONE</i> , 2015, 10, e0142197.	1.1	101
156	FRAS1-related extracellular matrix 3 (FREM3) single-nucleotide polymorphism effects on gene expression, amygdala reactivity and perceptual processing speed: An accelerated aging pathway of depression risk. <i>Frontiers in Psychology</i> , 2015, 6, 1377.	1.1	17
158	First robust genetic links to depression emerge. <i>Nature</i> , 2015, 523, 268-269.	13.7	15
159	Genetic Determinants of Depression. <i>Harvard Review of Psychiatry</i> , 2015, 23, 1-18.	0.9	132
160	Meta-analysis of Genome-wide Association Studies for Neuroticism, and the Polygenic Association With Major Depressive Disorder. <i>JAMA Psychiatry</i> , 2015, 72, 642.	6.0	289

#	ARTICLE	IF	CITATIONS
161	The association between lower educational attainment and depression owing to shared genetic effects? Results in ~25â€‰%000 subjects. <i>Molecular Psychiatry</i> , 2015, 20, 735-743.	4.1	59
162	The Genetics of Loneliness. <i>Perspectives on Psychological Science</i> , 2015, 10, 213-226.	5.2	80
163	GeneÃƒâ€šEnvironment Interaction in Major Depression: Focus on Experience-Dependent Biological Systems. <i>Frontiers in Psychiatry</i> , 2015, 6, 68.	1.3	113
164	An association study of the m6A genes with major depressive disorder in Chinese Han population. <i>Journal of Affective Disorders</i> , 2015, 183, 279-286.	2.0	93
165	Schizophrenia and Depression Co-Morbidity: What We have Learned from Animal Models. <i>Frontiers in Psychiatry</i> , 2015, 6, 13.	1.3	55
167	Proteasome system dysregulation and treatment resistance mechanisms in major depressive disorder. <i>Translational Psychiatry</i> , 2015, 5, e687-e687.	2.4	26
168	Analyzing pathways from childhood maltreatment to internalizing symptoms and disorders in children and adolescents (AMIS): a study protocol. <i>BMC Psychiatry</i> , 2015, 15, 126.	1.1	14
169	What have we learned from the Psychiatric Genomics Consortium. <i>World Psychiatry</i> , 2015, 14, 291-293.	4.8	29
170	DISTMIX: direct imputation of summary statistics for unmeasured SNPs from mixed ethnicity cohorts. <i>Bioinformatics</i> , 2015, 31, 3099-3104.	1.8	25
171	Genetic Differences in the Immediate Transcriptome Response to Stress Predict Risk-Related Brain Function and Psychiatric Disorders. <i>Neuron</i> , 2015, 86, 1189-1202.	3.8	102
172	Expression of genes in the brain associated with depression. <i>Russian Journal of Genetics: Applied Research</i> , 2015, 5, 582-588.	0.4	1
173	The successful search for genetic loci associated with depression. <i>Genome Medicine</i> , 2015, 7, 92.	3.6	3
174	DNA Modification Study of Major Depressive Disorder: Beyond Locus-by-Locus Comparisons. <i>Biological Psychiatry</i> , 2015, 77, 246-255.	0.7	66
175	Integrating NIMH Research Domain Criteria (RDoC) into depression research. <i>Current Opinion in Psychology</i> , 2015, 4, 6-12.	2.5	139
176	The role of the potassium channel gene KCNK2 in major depressive disorder. <i>Psychiatry Research</i> , 2015, 225, 489-492.	1.7	10
177	The mathematical limits of genetic prediction for complex chronic disease. <i>Journal of Epidemiology and Community Health</i> , 2015, 69, 574-579.	2.0	21
178	Effect of variation in BDNF Val66Met polymorphism, smoking, and nicotine dependence on symptom severity of depressive and anxiety disorders. <i>Drug and Alcohol Dependence</i> , 2015, 148, 150-157.	1.6	14
179	New insights into the neurobiological mechanisms of major depressive disorders. <i>General Hospital Psychiatry</i> , 2015, 37, 172-177.	1.2	33

#	ARTICLE	IF	CITATIONS
180	Genetic pleiotropy between multiple sclerosis and schizophrenia but not bipolar disorder: differential involvement of immune-related gene loci. <i>Molecular Psychiatry</i> , 2015, 20, 207-214.	4.1	173
181	GABAergic Control of Depression-Related Brain States. <i>Advances in Pharmacology</i> , 2015, 73, 97-144.	1.2	107
182	LD Score regression distinguishes confounding from polygenicity in genome-wide association studies. <i>Nature Genetics</i> , 2015, 47, 291-295.	9.4	3,905
183	Psychiatric genome-wide association study analyses implicate neuronal, immune and histone pathways. <i>Nature Neuroscience</i> , 2015, 18, 199-209.	7.1	701
184	Src kinase as a mediator of convergent molecular abnormalities leading to NMDAR hypoactivity in schizophrenia. <i>Molecular Psychiatry</i> , 2015, 20, 1091-1100.	4.1	56
185	Epigenetics and depressive disorders: a review of current progress and future directions. <i>International Journal of Epidemiology</i> , 2015, 44, 1364-1387.	0.9	84
186	The Promise of Biomarkers in Diagnosing Major Depression in Primary Care: the Present and Future. <i>Current Psychiatry Reports</i> , 2015, 17, 601.	2.1	17
187	The GSK3B gene confers risk for both major depressive disorder and schizophrenia in the Han Chinese population. <i>Journal of Affective Disorders</i> , 2015, 185, 149-155.	2.0	34
188	The NVL gene confers risk for both major depressive disorder and schizophrenia in the Han Chinese population. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2015, 62, 7-13.	2.5	17
189	Functional characterization of the PCLO p.Ser4814Ala variant associated with major depressive disorder reveals cellular but not behavioral differences. <i>Neuroscience</i> , 2015, 300, 518-538.	1.1	13
190	Mental Illness, <i>Genetics of.</i> , 2015, , 209-215.		0
191	Associations with depression. <i>Nature</i> , 2015, 523, 539-540.	13.7	16
192	Sparse whole-genome sequencing identifies two loci for major depressive disorder. <i>Nature</i> , 2015, 523, 588-591.	13.7	777
193	Epigenetics of Stress-Related Psychiatric Disorders and Gene-Environment Interactions. <i>Neuron</i> , 2015, 86, 1343-1357.	3.8	271
194	Problematic assumptions have slowed down depression research: why symptoms, not syndromes are the way forward. <i>Frontiers in Psychology</i> , 2015, 6, 309.	1.1	222
195	Blood transcriptomic markers for major depression: from animal models to clinical settings. <i>Annals of the New York Academy of Sciences</i> , 2015, 1344, 37-49.	1.8	16
196	IGF-I in major depression and antidepressant treatment response. <i>European Neuropsychopharmacology</i> , 2015, 25, 864-872.	0.3	53
197	The inflammatory cytokines: molecular biomarkers for major depressive disorder?. <i>Biomarkers in Medicine</i> , 2015, 9, 169-180.	0.6	31

#	ARTICLE	IF	CITATIONS
198	An update on research and approaches in biological psychiatry. <i>Neuroscience Bulletin</i> , 2015, 31, 1-3.	1.5	0
199	Genes, environments and depressions in young people. <i>Archives of Disease in Childhood</i> , 2015, 100, 1064-1069.	1.0	11
200	Depression sum-scores don't add up: why analyzing specific depression symptoms is essential. <i>BMC Medicine</i> , 2015, 13, 72.	2.3	528
201	Longevity Genes. <i>Advances in Experimental Medicine and Biology</i> , 2015, , .	0.8	5
202	Models to Explore Genetics of Human Aging. <i>Advances in Experimental Medicine and Biology</i> , 2015, 847, 141-161.	0.8	4
203	Mendelian Randomization: New Applications in the Coming Age of Hypothesis-Free Causality. <i>Annual Review of Genomics and Human Genetics</i> , 2015, 16, 327-350.	2.5	298
204	Genetics of anxiety disorders: Genetic epidemiological and molecular studies in humans. <i>Psychiatry and Clinical Neurosciences</i> , 2015, 69, 388-401.	1.0	109
205	Direct, indirect and pleiotropic effects of candidate genes on internalizing disorder psychopathology. <i>Psychological Medicine</i> , 2015, 45, 2227-2236.	2.7	20
206	Polygenic risk, stressful life events and depressive symptoms in older adults: a polygenic score analysis. <i>Psychological Medicine</i> , 2015, 45, 1709-1720.	2.7	98
207	Genetics and genomics of psychiatric disease. <i>Science</i> , 2015, 349, 1489-1494.	6.0	337
208	Interaction of tryptophan hydroxylase 2 gene and life events in susceptibility to major depression in a Chinese Han population. <i>Journal of Affective Disorders</i> , 2015, 188, 304-309.	2.0	18
209	Serum proteomic profiling of major depressive disorder. <i>Translational Psychiatry</i> , 2015, 5, e599-e599.	2.4	100
210	Circadian Disruption in Psychiatric Disorders. <i>Sleep Medicine Clinics</i> , 2015, 10, 481-493.	1.2	65
211	BDNF promoter methylation and genetic variation in late-life depression. <i>Translational Psychiatry</i> , 2015, 5, e619-e619.	2.4	111
212	Identification of type 2 diabetes subgroups through topological analysis of patient similarity. <i>Science Translational Medicine</i> , 2015, 7, 311ra174.	5.8	426
213	Lack of association between type 2 diabetes and major depression: epidemiologic and genetic evidence in a multiethnic population. <i>Translational Psychiatry</i> , 2015, 5, e618-e618.	2.4	32
214	Adjusting heterogeneous ascertainment bias for genetic association analysis with extended families. <i>BMC Medical Genetics</i> , 2015, 16, 62.	2.1	9
215	Major depressive disorder and current psychological distress moderate the effect of polygenic risk for obesity on body mass index. <i>Translational Psychiatry</i> , 2015, 5, e592-e592.	2.4	24

#	ARTICLE	IF	CITATIONS
216	Single Nucleotide Polymorphism Heritability of Behavior Problems in Childhood: Genome-Wide Complex Trait Analysis. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2015, 54, 737-744.	0.3	40
217	Psychometric precision in phenotype definition is a useful step in molecular genetic investigation of psychiatric disorders. <i>Translational Psychiatry</i> , 2015, 5, e593-e593.	2.4	14
218	Translating depression biomarkers for improved targeted therapies. <i>Neuroscience and Biobehavioral Reviews</i> , 2015, 59, 1-15.	2.9	19
219	Genome-wide methylation study on depression: differential methylation and variable methylation in monozygotic twins. <i>Translational Psychiatry</i> , 2015, 5, e557-e557.	2.4	98
220	Psychiatric genomics: outlook for 2015 and challenges for 2020. <i>Current Opinion in Behavioral Sciences</i> , 2015, 2, 102-107.	2.0	3
221	JEPEG: a summary statistics based tool for gene-level joint testing of functional variants. <i>Bioinformatics</i> , 2015, 31, 1176-1182.	1.8	27
222	Identification of genes and gene pathways associated with major depressive disorder by integrative brain analysis of rat and human prefrontal cortex transcriptomes. <i>Translational Psychiatry</i> , 2015, 5, e519-e519.	2.4	43
223	Increased serum levels of sortilin are associated with depression and correlated with BDNF and VEGF. <i>Translational Psychiatry</i> , 2015, 5, e677-e677.	2.4	39
224	Disruption of sonic hedgehog signaling in Ellis-van Creveld dwarfism confers protection against bipolar affective disorder. <i>Molecular Psychiatry</i> , 2015, 20, 1212-1218.	4.1	8
225	Novel loci associated with usual sleep duration: the CHARGE Consortium Genome-Wide Association Study. <i>Molecular Psychiatry</i> , 2015, 20, 1232-1239.	4.1	112
226	Common variants of the PINK1 and PARL genes do not confer genetic susceptibility to schizophrenia in Han Chinese. <i>Molecular Genetics and Genomics</i> , 2015, 290, 585-592.	1.0	2
227	Schizophrenia as a Disorder of Molecular Pathways. <i>Biological Psychiatry</i> , 2015, 77, 22-28.	0.7	80
228	Coordinated Messenger RNA/MicroRNA Changes in Fibroblasts of Patients with Major Depression. <i>Biological Psychiatry</i> , 2015, 77, 256-265.	0.7	57
229	Heterogeneity and Individuality: microRNAs in Mental Disorders. <i>Journal of Neural Transmission</i> , 2015, 122, 79-97.	1.4	49
230	Caseâ€“Control Genome-Wide Association Study of Persistent Attention-Deficit Hyperactivity Disorder Identifies FBXO33 as a Novel Susceptibility Gene for the Disorder. <i>Neuropsychopharmacology</i> , 2015, 40, 915-926.	2.8	59
231	Medical Epigenetics and Twins. , 2016, , 147-158.		0
232	Epigenetics of Psychiatric Disorders. , 2016, , 335-350.		2
233	Estimating Effect Sizes and Expected Replication Probabilities from GWAS Summary Statistics. <i>Frontiers in Genetics</i> , 2016, 7, 15.	1.1	40

#	ARTICLE	IF	CITATIONS
234	Genetic Pathways to Insomnia. <i>Brain Sciences</i> , 2016, 6, 64.	1.1	54
235	Pathogenetic and Therapeutic Applications of Tumor Necrosis Factor- α (TNF- α) in Major Depressive Disorder: A Systematic Review. <i>International Journal of Molecular Sciences</i> , 2016, 17, 733.	1.8	134
236	Epigenetic Modifications of Major Depressive Disorder. <i>International Journal of Molecular Sciences</i> , 2016, 17, 1279.	1.8	81
237	Predicting gene targets from integrative analyses of summary data from GWAS and eQTL studies for 28 human complex traits. <i>Genome Medicine</i> , 2016, 8, 84.	3.6	91
238	Human Bacterial Artificial Chromosome (BAC) Transgenesis Fully Rescues Noradrenergic Function in Dopamine β -Hydroxylase Knockout Mice. <i>PLoS ONE</i> , 2016, 11, e0154864.	1.1	12
239	The First Pilot Genome-Wide Gene-Environment Study of Depression in the Japanese Population. <i>PLoS ONE</i> , 2016, 11, e0160823.	1.1	30
240	Is there Progress? An Overview of Selecting Biomarker Candidates for Major Depressive Disorder. <i>Frontiers in Psychiatry</i> , 2016, 7, 72.	1.3	53
241	Effect of the interaction between childhood abuse and rs1360780 of the <i>FKBP5</i> gene on gray matter volume in a general population sample. <i>Human Brain Mapping</i> , 2016, 37, 1602-1613.	1.9	62
242	Common variants in <i>CACNA1C</i> and MDD susceptibility: A comprehensive meta-analysis. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2016, 171, 896-903.	1.1	33
243	An epigenetic resolution of the lek paradox. <i>BioEssays</i> , 2016, 38, 355-366.	1.2	18
244	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, 265-280.	2.0	99
245	Computational meta-analysis of statistical parametric maps in major depression. <i>Human Brain Mapping</i> , 2016, 37, 1393-1404.	1.9	158
246	DISC1 as a Possible Genetic Contribution to Opioid Dependence in a Polish Sample. <i>Journal of Studies on Alcohol and Drugs</i> , 2016, 77, 220-226.	0.6	3
248	Metabolomic biosignature differentiates melancholic depressive patients from healthy controls. <i>BMC Genomics</i> , 2016, 17, 669.	1.2	35
249	Controlling the joint local false discovery rate is more powerful than meta-analysis methods in joint analysis of summary statistics from multiple genome-wide association studies. <i>Bioinformatics</i> , 2017, 33, 500-507.	1.8	21
250	Peripheral blood gene expression profiles linked to monoamine metabolite levels in cerebrospinal fluid. <i>Translational Psychiatry</i> , 2016, 6, e983-e983.	2.4	14
251	Association of ATP6V1B2 rs1106634 with lifetime risk of depression and hippocampal neurocognitive deficits: possible novel mechanisms in the etiopathology of depression. <i>Translational Psychiatry</i> , 2016, 6, e945-e945.	2.4	12
252	Alterations in the neuropeptide galanin system in major depressive disorder involve levels of transcripts, methylation, and peptide. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E8472-E8481.	3.3	43

#	ARTICLE	IF	CITATIONS
253	Genome-wide association study of response to cognitive-behavioural therapy in children with anxiety disorders. <i>British Journal of Psychiatry</i> , 2016, 209, 236-243.	1.7	39
254	Somatic, positive and negative domains of the Center for Epidemiological Studies Depression (CES-D) scale: a meta-analysis of genome-wide association studies. <i>Psychological Medicine</i> , 2016, 46, 1613-1623.	2.7	17
255	Genome-wide association analysis of pain severity in dysmenorrhea identifies association at chromosome 1p13.2, near the nerve growth factor locus. <i>Pain</i> , 2016, 157, 2571-2581.	2.0	36
256	The Twin Study of Negative Valence Emotional Constructs. <i>Twin Research and Human Genetics</i> , 2016, 19, 456-464.	0.3	20
257	Body mass index and psychiatric disorders: a Mendelian randomization study. <i>Scientific Reports</i> , 2016, 6, 32730.	1.6	47
258	The potential of genetic and gene expression analysis in the diagnosis of neuropsychiatric disorders. <i>Expert Review of Molecular Diagnostics</i> , 2016, 16, 677-695.	1.5	5
259	A method to decipher pleiotropy by detecting underlying heterogeneity driven by hidden subgroups applied to autoimmune and neuropsychiatric diseases. <i>Nature Genetics</i> , 2016, 48, 803-810.	9.4	62
260	Polygenic interactions with environmental adversity in the aetiology of major depressive disorder. <i>Psychological Medicine</i> , 2016, 46, 759-770.	2.7	176
261	The effects of the interplay of genetics and early environmental risk on the course of internalizing symptoms from late childhood through adolescence. <i>Development and Psychopathology</i> , 2016, 28, 225-237.	1.4	14
262	Distinct effects of folate pathway genes MTHFR and MTHFD1L on ruminative response style: a potential risk mechanism for depression. <i>Translational Psychiatry</i> , 2016, 6, e745-e745.	2.4	23
263	Genetic variants associated with subjective well-being, depressive symptoms, and neuroticism identified through genome-wide analyses. <i>Nature Genetics</i> , 2016, 48, 624-633.	9.4	870
264	Genetic and Environmental Factors Associated with Cannabis Involvement. <i>Current Addiction Reports</i> , 2016, 3, 199-213.	1.6	13
265	Genome-wide analysis of over 106,000 individuals identifies 9 neuroticism-associated loci. <i>Molecular Psychiatry</i> , 2016, 21, 749-757.	4.1	220
266	Genome-wide Association Studies of Posttraumatic Stress Disorder in 2 Cohorts of US Army Soldiers. <i>JAMA Psychiatry</i> , 2016, 73, 695.	6.0	158
267	Guidelines for Reporting Articles on Psychiatry and Heart rate variability (GRAPH): recommendations to advance research communication. <i>Translational Psychiatry</i> , 2016, 6, e803-e803.	2.4	289
268	Identification of SLC25A37 as a major depressive disorder risk gene. <i>Journal of Psychiatric Research</i> , 2016, 83, 168-175.	1.5	24
269	MicroRNA 101b Is Downregulated in the Prefrontal Cortex of a Genetic Model of Depression and Targets the Glutamate Transporter SLC1A1 (EAAT3) <i>in Vitro</i> . <i>International Journal of Neuropsychopharmacology</i> , 2016, 19, pyw069.	1.0	22
271	Major depressive disorder. <i>Nature Reviews Disease Primers</i> , 2016, 2, 16065.	18.1	1,171

#	ARTICLE	IF	CITATIONS
272	Identification of 15 genetic loci associated with risk of major depression in individuals of European descent. <i>Nature Genetics</i> , 2016, 48, 1031-1036.	9.4	655
273	A current snapshot of common genomic variants contribution in psychiatric disorders. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2016, 171, 997-1005.	1.1	6
274	Depression, marijuana use and early-onset marijuana use conferred unique effects on neural connectivity and cognition. <i>Acta Psychiatrica Scandinavica</i> , 2016, 134, 399-409.	2.2	19
275	Gene and Network Analysis of Common Variants Reveals Novel Associations in Multiple Complex Diseases. <i>Genetics</i> , 2016, 204, 783-798.	1.2	56
276	Effects of corticotropin-releasing hormone receptor 1 SNPs on major depressive disorder are influenced by sex and smoking status. <i>Journal of Affective Disorders</i> , 2016, 205, 282-288.	2.0	11
277	Habitual coffee consumption and risk of type 2 diabetes, ischemic heart disease, depression and Alzheimer's disease: a Mendelian randomization study. <i>Scientific Reports</i> , 2016, 6, 36500.	1.6	55
278	Serum proteomic profiles of depressive subtypes. <i>Translational Psychiatry</i> , 2016, 6, e851-e851.	2.4	56
279	Analysis of 23andMe antidepressant efficacy survey data: implication of circadian rhythm and neuroplasticity in bupropion response. <i>Translational Psychiatry</i> , 2016, 6, e889-e889.	2.4	56
280	Consistently altered expression of gene sets in postmortem brains of individuals with major psychiatric disorders. <i>Translational Psychiatry</i> , 2016, 6, e890-e890.	2.4	63
281	Neuropsychological and neuroimaging evidence for the involvement of the frontal lobes in depression: 20 years on. <i>Journal of Psychopharmacology</i> , 2016, 30, 1090-1094.	2.0	11
282	Seeking an objective diagnosis of depression. <i>Biomarkers in Medicine</i> , 2016, 10, 861-875.	0.6	32
283	Contribution of Genetic Epidemiology to Our Understanding of Psychiatric Disorders. , 2016, , 27-50.		0
284	Conceptualizing Major Depression. , 2016, , 487-501.		0
285	Changing Polygenic Penetrance on Phenotypes in the 20th Century Among Adults in the US Population. <i>Scientific Reports</i> , 2016, 6, 30348.	1.6	40
286	Pupillary response abnormalities in depressive disorders. <i>Psychiatry Research</i> , 2016, 246, 492-499.	1.7	27
287	Shared Genetics and Couple-Associated Environment Are Major Contributors to the Risk of Both Clinical and Self-Declared Depression. <i>EBioMedicine</i> , 2016, 14, 161-167.	2.7	32
288	DNA co-methylation modules in postmortem prefrontal cortex tissues of European Australians with alcohol use disorders. <i>Scientific Reports</i> , 2016, 6, 19430.	1.6	68
289	Common variants in FKBP5 gene and major depressive disorder (MDD) susceptibility: a comprehensive meta-analysis. <i>Scientific Reports</i> , 2016, 6, 32687.	1.6	48

#	ARTICLE	IF	CITATIONS
290	Association of Genetic Risk Variants With Attention-Deficit/Hyperactivity Disorder Trajectories in the General Population. <i>JAMA Psychiatry</i> , 2016, 73, 1285.	6.0	115
291	Genetic variants linked to education predict longevity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 13366-13371.	3.3	110
292	Genetic Correlation Analysis Suggests Association between Increased Self-Reported Sleep Duration in Adults and Schizophrenia and Type 2 Diabetes. <i>Sleep</i> , 2016, 39, 1853-1857.	0.6	19
293	Heritability and Genome-Wide Association Analyses of Sleep Duration in Children: The EAGLE Consortium. <i>Sleep</i> , 2016, 39, 1859-1869.	0.6	34
294	Genetic and environmental components of female depression as a function of the severity of the disorder. <i>Brain and Behavior</i> , 2016, 6, e00519.	1.0	3
295	Genetic and Functional Study of $SLC6A4$-Type Amino Acid Transporter 1 in Schizophrenia. <i>Neuropsychobiology</i> , 2016, 74, 96-103.	0.9	7
296	On combining family- and population-based sequencing data. <i>BMC Proceedings</i> , 2016, 10, 175-179.	1.8	1
298	Quantitative Trait Loci and a Novel Genetic Candidate for Fear Learning. <i>Journal of Neuroscience</i> , 2016, 36, 6258-6268.	1.7	23
300	What Cure Models Can Teach us About Genome-Wide Survival Analysis. <i>Behavior Genetics</i> , 2016, 46, 269-280.	1.4	5
301	Telomere length and health outcomes: A two-sample genetic instrumental variables analysis. <i>Experimental Gerontology</i> , 2016, 82, 88-94.	1.2	22
302	Assortative mating and differential fertility by phenotype and genotype across the 20th century. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 6647-6652.	3.3	82
303	Nonsynonymous HTR2C polymorphism predicts cortisol response to psychosocial stress I: Effects in males and females. <i>Psychoneuroendocrinology</i> , 2016, 70, 134-141.	1.3	21
304	Identification of the BRD1 interaction network and its impact on mental disorder risk. <i>Genome Medicine</i> , 2016, 8, 53.	3.6	29
305	BDNF Val66Met polymorphism and bipolar disorder in European populations: A risk association in case-control, family-based and GWAS studies. <i>Neuroscience and Biobehavioral Reviews</i> , 2016, 68, 218-233.	2.9	69
306	Epigenetic differences in monozygotic twins discordant for major depressive disorder. <i>Translational Psychiatry</i> , 2016, 6, e839-e839.	2.4	38
307	Genome-wide association study reveals greater polygenic loading for schizophrenia in cases with a family history of illness. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2016, 171, 276-289.	1.1	28
308	Gene expression in major depressive disorder. <i>Molecular Psychiatry</i> , 2016, 21, 339-347.	4.1	178
309	Polygenic associations of neurodevelopmental genes in suicide attempt. <i>Molecular Psychiatry</i> , 2016, 21, 1381-1390.	4.1	84

#	ARTICLE	IF	CITATIONS
310	MetaKTSP: a meta-analytic top scoring pair method for robust cross-study validation of omics prediction analysis. <i>Bioinformatics</i> , 2016, 32, 1966-1973.	1.8	35
311	Recurrent major depression and right hippocampal volume: A bivariate linkage and association study. <i>Human Brain Mapping</i> , 2016, 37, 191-202.	1.9	21
312	Decoding the non-coding genome: elucidating genetic risk outside the coding genome. <i>Genes, Brain and Behavior</i> , 2016, 15, 187-204.	1.1	32
313	Impact of a cis-associated gene expression SNP on chromosome 20q11.22 on bipolar disorder susceptibility, hippocampal structure and cognitive performance. <i>British Journal of Psychiatry</i> , 2016, 208, 128-137.	1.7	11
314	The interaction of combined effects of the BDNF and PRKCG genes and negative life events in major depressive disorder. <i>Psychiatry Research</i> , 2016, 237, 72-77.	1.7	7
315	Survival in somatoform disorders. <i>British Journal of Psychiatry</i> , 2016, 208, 127-127.	1.7	1
316	Disease and Polygenic Architecture: Avoid Trio Design and Appropriately Account for Unscreened Control Subjects for Common Disease. <i>American Journal of Human Genetics</i> , 2016, 98, 382-391.	2.6	41
317	Depression as a risk factor for Alzheimer's disease: Genes, steroids, cytokines and neurogenesis – What do we need to know?. <i>Frontiers in Neuroendocrinology</i> , 2016, 41, 153-171.	2.5	102
318	Genetically reduced FAAH activity may be a risk for the development of anxiety and depression in persons with repetitive childhood trauma. <i>European Neuropsychopharmacology</i> , 2016, 26, 1020-1028.	0.3	60
319	Genetic epidemiology of migraine and depression. <i>Cephalalgia</i> , 2016, 36, 679-691.	1.8	46
320	Polygenic Risk of Schizophrenia and Cognition in a Population-Based Survey of Older Adults. <i>Schizophrenia Bulletin</i> , 2016, 42, 984-991.	2.3	44
321	Genome-wide linkage on chromosome 10q26 for a dimensional scale of major depression. <i>Journal of Affective Disorders</i> , 2016, 191, 123-131.	2.0	20
322	Transcriptomics analysis of iPSC-derived neurons and modeling of neuropsychiatric disorders. <i>Molecular and Cellular Neurosciences</i> , 2016, 73, 32-42.	1.0	33
323	Genetical Genomics of Behavior: A Novel Chicken Genomic Model for Anxiety Behavior. <i>Genetics</i> , 2016, 202, 327-340.	1.2	51
324	The specificity of the familial aggregation of early-onset bipolar disorder : A controlled 10-year follow-up study of offspring of parents with mood disorders. <i>Journal of Affective Disorders</i> , 2016, 190, 26-33.	2.0	54
325	The identification of symptom-based subtypes of depression: A nationally representative cohort study. <i>Journal of Affective Disorders</i> , 2016, 190, 395-406.	2.0	50
326	Age-Dependent Pleiotropy Between General Cognitive Function and Major Psychiatric Disorders. <i>Biological Psychiatry</i> , 2016, 80, 266-273.	0.7	71
327	The Genetics of Stress-Related Disorders: PTSD, Depression, and Anxiety Disorders. <i>Neuropsychopharmacology</i> , 2016, 41, 297-319.	2.8	332

#	ARTICLE	IF	CITATIONS
328	A haplotype in the 5'-upstream region of the NDUFV2 gene is associated with major depressive disorder in Han Chinese. <i>Journal of Affective Disorders</i> , 2016, 190, 329-332.	2.0	13
329	Models of Intergenerational and Transgenerational Transmission of Risk for Psychopathology in Mice. <i>Neuropsychopharmacology</i> , 2016, 41, 219-231.	2.8	91
330	Heritability of Perinatal Depression and Genetic Overlap With Nonperinatal Depression. <i>American Journal of Psychiatry</i> , 2016, 173, 158-165.	4.0	102
331	Phenome-wide analysis of genome-wide polygenic scores. <i>Molecular Psychiatry</i> , 2016, 21, 1188-1193.	4.1	154
332	Gene × Environment Determinants of Stress- and Anxiety-Related Disorders. <i>Annual Review of Psychology</i> , 2016, 67, 239-261.	9.9	106
333	Polygenic dissection of major depression clinical heterogeneity. <i>Molecular Psychiatry</i> , 2016, 21, 516-522.	4.1	154
334	High-throughput sequencing of the synaptome in major depressive disorder. <i>Molecular Psychiatry</i> , 2016, 21, 650-655.	4.1	31
335	Translating Neurogenomics Into New Medicines. <i>Biological Psychiatry</i> , 2016, 79, 650-656.	0.7	12
336	Subcortical brain volume abnormalities in 2028 individuals with schizophrenia and 2540 healthy controls via the ENIGMA consortium. <i>Molecular Psychiatry</i> , 2016, 21, 547-553.	4.1	820
337	Phenotypic Association Analyses With Copy Number Variation in Recurrent Depressive Disorder. <i>Biological Psychiatry</i> , 2016, 79, 329-336.	0.7	21
338	A Combined Pathway and Regional Heritability Analysis Indicates NETRIN1 Pathway Is Associated With Major Depressive Disorder. <i>Biological Psychiatry</i> , 2017, 81, 336-346.	0.7	32
339	Genome-wide Association for Major Depression Through Age at Onset Stratification: Major Depressive Disorder Working Group of the Psychiatric Genomics Consortium. <i>Biological Psychiatry</i> , 2017, 81, 325-335.	0.7	175
340	The Association of Cigarette Smoking With Depression and Anxiety: A Systematic Review. <i>Nicotine and Tobacco Research</i> , 2017, 19, 3-13.	1.4	730
341	Exploring the predictive power of polygenic scores derived from genome-wide association studies: a study of 10 complex traits. <i>Bioinformatics</i> , 2017, 33, 886-892.	1.8	39
342	Evidence for genetic heterogeneity between clinical subtypes of bipolar disorder. <i>Translational Psychiatry</i> , 2017, 7, e993-e993.	2.4	162
343	A New Hope for Biological Insights Into Depression. <i>Biological Psychiatry</i> , 2017, 81, 280-281.	0.7	3
344	Case-control association mapping by proxy using family history of disease. <i>Nature Genetics</i> , 2017, 49, 325-331.	9.4	192
345	Adolescent depression linked to socioeconomic status? Molecular approaches for revealing premorbid risk factors. <i>BioEssays</i> , 2017, 39, 1600194.	1.2	12

#	ARTICLE	IF	CITATIONS
346	Systematic review and meta-analysis of genetic studies of late-life depression. <i>Neuroscience and Biobehavioral Reviews</i> , 2017, 75, 129-139.	2.9	41
347	Association of polygenic risk for major psychiatric illness with subcortical volumes and white matter integrity in UK Biobank. <i>Scientific Reports</i> , 2017, 7, 42140.	1.6	98
348	An examination of the etiologic overlap between the genetic and environmental influences on insomnia and common psychopathology. <i>Depression and Anxiety</i> , 2017, 34, 453-462.	2.0	41
349	Personalized medicine: Genetic risk prediction of drug response. , 2017, 175, 75-90.		47
350	The NCAM1 gene set is linked to depressive symptoms and their brain structural correlates in healthy individuals. <i>Journal of Psychiatric Research</i> , 2017, 91, 116-123.	1.5	14
351	Association Between Telomere Length and Risk of Cancer and Non-Neoplastic Diseases. <i>JAMA Oncology</i> , 2017, 3, 636.	3.4	376
352	Pharmacogenetics of antidepressant response: A polygenic approach. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2017, 75, 128-134.	2.5	71
354	Polygenic risk for five psychiatric disorders and cross-disorder and disorder-specific neural connectivity in two independent populations. <i>NeuroImage: Clinical</i> , 2017, 14, 441-449.	1.4	81
355	Effects of Antenatal Maternal Depressive Symptoms and Socio-Economic Status on Neonatal Brain Development are Modulated by Genetic Risk. <i>Cerebral Cortex</i> , 2017, 27, 3080-3092.	1.6	90
356	Spatial and temporal expression patterns of genes around nine neuroticism-associated loci. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2017, 77, 164-171.	2.5	10
357	Assessing the presence of shared genetic architecture between Alzheimer's disease and major depressive disorder using genome-wide association data. <i>Translational Psychiatry</i> , 2017, 7, e1094-e1094.	2.4	38
358	Posttraumatic stress disorder symptom severity is associated with reduced default mode network connectivity in individuals with elevated genetic risk for psychopathology. <i>Depression and Anxiety</i> , 2017, 34, 632-640.	2.0	23
359	New directions for the treatment of depression: Targeting the photic regulation of arousal and mood (PRAM) pathway. <i>Depression and Anxiety</i> , 2017, 34, 588-595.	2.0	22
360	Interactions of early-life stress with the genome and epigenome: from prenatal stress to psychiatric disorders. <i>Current Opinion in Behavioral Sciences</i> , 2017, 14, 167-171.	2.0	18
361	Polymorphism of ERK/PTPRR Genes in Major Depressive Disorder at Resting-State Brain Function. <i>Developmental Neuropsychology</i> , 2017, 42, 231-240.	1.0	10
362	A comprehensive regional analysis of genome-wide expression profiles for major depressive disorder. <i>Journal of Affective Disorders</i> , 2017, 218, 86-92.	2.0	33
363	Inference in the age of big data: Future perspectives on neuroscience. <i>NeuroImage</i> , 2017, 155, 549-564.	2.1	161
364	Genome-wide analyses for personality traits identify six genomic loci and show correlations with psychiatric disorders. <i>Nature Genetics</i> , 2017, 49, 152-156.	9.4	350

#	ARTICLE	IF	CITATIONS
365	Resting-state connectivity biomarkers define neurophysiological subtypes of depression. <i>Nature Medicine</i> , 2017, 23, 28-38.	15.2	1,554
366	Genetic Approaches to Understanding Psychiatric Disease. <i>Neurotherapeutics</i> , 2017, 14, 564-581.	2.1	6
367	A continuum of genetic liability for minor and major depression. <i>Translational Psychiatry</i> , 2017, 7, e1131-e1131.	2.4	47
368	Pleiotropic Effects of Trait-Associated Genetic Variation on DNA Methylation: Utility for Refining GWAS Loci. <i>American Journal of Human Genetics</i> , 2017, 100, 954-959.	2.6	77
369	Runs of homozygosity, copy number variation, and risk for depression and suicidal behavior in an Arab Bedouin kindred. <i>Psychiatric Genetics</i> , 2017, 27, 169-177.	0.6	5
370	Polygenic risk for depression and the neural correlates of working memory in healthy subjects. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2017, 79, 67-76.	2.5	41
371	Mild traumatic brain injury is associated with reduced cortical thickness in those at risk for Alzheimer's disease. <i>Brain</i> , 2017, 140, aww344.	3.7	65
372	HPA Axis Genes, and Their Interaction with Childhood Maltreatment, are Related to Cortisol Levels and Stress-Related Phenotypes. <i>Neuropsychopharmacology</i> , 2017, 42, 2446-2455.	2.8	69
373	Recent advances in applying mass spectrometry and systems biology to determine brain dynamics. <i>Expert Review of Proteomics</i> , 2017, 14, 545-559.	1.3	12
374	Genome-wide association study of facial emotion recognition in children and association with polygenic risk for mental health disorders. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 701-711.	1.1	26
375	Genetics of Depression: Progress at Last. <i>Current Psychiatry Reports</i> , 2017, 19, 43.	2.1	101
376	Reproductive fitness and genetic risk of psychiatric disorders in the general population. <i>Nature Communications</i> , 2017, 8, 15833.	5.8	30
377	Genetic Heterogeneity in Depressive Symptoms Following the Death of a Spouse: Polygenic Score Analysis of the U.S. Health and Retirement Study. <i>American Journal of Psychiatry</i> , 2017, 174, 963-970.	4.0	37
378	The Relationship Between Mental Health, Disease Severity, and Genetic Risk for Depression in Early Rheumatoid Arthritis. <i>Psychosomatic Medicine</i> , 2017, 79, 638-645.	1.3	35
379	The association of childhood maltreatment with depression and anxiety is not moderated by the oxytocin receptor gene. <i>European Archives of Psychiatry and Clinical Neuroscience</i> , 2017, 267, 517-526.	1.8	32
380	Genetic effects influencing risk for major depressive disorder in China and Europe. <i>Translational Psychiatry</i> , 2017, 7, e1074-e1074.	2.4	64
381	Neurocognitive performance as an endophenotype for mood disorder subgroups. <i>Journal of Affective Disorders</i> , 2017, 215, 163-171.	2.0	13
382	Intergenerational Transmission of Maternal Childhood Maltreatment Exposure: Implications for Fetal Brain Development. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2017, 56, 373-382.	0.3	181

#	ARTICLE	IF	CITATIONS
383	Genetic correlation between amyotrophic lateral sclerosis and schizophrenia. <i>Nature Communications</i> , 2017, 8, 14774.	5.8	114
385	The Genetic Architecture of Major Depressive Disorder in Han Chinese Women. <i>JAMA Psychiatry</i> , 2017, 74, 162.	6.0	82
386	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.	0.7	84
387	Highly polygenic architecture of antidepressant treatment response: Comparative analysis of SSRI and NRI treatment in an animal model of depression. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 235-250.	1.1	10
388	Validity of LIDAS (Lifetime Depression Assessment Self-report): a self-report online assessment of lifetime major depressive disorder. <i>Psychological Medicine</i> , 2017, 47, 279-289.	2.7	29
389	Epigenetic and genetic variants in the HTR1B gene and clinical improvement in children and adolescents treated with fluoxetine. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2017, 75, 28-34.	2.5	28
390	Methylomic profiling of cortex samples from completed suicide cases implicates a role for PSORS1C3 in major depression and suicide. <i>Translational Psychiatry</i> , 2017, 7, e989-e989.	2.4	64
391	Genome-wide Regional Heritability Mapping Identifies a Locus Within the TOX2 Gene Associated With Major Depressive Disorder. <i>Biological Psychiatry</i> , 2017, 82, 312-321.	0.7	26
392	Copy Number Variation in Syndromic Forms of Psychiatric Illness: The Emerging Value of Clinical Genetic Testing in Psychiatry. <i>American Journal of Psychiatry</i> , 2017, 174, 1036-1050.	4.0	16
393	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.	1.1	17
394	Choosing Prediction Over Explanation in Psychology: Lessons From Machine Learning. <i>Perspectives on Psychological Science</i> , 2017, 12, 1100-1122.	5.2	1,063
395	Do regional brain volumes and major depressive disorder share genetic architecture? A study of Generation Scotland (n=19,762), UK Biobank (n=24,048) and the English Longitudinal Study of Ageing (n=5766). <i>Translational Psychiatry</i> , 2017, 7, e1205-e1205.	2.4	45
396	Genomewide association studies of suicide attempts in US soldiers. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 786-797.	1.1	52
397	Thyroid function and ischemic heart disease: a Mendelian randomization study. <i>Scientific Reports</i> , 2017, 7, 8515.	1.6	31
398	Gene-wide Association Study Reveals RNF122 Ubiquitin Ligase as a Novel Susceptibility Gene for Attention Deficit Hyperactivity Disorder. <i>Scientific Reports</i> , 2017, 7, 5407.	1.6	11
399	Expert and self-assessment of lifetime symptoms and diagnosis of major depressive disorder in large-scale genetic studies in the general population. <i>Psychiatric Genetics</i> , 2017, 27, 187-196.	0.6	10
400	Gene-based interaction analysis shows GABAergic genes interacting with parenting in adolescent depressive symptoms. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2017, 58, 1301-1309.	3.1	16
401	High-Coverage Whole-Exome Sequencing Identifies Candidate Genes for Suicide in Victims with Major Depressive Disorder. <i>Scientific Reports</i> , 2017, 7, 7106.	1.6	56

#	ARTICLE	IF	CITATIONS
402	The utility of empirically assigning ancestry groups in cross-population genetic studies of addiction. <i>American Journal on Addictions</i> , 2017, 26, 494-501.	1.3	46
403	Large normal-range TBP and ATXN7 CAG repeat lengths are associated with increased lifetime risk of depression. <i>Translational Psychiatry</i> , 2017, 7, e1143-e1143.	2.4	20
405	Neuronal-expressed microRNA-targeted pseudogenes compete with coding genes in the human brain. <i>Translational Psychiatry</i> , 2017, 7, e1199-e1199.	2.4	18
406	Polygenic Scores for Major Depressive Disorder and Risk of Alcohol Dependence. <i>JAMA Psychiatry</i> , 2017, 74, 1153.	6.0	73
407	Analysis of genome-wide association data highlights candidates for drug repositioning in psychiatry. <i>Nature Neuroscience</i> , 2017, 20, 1342-1349.	7.1	135
408	Common variants at 2q11.2, 8q21.3, and 11q13.2 are associated with major mood disorders. <i>Translational Psychiatry</i> , 2017, 7, 1273.	2.4	9
409	Huntingtin gene repeat size variations affect risk of lifetime depression. <i>Translational Psychiatry</i> , 2017, 7, 1277.	2.4	37
410	Genome-wide analysis in UK Biobank identifies four loci associated with mood instability and genetic correlation with major depressive disorder, anxiety disorder and schizophrenia. <i>Translational Psychiatry</i> , 2017, 7, 1264.	2.4	69
411	Genome-wide haplotype-based association analysis of major depressive disorder in Generation Scotland and UK Biobank. <i>Translational Psychiatry</i> , 2017, 7, 1263.	2.4	23
412	Association between neuropeptide Y receptor Y2 promoter variant rs6857715 and major depressive disorder. <i>Psychiatric Genetics</i> , 2017, 27, 34-37.	0.6	13
413	Interaction between the <i>FTO</i> gene, body mass index and depression: meta-analysis of 13701 individuals. <i>British Journal of Psychiatry</i> , 2017, 211, 70-76.	1.7	49
414	Expression quantitative trait loci (eQTLs) in human placentas suggest developmental origins of complex diseases. <i>Human Molecular Genetics</i> , 2017, 26, 3432-3441.	1.4	58
415	Investigating the relationship between iron and depression. <i>Journal of Psychiatric Research</i> , 2017, 94, 148-155.	1.5	10
416	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.	1.7	21
417	Predicting Cognitive Executive Functioning with Polygenic Risk Scores for Psychiatric Disorders. <i>Behavior Genetics</i> , 2017, 47, 11-24.	1.4	20
418	The PHF21B gene is associated with major depression and modulates the stress response. <i>Molecular Psychiatry</i> , 2017, 22, 1015-1025.	4.1	56
419	The genetic basis of the comorbidity between cannabis use and major depression. <i>Addiction</i> , 2017, 112, 113-123.	1.7	28
420	Investigating shared aetiology between type 2 diabetes and major depressive disorder in a population based cohort. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 227-234.	1.1	27

#	ARTICLE	IF	CITATIONS
421	Genetic Contributions of Inflammation to Depression. <i>Neuropsychopharmacology</i> , 2017, 42, 81-98.	2.8	174
422	Personalized medicine in psychiatry. <i>Nordic Journal of Psychiatry</i> , 2017, 71, 12-19.	0.7	46
423	<sc>HLA</sc> – <sc>DPB</sc> 1 variant rs3117242 is associated with anti–neutrophil cytoplasmic antibody–associated vasculitides in a Han Chinese population. <i>International Journal of Rheumatic Diseases</i> , 2017, 20, 1009-1015.	0.9	12
424	The Association of Genetic Predisposition to Depressive Symptoms with Non-suicidal and Suicidal Self-Injuries. <i>Behavior Genetics</i> , 2017, 47, 3-10.	1.4	24
425	Using Clinical Characteristics to Identify Which Patients With Major Depressive Disorder Have a Higher Genetic Load for Three Psychiatric Disorders. <i>Biological Psychiatry</i> , 2017, 81, 316-324.	0.7	31
426	Gene – Environment Interactions: From Molecular Mechanisms to Behavior. <i>Annual Review of Psychology</i> , 2017, 68, 215-241.	9.9	179
427	LD Hub: a centralized database and web interface to perform LD score regression that maximizes the potential of summary level GWAS data for SNP heritability and genetic correlation analysis. <i>Bioinformatics</i> , 2017, 33, 272-279.	1.8	822
428	Exome-sequencing in a large population-based study reveals a rare Asn396Ser variant in the LIPC gene associated with depressive symptoms. <i>Molecular Psychiatry</i> , 2017, 22, 537-543.	4.1	49
429	Why is depression more common among women than among men?. <i>Lancet Psychiatry</i> , 2017, 4, 146-158.	3.7	850
430	Genome-Wide Association Study of Loneliness Demonstrates a Role for Common Variation. <i>Neuropsychopharmacology</i> , 2017, 42, 811-821.	2.8	75
431	Can clinical subtypes contribute to genetic studies on major depression?. <i>Australasian Psychiatry</i> , 2017, 25, 633-634.	0.4	0
432	Plasma levels of the anti-coagulation protein C and the risk of ischaemic heart disease. <i>Thrombosis and Haemostasis</i> , 2017, 117, 262-268.	1.8	7
433	Statistical methods to detect pleiotropy in human complex traits. <i>Open Biology</i> , 2017, 7, 170125.	1.5	113
434	Significant Need for a French Network of Expert Centers Enabling a Better Characterization and Management of Treatment-Resistant Depression (Fondation FondaMental). <i>Frontiers in Psychiatry</i> , 2017, 8, 244.	1.3	11
435	Biomarkers for depression: recent insights, current challenges and future prospects. <i>Neuropsychiatric Disease and Treatment</i> , 2017, Volume 13, 1245-1262.	1.0	242
436	Multiple Testing in the Context of Gene Discovery in Sickle Cell Disease Using Genome-Wide Association Studies. <i>Genomics Insights</i> , 2017, 10, 117863101772117.	3.0	11
437	Epigenomics of Major Depressive Disorders and Schizophrenia: Early Life Decides. <i>International Journal of Molecular Sciences</i> , 2017, 18, 1711.	1.8	49
438	A bidirectional relationship between depression and the autoimmune disorders – New perspectives from the National Child Development Study. <i>PLoS ONE</i> , 2017, 12, e0173015.	1.1	76

#	ARTICLE	IF	CITATIONS
439	Social inequality, scientific inequality, and the future of mental illness. <i>Philosophy, Ethics, and Humanities in Medicine</i> , 2017, 12, 10.	0.7	13
440	Genetics of depressive symptoms in adolescence. <i>BMC Psychiatry</i> , 2017, 17, 321.	1.1	11
441	Investigating the genetic underpinnings of early-life irritability. <i>Translational Psychiatry</i> , 2017, 7, e1241-e1241.	2.4	42
442	Causal Inference in Psychopathology: A Systematic Review of Mendelian Randomisation Studies Aiming to Identify Environmental Risk Factors for Psychopathology. <i>Psychopathology Review</i> , 2017, a4, 4-25.	0.9	18
443	Association study of δ and μ promoter polymorphisms with suicide completers in the Japanese population. <i>Neuropsychiatric Disease and Treatment</i> , 2017, Volume 13, 899-908.	1.0	9
444	Genome-wide analysis of insomnia disorder. <i>Molecular Psychiatry</i> , 2018, 23, 2238-2250.	4.1	71
445	Genetic contribution to "theory of mind" in adolescence. <i>Scientific Reports</i> , 2018, 8, 3465.	1.6	24
446	Molecular Genetic Analysis Subdivided by Adversity Exposure Suggests Etiologic Heterogeneity in Major Depression. <i>American Journal of Psychiatry</i> , 2018, 175, 545-554.	4.0	69
447	Association of copy number variation across the genome with neuropsychiatric traits in the general population. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018, 177, 489-502.	1.1	26
448	Genome-wide association study of depression phenotypes in UK Biobank identifies variants in excitatory synaptic pathways. <i>Nature Communications</i> , 2018, 9, 1470.	5.8	415
449	The role of genetic variation of human metabolism for BMI, mental traits and mental disorders. <i>Molecular Metabolism</i> , 2018, 12, 1-11.	3.0	19
450	Association of Methylation Signals With Incident Coronary Heart Disease in an Epigenome-Wide Assessment of Circulating Tumor Necrosis Factor α . <i>JAMA Cardiology</i> , 2018, 3, 463.	3.0	33
451	Network Neuroscience: A Framework for Developing Biomarkers in Psychiatry. <i>Current Topics in Behavioral Neurosciences</i> , 2018, 40, 79-109.	0.8	16
452	Imaging and Genetic Biomarkers Predicting Transition to Psychosis. <i>Current Topics in Behavioral Neurosciences</i> , 2018, 40, 353-388.	0.8	13
453	Genome-wide analysis of adolescent psychotic-like experiences shows genetic overlap with psychiatric disorders. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018, 177, 416-425.	1.1	74
454	The effect of 5-HTTLPR and a serotonergic multi-marker score on amygdala, prefrontal and anterior cingulate cortex reactivity and habituation in a large, healthy fMRI cohort. <i>European Neuropsychopharmacology</i> , 2018, 28, 415-427.	0.3	25
455	Genome-wide analysis of self-reported risk-taking behaviour and cross-disorder genetic correlations in the UK Biobank cohort. <i>Translational Psychiatry</i> , 2018, 8, 39.	2.4	57
456	The use of polygenic risk scores to identify phenotypes associated with genetic risk of bipolar disorder and depression: A systematic review. <i>Journal of Affective Disorders</i> , 2018, 234, 148-155.	2.0	97

#	ARTICLE	IF	CITATIONS
457	Shared molecular neuropathology across major psychiatric disorders parallels polygenic overlap. <i>Science</i> , 2018, 359, 693-697.	6.0	851
458	Polygenic Risk for Depression Increases Risk of Ischemic Stroke. <i>Stroke</i> , 2018, 49, 543-548.	1.0	23
459	The long hard road to the doability of interdisciplinary research projects: the case of biosocial criminology. <i>New Genetics and Society</i> , 2018, 37, 21-43.	0.7	4
460	The Impact of COMT and Childhood Maltreatment on Suicidal Behaviour in Affective Disorders. <i>Scientific Reports</i> , 2018, 8, 692.	1.6	12
461	How can genetics help understand the relationship between cognitive dysfunction and schizophrenia?. <i>Scandinavian Journal of Psychology</i> , 2018, 59, 26-31.	0.8	9
462	Issues on the diagnosis and etiopathogenesis of mood disorders: reconsidering DSM-5. <i>Journal of Neural Transmission</i> , 2018, 125, 211-222.	1.4	17
463	Understanding the pathophysiology of depression: From monoamines to the neurogenesis hypothesis model - are we there yet?. <i>Behavioural Brain Research</i> , 2018, 341, 79-90.	1.2	219
464	The multimodal antidepressant vortioxetine may facilitate pyramidal cell firing by inhibition of 5-HT ₃ receptor expressing interneurons: An in vitro study in rat hippocampus slices. <i>Brain Research</i> , 2018, 1689, 1-11.	1.1	20
465	Multi-trait analysis of genome-wide association summary statistics using MTAG. <i>Nature Genetics</i> , 2018, 50, 229-237.	9.4	700
466	Gene-Environment Interplay and Individual Differences in Behavior. <i>Mind, Brain, and Education</i> , 2018, 12, 200-211.	0.9	26
467	Genome-wide meta-analyses of stratified depression in Generation Scotland and UK Biobank. <i>Translational Psychiatry</i> , 2018, 8, 9.	2.4	66
468	Beta-defensin 1, aryl hydrocarbon receptor and plasma kynurenine in major depressive disorder: metabolomics-informed genomics. <i>Translational Psychiatry</i> , 2018, 8, 10.	2.4	59
469	XPAT: a toolkit to conduct cross-platform association studies with heterogeneous sequencing datasets. <i>Nucleic Acids Research</i> , 2018, 46, e32-e32.	6.5	6
470	Biological Markers to Differentiate the Subtypes of Depression. , 2018, , 115-128.		0
471	Genome-Wide DNA Methylation Changes Associated with Intermittent Explosive Disorder: A Gene-Based Functional Enrichment Analysis. <i>International Journal of Neuropsychopharmacology</i> , 2018, 21, 12-20.	1.0	17
472	FoxO1, A2M, and TGF- β 1: three novel genes predicting depression in gene X environment interactions are identified using cross-species and cross-tissues transcriptomic and miRNomic analyses. <i>Molecular Psychiatry</i> , 2018, 23, 2192-2208.	4.1	73
473	Local adaptation in European populations affected the genetics of psychiatric disorders and behavioral traits. <i>Genome Medicine</i> , 2018, 10, 24.	3.6	5
474	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. <i>Nature Genetics</i> , 2018, 50, 668-681.	9.4	2,224

#	ARTICLE	IF	CITATIONS
475	Phenotypic and genetic analysis of cognitive performance in Major Depressive Disorder in the Generation Scotland: Scottish Family Health Study. <i>Translational Psychiatry</i> , 2018, 8, 63.	2.4	11
476	Polygenic Risk Scores in Clinical Psychology: Bridging Genomic Risk to Individual Differences. <i>Annual Review of Clinical Psychology</i> , 2018, 14, 119-157.	6.3	110
477	Genome-wide scan of depressive symptomatology in two representative cohorts in the United States and the United Kingdom. <i>Journal of Psychiatric Research</i> , 2018, 100, 63-70.	1.5	3
478	Largest GWAS of PTSD (N=20,070) yields genetic overlap with schizophrenia and sex differences in heritability. <i>Molecular Psychiatry</i> , 2018, 23, 666-673.	4.1	374
479	The protocadherin 17 gene affects cognition, personality, amygdala structure and function, synapse development and risk of major mood disorders. <i>Molecular Psychiatry</i> , 2018, 23, 400-412.	4.1	60
480	DNA methylation and clinical response to antidepressant medication in major depressive disorder: A review and recommendations. <i>Neuroscience Letters</i> , 2018, 669, 14-23.	1.0	54
481	Collaborative meta-analysis finds no evidence of a strong interaction between stress and 5-HTTLPR genotype contributing to the development of depression. <i>Molecular Psychiatry</i> , 2018, 23, 133-142.	4.1	247
482	A rare missense variant in RCL1 segregates with depression in extended families. <i>Molecular Psychiatry</i> , 2018, 23, 1120-1126.	4.1	34
483	Comparative genomic evidence for the involvement of schizophrenia risk genes in antipsychotic effects. <i>Molecular Psychiatry</i> , 2018, 23, 708-712.	4.1	27
484	Replicable and Coupled Changes in Innate and Adaptive Immune Gene Expression in Two Case-Control Studies of Blood Microarrays in Major Depressive Disorder. <i>Biological Psychiatry</i> , 2018, 83, 70-80.	0.7	158
485	A direct test of the diathesis-stress model for depression. <i>Molecular Psychiatry</i> , 2018, 23, 1590-1596.	4.1	187
486	Comorbidity classes and associated impairment, demographics and 9/11-exposures in 8,236 children and adolescents. <i>Journal of Psychiatric Research</i> , 2018, 96, 171-177.	1.5	7
487	Diagnostic and genetic overlap of three common mental disorders in structured interviews and health registries. <i>Acta Psychiatrica Scandinavica</i> , 2018, 137, 54-64.	2.2	45
488	The Gene Encoding Protocadherin 9 (PCDH9), a Novel Risk Factor for Major Depressive Disorder. <i>Neuropsychopharmacology</i> , 2018, 43, 1128-1137.	2.8	35
489	Transcriptome Alterations in Posttraumatic Stress Disorder. <i>Biological Psychiatry</i> , 2018, 83, 840-848.	0.7	36
490	Treatment resistant depression: A multi-scale, systems biology approach. <i>Neuroscience and Biobehavioral Reviews</i> , 2018, 84, 272-288.	2.9	319
491	Interaction between childhood maltreatment on immunogenetic risk in depression: Discovery and replication in clinical case-control samples. <i>Brain, Behavior, and Immunity</i> , 2018, 67, 203-210.	2.0	31
492	The role of <i>CLOCK</i> gene in psychiatric disorders: Evidence from human and animal research. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018, 177, 181-198.	1.1	50

#	ARTICLE	IF	CITATIONS
493	Trauma exposure interacts with the genetic risk of bipolar disorder in alcohol misuse of <sc>US</sc> soldiers. <i>Acta Psychiatrica Scandinavica</i> , 2018, 137, 148-156.	2.2	14
494	The impact of schizophrenia and mood disorder risk alleles on emotional problems: investigating change from childhood to middle age. <i>Psychological Medicine</i> , 2018, 48, 2153-2158.	2.7	24
495	Genetics of Post-traumatic Stress Disorder and Sleep Disturbance. , 2018, , 89-110.		0
496	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.	1.5	15
497	Genetic risk of major depressive disorder: the moderating and mediating effects of neuroticism and psychological resilience on clinical and self-reported depression. <i>Psychological Medicine</i> , 2018, 48, 1890-1899.	2.7	36
498	Association between a functional interleukin 6 receptor genetic variant and risk of depression and psychosis in a population-based birth cohort. <i>Brain, Behavior, and Immunity</i> , 2018, 69, 264-272.	2.0	86
499	The P2RX7 polymorphism rs2230912 is associated with depression: A meta-analysis. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2018, 82, 272-277.	2.5	34
500	Replication in Imaging Genetics: The Case of Threat-Related Amygdala Reactivity. <i>Biological Psychiatry</i> , 2018, 84, 148-159.	0.7	25
501	The cAMP responsive element-binding (CREB)-1 gene increases risk of major psychiatric disorders. <i>Molecular Psychiatry</i> , 2018, 23, 1957-1967.	4.1	38
502	Commentary for Special Issue of Prevention Science â€œUsing Genetics in Prevention: Science Fiction or Science Fact?â€. <i>Prevention Science</i> , 2018, 19, 101-108.	1.5	15
503	Serotonin in psychiatry: in vitro disease modeling using patient-derived neurons. <i>Cell and Tissue Research</i> , 2018, 371, 161-170.	1.5	36
504	Genetic Risks for Chronic Conditions: Implications for Long-term Wellbeing. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2018, 73, 477-483.	1.7	19
505	An Argument for the Foundations of Population Mental Health. <i>Frontiers in Psychiatry</i> , 2018, 9, 600.	1.3	7
506	Neuropeptide and Small Transmitter Coexistence: Fundamental Studies and Relevance to Mental Illness. <i>Frontiers in Neural Circuits</i> , 2018, 12, 106.	1.4	87
507	TPH-2 Gene Polymorphism in Major Depressive Disorder Patients With Early-Wakening Symptom. <i>Frontiers in Neuroscience</i> , 2018, 12, 827.	1.4	26
508	Neurobiological substrates underlying the effect of genomic risk for depression on the conversion of amnesic mild cognitive impairment. <i>Brain</i> , 2018, 141, 3457-3471.	3.7	18
509	Partial loss of psychiatric risk gene Mir137 in mice causes repetitive behavior and impairs sociability and learning via increased Pde10a. <i>Nature Neuroscience</i> , 2018, 21, 1689-1703.	7.1	127
510	Heritability informed power optimization (HIPO) leads to enhanced detection of genetic associations across multiple traits. <i>PLoS Genetics</i> , 2018, 14, e1007549.	1.5	36

#	ARTICLE	IF	CITATIONS
511	Comprehensive cross-disorder analyses of CNTNAP2 suggest it is unlikely to be a primary risk gene for psychiatric disorders. <i>PLoS Genetics</i> , 2018, 14, e1007535.	1.5	27
512	Dysregulation of the Lateral Habenula in Major Depressive Disorder. <i>Frontiers in Synaptic Neuroscience</i> , 2018, 10, 46.	1.3	71
513	Association of Genetic Variation at AQP4 Locus with Vascular Depression. <i>Biomolecules</i> , 2018, 8, 164.	1.8	14
514	A case-control genome-wide association study of ADHD discovers a novel association with the tenascin R (TNR) gene. <i>Translational Psychiatry</i> , 2018, 8, 284.	2.4	20
515	Precision pharmacotherapy: psychiatry's future direction in preventing, diagnosing, and treating mental disorders. <i>Pharmacogenomics and Personalized Medicine</i> , 2018, Volume 11, 211-222.	0.4	31
516	PPD ACT: an app-based genetic study of postpartum depression. <i>Translational Psychiatry</i> , 2018, 8, 260.	2.4	18
517	Dvl3 polymorphism interacts with life events and pro-inflammatory cytokines to influence major depressive disorder susceptibility. <i>Scientific Reports</i> , 2018, 8, 14181.	1.6	4
518	Phenotype-Specific Enrichment of Mendelian Disorder Genes near GWAS Regions across 62 Complex Traits. <i>American Journal of Human Genetics</i> , 2018, 103, 535-552.	2.6	90
519	Depression. <i>Lancet, The</i> , 2018, 392, 2299-2312.	6.3	2,026
520	Comparative Evaluation of Machine Learning Strategies for Analyzing Big Data in Psychiatry. <i>International Journal of Molecular Sciences</i> , 2018, 19, 3387.	1.8	18
521	Genetic correlations among psychiatric and immune-related phenotypes based on genome-wide association data. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018, 177, 641-657.	1.1	158
522	Variability in Resting State Network and Functional Network Connectivity Associated With Schizophrenia Genetic Risk: A Pilot Study. <i>Frontiers in Neuroscience</i> , 2018, 12, 114.	1.4	17
523	Additional common variants associated with type 2 diabetes and coronary artery disease detected using a pleiotropic cFDR method. <i>Journal of Diabetes and Its Complications</i> , 2018, 32, 1105-1112.	1.2	5
524	Rigor and reproducibility in genetic research on eating disorders. <i>International Journal of Eating Disorders</i> , 2018, 51, 593-607.	2.1	17
525	PRS-on-Spark (PRSoS): a novel, efficient and flexible approach for generating polygenic risk scores. <i>BMC Bioinformatics</i> , 2018, 19, 295.	1.2	20
526	Offspring of parents with mood disorders. <i>Current Opinion in Psychiatry</i> , 2018, 31, 349-357.	3.1	35
527	Method to estimate the approximate samples size that yield a certain number of significant GWAS signals in polygenic traits. <i>Genetic Epidemiology</i> , 2018, 42, 488-496.	0.6	2
528	PolyGEE: a generalized estimating equation approach to the efficient and robust estimation of polygenic effects in large-scale association studies. <i>Biostatistics</i> , 2018, 19, 295-306.	0.9	5

#	ARTICLE	IF	CITATIONS
529	Cross-species evidence from human and rat brain transcriptome for growth factor signaling pathway dysregulation in major depression. <i>Neuropsychopharmacology</i> , 2018, 43, 2134-2145.	2.8	25
530	Meta-analysis of genome-wide association studies for neuroticism in 449,484 individuals identifies novel genetic loci and pathways. <i>Nature Genetics</i> , 2018, 50, 920-927.	9.4	564
531	The GWAS Risk Genes for Depression May Be Actively Involved in Alzheimer's Disease. <i>Journal of Alzheimer's Disease</i> , 2018, 64, 1149-1161.	1.2	43
532	Sample Size for Successful Genome-Wide Association Study of Major Depressive Disorder. <i>Frontiers in Genetics</i> , 2018, 9, 227.	1.1	31
533	Absence of Wdr13 Gene Predisposes Mice to Mild Social Isolation – Chronic Stress, Leading to Depression-Like Phenotype Associated With Differential Expression of Synaptic Proteins. <i>Frontiers in Molecular Neuroscience</i> , 2018, 11, 133.	1.4	6
534	DNA Methylation Signatures of Depressive Symptoms in Middle-aged and Elderly Persons. <i>JAMA Psychiatry</i> , 2018, 75, 949.	6.0	78
535	Molecular genetic overlap between migraine and major depressive disorder. <i>European Journal of Human Genetics</i> , 2018, 26, 1202-1216.	1.4	56
536	Common variants on 6q16.2, 12q24.31 and 16p13.3 are associated with major depressive disorder. <i>Neuropsychopharmacology</i> , 2018, 43, 2146-2153.	2.8	36
537	Polygenic risk score: use in migraine research. <i>Journal of Headache and Pain</i> , 2018, 19, 29.	2.5	24
538	A systematic review of genome-wide research on psychotic experiences and negative symptom traits: new revelations and implications for psychiatry. <i>Human Molecular Genetics</i> , 2018, 27, R136-R152.	1.4	27
539	Genetic Study of White Matter Integrity in UK Biobank (N=8448) and the Overlap With Stroke, Depression, and Dementia. <i>Stroke</i> , 2018, 49, 1340-1347.	1.0	63
540	Circulating vitamin E levels and Alzheimer's disease: a Mendelian randomization study. <i>Neurobiology of Aging</i> , 2018, 72, 189.e1-189.e9.	1.5	53
541	Identification of expression quantitative trait loci associated with schizophrenia and affective disorders in normal brain tissue. <i>PLoS Genetics</i> , 2018, 14, e1007607.	1.5	34
542	Genetic association and meta-analysis of a schizophrenia GWAS variant rs10489202 in East Asian populations. <i>Translational Psychiatry</i> , 2018, 8, 144.	2.4	7
543	Interplay between the APOE Genotype and Possible Plasma Biomarkers in Alzheimer's Disease. <i>Current Alzheimer Research</i> , 2018, 15, 938-950.	0.7	15
544	Genetics Factors in Major Depression Disease. <i>Frontiers in Psychiatry</i> , 2018, 9, 334.	1.3	155
545	Shared genetic etiology between alcohol dependence and major depressive disorder. <i>Psychiatric Genetics</i> , 2018, 28, 66-70.	0.6	19
546	Biomarkers for Depression: Recent Insights, Current Challenges and Future Prospects. <i>Focus (American Psychiatric Publishing)</i> , 2018, 16, 194-209.	0.4	19

#	ARTICLE	IF	CITATIONS
547	DNA sequence-level analyses reveal potential phenotypic modifiers in a large family with psychiatric disorders. <i>Molecular Psychiatry</i> , 2018, 23, 2254-2265.	4.1	19
548	Further Evidence of an Association between <i>NCAN</i> rs1064395 and Bipolar Disorder. <i>Molecular Neuropsychiatry</i> , 2018, 4, 30-34.	3.0	10
549	Polygenic Scores for Neuropsychiatric Traits and White Matter Microstructure in the Pediatric Population. <i>Biological Psychiatry: Cognitive Neuroscience and Neuroimaging</i> , 2019, 4, 243-250.	1.1	11
550	Multivariate analysis of genome-wide data to identify potential pleiotropic genes for five major psychiatric disorders using MetaCCA. <i>Journal of Affective Disorders</i> , 2019, 242, 234-243.	2.0	24
551	Do metabolic HAD phosphatases moonlight as protein phosphatases?. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2019, 1866, 153-166.	1.9	23
552	GWAS of Behavioral Traits. <i>Current Topics in Behavioral Neurosciences</i> , 2019, 42, 1-34.	0.8	0
553	Practical recommendations to conduct a neuroimaging meta-analysis for neuropsychiatric disorders. <i>Human Brain Mapping</i> , 2019, 40, 5142-5154.	1.9	77
554	The genetic relationship between female reproductive traits and six psychiatric disorders. <i>Scientific Reports</i> , 2019, 9, 12041.	1.6	18
555	Polymorphisms in the <i>BDNF</i> and <i>BDNFOS</i> genes are associated with hypothalamus-pituitary axis regulation in major depression. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2019, 95, 109686.	2.5	29
556	Emotional Development and Depression. , 2019, , 695-748.		2
557	Protein Biomarkers in Major Depressive Disorder: An Update. <i>Advances in Experimental Medicine and Biology</i> , 2019, 1140, 585-600.	0.8	5
558	Neuroimaging insights into the link between depression and Insomnia: A systematic review. <i>Journal of Affective Disorders</i> , 2019, 258, 133-143.	2.0	44
559	HOPS: a quantitative score reveals pervasive horizontal pleiotropy in human genetic variation is driven by extreme polygenicity of human traits and diseases. <i>Genome Biology</i> , 2019, 20, 222.	3.8	47
560	The association of depression and anxiety with cardiac autonomic activity: The role of confounding effects of antidepressants. <i>Depression and Anxiety</i> , 2019, 36, 1163-1172.	2.0	36
561	Addressing heterogeneity (and homogeneity) in treatment mechanisms in depression and the potential to develop diagnostic and predictive biomarkers. <i>NeuroImage: Clinical</i> , 2019, 24, 101997.	1.4	16
562	Unraveling the genetic architecture of major depressive disorder: merits and pitfalls of the approaches used in genome-wide association studies. <i>Psychological Medicine</i> , 2019, 49, 2646-2656.	2.7	29
563	A model of human endogenous retrovirus (HERV) activation in mental health and illness. <i>Medical Hypotheses</i> , 2019, 133, 109404.	0.8	7
564	Depression genetic risk score is associated with anhedonia-related markers across units of analysis. <i>Translational Psychiatry</i> , 2019, 9, 236.	2.4	14

#	ARTICLE	IF	CITATIONS
565	A genome-wide association study of shared risk across psychiatric disorders implicates gene regulation during fetal neurodevelopment. <i>Nature Neuroscience</i> , 2019, 22, 353-361.	7.1	173
566	Genome-wide association analyses of chronotype in 697,828 individuals provides insights into circadian rhythms. <i>Nature Communications</i> , 2019, 10, 343.	5.8	417
567	Association of Polygenic Liabilities for Major Depression, Bipolar Disorder, and Schizophrenia With Risk for Depression in the Danish Population. <i>JAMA Psychiatry</i> , 2019, 76, 516.	6.0	78
568	Progressing Polygenic Medicine in Psychiatry Through Electronic Health Records. <i>JAMA Psychiatry</i> , 2019, 76, 470.	6.0	6
570	Genetics of Eating Disorders. <i>Psychiatric Clinics of North America</i> , 2019, 42, 59-73.	0.7	49
571	Quantifying between-cohort and between-sex genetic heterogeneity in major depressive disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019, 180, 439-447.	1.1	35
572	The bidirectional causal relationships of insomnia with five major psychiatric disorders: A Mendelian randomization study. <i>European Psychiatry</i> , 2019, 60, 79-85.	0.1	45
573	Examining sex differences in pleiotropic effects for depression and smoking using polygenic and gene-region aggregation techniques. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019, 180, 448-468.	1.1	5
574	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.	1.1	34
575	The involvement of sleep in the relationship between the serotonin transporter gene-linked polymorphic region (5-HTTLPR) and depression: A systematic review. <i>Journal of Affective Disorders</i> , 2019, 256, 205-212.	2.0	11
576	Novel susceptibility genes were found in a targeted sequencing of stroke patients with or without depression in the Chinese Han population. <i>Journal of Affective Disorders</i> , 2019, 255, 1-9.	2.0	11
577	Vitamin D and the Risk of Depression: A Causal Relationship? Findings from a Mendelian Randomization Study. <i>Nutrients</i> , 2019, 11, 1085.	1.7	45
578	Powerful and Efficient Strategies for Genetic Association Testing of Symptom and Questionnaire Data in Psychiatric Genetic Studies. <i>Scientific Reports</i> , 2019, 9, 7523.	1.6	2
579	Integrative analysis of Dupuytren's disease identifies novel risk locus and reveals a shared genetic etiology with BMI. <i>Genetic Epidemiology</i> , 2019, 43, 629-645.	0.6	13
580	Translational Potential of Neuroimaging Genomic Analyses to Diagnosis and Treatment in Mental Disorders. <i>Proceedings of the IEEE</i> , 2019, 107, 912-927.	16.4	4
581	Multi-tissue transcriptome analyses identify genetic mechanisms underlying neuropsychiatric traits. <i>Nature Genetics</i> , 2019, 51, 933-940.	9.4	77
582	Depression in neurodegenerative diseases: Common mechanisms and current treatment options. <i>Neuroscience and Biobehavioral Reviews</i> , 2019, 102, 56-84.	2.9	159
583	Clinical Genetic Testing and Counseling in Psychiatry. , 2019, , 181-202.		0

#	ARTICLE	IF	CITATIONS
584	Systems Approach to Identify Common Genes and Pathways Associated with Response to Selective Serotonin Reuptake Inhibitors and Major Depression Risk. <i>International Journal of Molecular Sciences</i> , 2019, 20, 1993.	1.8	14
585	Replicability analysis in genome-wide association studies via Cartesian hidden Markov models. <i>BMC Bioinformatics</i> , 2019, 20, 146.	1.2	8
586	Shared Molecular Neuropathology Across Major Psychiatric Disorders Parallels Polygenic Overlap. <i>Focus (American Psychiatric Publishing)</i> , 2019, 17, 66-72.	0.4	20
587	Genome-wide association analysis reveals KCTD12 and miR-383-binding genes in the background of rumination. <i>Translational Psychiatry</i> , 2019, 9, 119.	2.4	18
588	Genome-wide Burden of Rare Short Deletions Is Enriched in Major Depressive Disorder in Four Cohorts. <i>Biological Psychiatry</i> , 2019, 85, 1065-1073.	0.7	25
589	Genomic structural equation modelling provides insights into the multivariate genetic architecture of complex traits. <i>Nature Human Behaviour</i> , 2019, 3, 513-525.	6.2	511
590	Using Mendelian randomisation to assess causality in observational studies. <i>Evidence-Based Mental Health</i> , 2019, 22, 67-71.	2.2	49
591	Uncovering the Genetic Architecture of Major Depression. <i>Neuron</i> , 2019, 102, 91-103.	3.8	113
592	Polygenic Risk: Predicting Depression Outcomes in Clinical and Epidemiological Cohorts of Youths. <i>American Journal of Psychiatry</i> , 2019, 176, 615-625.	4.0	67
593	Co-shared genetics and possible risk gene pathway partially explain the comorbidity of schizophrenia, major depressive disorder, type 2 diabetes, and metabolic syndrome. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019, 180, 186-203.	1.1	86
594	Depression Depresses Vasodilation. <i>Circulation Research</i> , 2019, 124, 465-466.	2.0	1
595	Genome-wide association scan identifies new variants associated with a cognitive predictor of dyslexia. <i>Translational Psychiatry</i> , 2019, 9, 77.	2.4	82
596	Telomeres in neurological disorders. <i>Advances in Clinical Chemistry</i> , 2019, 90, 81-132.	1.8	32
597	Treating Depression in the Era of Precision Medicine: Challenges and Perspectives. , 2019, , 265-275.		3
598	Pediatric Depression. , 2019, , 415-424.		0
599	Integration of GWAS and brain eQTL identifies FLOT1 as a risk gene for major depressive disorder. <i>Neuropsychopharmacology</i> , 2019, 44, 1542-1551.	2.8	37
600	Integrative analyses of major histocompatibility complex loci in the genome-wide association studies of major depressive disorder. <i>Neuropsychopharmacology</i> , 2019, 44, 1552-1561.	2.8	27
601	The effect of genetic vulnerability and military deployment on the development of post-traumatic stress disorder and depressive symptoms. <i>European Neuropsychopharmacology</i> , 2019, 29, 405-415.	0.3	11

#	ARTICLE	IF	CITATIONS
602	Genome-Wide Association Study of Sleep Disturbances in Depressive Disorders. <i>Molecular Neuropsychiatry</i> , 2019, 5, 34-43.	3.0	1
603	Combining multi-population datasets for joint genome-wide association and meta-analyses: The case of bovine milk fat composition traits. <i>Journal of Dairy Science</i> , 2019, 102, 11124-11141.	1.4	12
604	Comparative genetic architectures of schizophrenia in East Asian and European populations. <i>Nature Genetics</i> , 2019, 51, 1670-1678.	9.4	440
606	Depressive Disorders: Mechanisms, Measurement and Management. <i>Advances in Experimental Medicine and Biology</i> , 2019, , .	0.8	6
607	Clinical association to FKBP5 rs1360780 in patients with depression. <i>Psychiatric Genetics</i> , 2019, 29, 220-225.	0.6	3
608	Insight into the genetic architecture of back pain and its risk factors from a study of 509,000 individuals. <i>Pain</i> , 2019, 160, 1361-1373.	2.0	74
609	A resting state fMRI analysis pipeline for pooling inference across diverse cohorts: an ENIGMA rs-fMRI protocol. <i>Brain Imaging and Behavior</i> , 2019, 13, 1453-1467.	1.1	49
610	Association of Whole-Genome and NETRIN1 Signaling Pathwayâ€Derived Polygenic Risk Scores for Major Depressive Disorder and White Matter Microstructure in the UK Biobank. <i>Biological Psychiatry: Cognitive Neuroscience and Neuroimaging</i> , 2019, 4, 91-100.	1.1	16
611	Brain proteome changes in female Brd1 mice unmask dendritic spine pathology and show enrichment for schizophrenia risk. <i>Neurobiology of Disease</i> , 2019, 124, 479-488.	2.1	14
612	HPA-axis multilocus genetic variation moderates associations between environmental stress and depressive symptoms among adolescents. <i>Development and Psychopathology</i> , 2019, 31, 1339-1352.	1.4	27
613	Sex differences in the genetic architecture of obsessiveâ€compulsive disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019, 180, 351-364.	1.1	41
614	Genomics of body fat percentage may contribute to sex bias in anorexia nervosa. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019, 180, 428-438.	1.1	87
615	Negr1 controls adult hippocampal neurogenesis and affective behaviors. <i>Molecular Psychiatry</i> , 2019, 24, 1189-1205.	4.1	59
616	Transmembrane protein 108 involves in adult neurogenesis in the hippocampal dentate gyrus. <i>Cell and Bioscience</i> , 2019, 9, 9.	2.1	25
617	Polygenic approaches to detect geneâ€environment interactions when external information is unavailable. <i>Briefings in Bioinformatics</i> , 2019, 20, 2236-2252.	3.2	14
618	The role of the individual in the coming era of process-based therapy. <i>Behaviour Research and Therapy</i> , 2019, 117, 40-53.	1.6	157
619	Differential associations of depressionâ€related phenotypes with cardiometabolic risks: Polygenic analyses and exploring shared genetic variants and pathways. <i>Depression and Anxiety</i> , 2019, 36, 330-344.	2.0	26
620	Continuity of genetic and environmental influences on clinically assessed major depression from ages 18 to 45. <i>Psychological Medicine</i> , 2019, 49, 2582-2590.	2.7	5

#	ARTICLE	IF	CITATIONS
621	Predictors of depressive symptoms in college students: A systematic review and meta-analysis of cohort studies. <i>Journal of Affective Disorders</i> , 2019, 244, 196-208.	2.0	88
622	Moody microbiome: Challenges and chances. <i>Journal of the Formosan Medical Association</i> , 2019, 118, S42-S54.	0.8	25
623	Genome-Wide Association Study Detected Novel Susceptibility Genes for Schizophrenia and Shared Trans-Populations/Diseases Genetic Effect. <i>Schizophrenia Bulletin</i> , 2019, 45, 824-834.	2.3	109
624	Polygenic risk for psychiatric disorders correlates with executive function in typical development. <i>Genes, Brain and Behavior</i> , 2019, 18, e12480.	1.1	16
625	Human brain arousal in the resting state: a genome-wide association study. <i>Molecular Psychiatry</i> , 2019, 24, 1599-1609.	4.1	26
626	Further evidence for the association between LRP8 and schizophrenia. <i>Schizophrenia Research</i> , 2020, 215, 499-505.	1.1	10
627	A joint study of whole exome sequencing and structural MRI analysis in major depressive disorder. <i>Psychological Medicine</i> , 2020, 50, 384-395.	2.7	19
628	Depression and coronary heart disease: 2018 position paper of the ESC working group on coronary pathophysiology and microcirculation. <i>European Heart Journal</i> , 2020, 41, 1687-1696.	1.0	203
629	Toward a framework for best practices and research guidelines for perinatal depression research. <i>Journal of Neuroscience Research</i> , 2020, 98, 1255-1267.	1.3	14
630	Genetic heterogeneity in self-reported depressive symptoms identified through genetic analyses of the PHQ-9. <i>Psychological Medicine</i> , 2020, 50, 2385-2396.	2.7	46
631	Genetic stratification of depression by neuroticism: revisiting a diagnostic tradition. <i>Psychological Medicine</i> , 2020, 50, 2526-2535.	2.7	27
632	Genetics of resilience: Implications from genome-wide association studies and candidate genes of the stress response system in posttraumatic stress disorder and depression. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2020, 183, 77-94.	1.1	54
633	Stratifying major depressive disorder by polygenic risk for schizophrenia in relation to structural brain measures. <i>Psychological Medicine</i> , 2020, 50, 1653-1662.	2.7	13
634	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. <i>Biological Psychiatry</i> , 2020, 87, 419-430.	0.7	27
635	Genomics of major depressive disorder. , 2020, , 187-200.		0
636	Evidence for causal effects of lifetime smoking on risk for depression and schizophrenia: a Mendelian randomisation study. <i>Psychological Medicine</i> , 2020, 50, 2435-2443.	2.7	324
637	Genetic correlations between pain phenotypes and depression and neuroticism. <i>European Journal of Human Genetics</i> , 2020, 28, 358-366.	1.4	52
638	Bivariate genome-wide association analyses of the broad depression phenotype combined with major depressive disorder, bipolar disorder or schizophrenia reveal eight novel genetic loci for depression. <i>Molecular Psychiatry</i> , 2020, 25, 1420-1429.	4.1	68

#	ARTICLE	IF	CITATIONS
639	Analyzing the genes and pathways related to major depressive disorder via a systems biology approach. <i>Brain and Behavior</i> , 2020, 10, e01502.	1.0	30
640	A study combining whole-exome sequencing and structural neuroimaging analysis for major depressive disorder. <i>Journal of Affective Disorders</i> , 2020, 262, 31-39.	2.0	10
641	Current progress and future direction in the genetics of PTSD: Focus on the development and contributions of the PGC-PTSD working group. , 2020, , 285-296.		0
642	The Psychiatric Genomics Consortium: History, development, and the future. , 2020, , 91-101.		6
643	Epigenetics of Major Depressive Disorder. , 2020, , 29-37.		1
644	Cannabis use, depression and self-harm: phenotypic and genetic relationships. <i>Addiction</i> , 2020, 115, 482-492.	1.7	29
645	Widespread Morphometric Abnormalities in Major Depression. <i>Neuroimaging Clinics of North America</i> , 2020, 30, 85-95.	0.5	4
646	Targeted exon sequencing in deceased schizophrenia patients in Denmark. <i>International Journal of Legal Medicine</i> , 2020, 134, 135-147.	1.2	2
647	Sex-specific association between infant caudate volumes and a polygenic risk score for major depressive disorder. <i>Journal of Neuroscience Research</i> , 2020, 98, 2529-2540.	1.3	10
648	Partial Support for an Interaction Between a Polygenic Risk Score for Major Depressive Disorder and Prenatal Maternal Depressive Symptoms on Infant Right Amygdalar Volumes. <i>Cerebral Cortex</i> , 2020, 30, 6121-6134.	1.6	21
649	Peripheral Markers of Depression. <i>Journal of Clinical Medicine</i> , 2020, 9, 3793.	1.0	99
650	Nature and Nurture: Effects of Affective Temperaments on Depressive Symptoms Are Markedly Modified by Stress Exposure. <i>Frontiers in Psychiatry</i> , 2020, 11, 599.	1.3	13
651	Genome-wide association study and polygenic risk score analysis of esketamine treatment response. <i>Scientific Reports</i> , 2020, 10, 12649.	1.6	24
652	Integrative omics analysis identifies differential biological pathways that are associated with regional grey matter volume changes in major depressive disorder. <i>Psychological Medicine</i> , 2022, 52, 924-935.	2.7	6
653	Common genetic risk variants identified in the SPARK cohort support DDHD2 as a candidate risk gene for autism. <i>Translational Psychiatry</i> , 2020, 10, 265.	2.4	56
654	Molecular Biological Aspects of Depressive Disorders: A Modern View. <i>Molecular Biology</i> , 2020, 54, 639-660.	0.4	7
655	Genome-wide association studies of antidepressant class response and treatment-resistant depression. <i>Translational Psychiatry</i> , 2020, 10, 360.	2.4	33
656	Genetics of Depressive Disorders: Candidate Genes and Genome-Wide Association Studies. <i>Russian Journal of Genetics</i> , 2020, 56, 903-915.	0.2	2

#	ARTICLE	IF	CITATIONS
657	Association of FKBP5 genotype with depressive symptoms in patients with coronary heart disease: a prospective study. <i>Journal of Neural Transmission</i> , 2020, 127, 1651-1662.	1.4	8
658	Depression-like behaviors induced by defective PTPRT activity through dysregulated synaptic functions and neurogenesis. <i>Journal of Cell Science</i> , 2020, 133, .	1.2	10
659	Neuropsychological and Psychiatric Features of Children and Adolescents Affected With Mitochondrial Diseases: A Systematic Review. <i>Frontiers in Psychiatry</i> , 2020, 11, 747.	1.3	12
660	Integrative genomics approach identifies conserved transcriptomic networks in Alzheimer's disease. <i>Human Molecular Genetics</i> , 2020, 29, 2899-2919.	1.4	50
661	Sleep, major depressive disorder, and Alzheimer disease. <i>Neurology</i> , 2020, 95, e1963-e1970.	1.5	45
662	The Emerging Role of Lhb CaMKII in the Comorbidity of Depressive and Alcohol Use Disorders. <i>International Journal of Molecular Sciences</i> , 2020, 21, 8123.	1.8	7
663	Genetic susceptibility, inflammation and specific types of depressive symptoms: evidence from the English Longitudinal Study of Ageing. <i>Translational Psychiatry</i> , 2020, 10, 140.	2.4	9
664	The Netrin-1/DCC Guidance Cue Pathway as a Molecular Target in Depression: Translational Evidence. <i>Biological Psychiatry</i> , 2020, 88, 611-624.	0.7	36
665	Heterogeneity and Polygenicity in Psychiatric Disorders: A Genome-Wide Perspective. <i>Chronic Stress</i> , 2020, 4, 247054702092484.	1.7	26
666	Human Herpesvirus 6B Greatly Increases Risk of Depression by Activating Hypothalamic-Pituitary-Adrenal Axis during Latent Phase of Infection. <i>IScience</i> , 2020, 23, 101187.	1.9	26
667	Epigenetic prediction of major depressive disorder. <i>Molecular Psychiatry</i> , 2021, 26, 5112-5123.	4.1	44
668	Genome-wide associations of human gut microbiome variation and implications for causal inference analyses. <i>Nature Microbiology</i> , 2020, 5, 1079-1087.	5.9	144
669	Nick Martin's Contribution to GxE Research. <i>Twin Research and Human Genetics</i> , 2020, 23, 131-134.	0.3	0
670	SLC1A3 C3590T but not BDNF G196A is a predisposition factor for stress as well as depression, in an adolescent eastern Indian population. <i>BMC Medical Genetics</i> , 2020, 21, 53.	2.1	3
671	Regulatory mechanisms of major depressive disorder risk variants. <i>Molecular Psychiatry</i> , 2020, 25, 1926-1945.	4.1	37
672	Neuroimaging Biomarkers for Predicting Treatment Response and Recurrence of Major Depressive Disorder. <i>International Journal of Molecular Sciences</i> , 2020, 21, 2148.	1.8	48
673	Minimal phenotyping yields genome-wide association signals of low specificity for major depression. <i>Nature Genetics</i> , 2020, 52, 437-447.	9.4	207
674	Sex differences in the genetic architecture of depression. <i>Scientific Reports</i> , 2020, 10, 9927.	1.6	50

#	ARTICLE	IF	CITATIONS
675	Traumas and Their Consequences According to Control-Mastery Theory. <i>Psychodynamic Psychiatry</i> , 2020, 48, 113-139.	0.1	17
676	Precision medicine in perinatal depression in light of the human microbiome. <i>Psychopharmacology</i> , 2020, 237, 915-941.	1.5	18
677	A polygenic predictor of treatment-resistant depression using whole exome sequencing and genome-wide genotyping. <i>Translational Psychiatry</i> , 2020, 10, 50.	2.4	33
678	Genomic influences on self-reported childhood maltreatment. <i>Translational Psychiatry</i> , 2020, 10, 38.	2.4	47
679	Large-Scale Exome Sequencing Study Implicates Both Developmental and Functional Changes in the Neurobiology of Autism. <i>Cell</i> , 2020, 180, 568-584.e23.	13.5	1,422
680	A Summary of Recent Updates on the Genetic Determinants of Depression. , 2020, , 1-27.		1
681	Hypothalamic-pituitary-adrenal axis-related genes and cognition in major mood disorders and schizophrenia: a systematic review. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2020, 101, 109929.	2.5	14
682	Genetic Contributions to Multivariate Data-Driven Brain Networks Constructed via Source-Based Morphometry. <i>Cerebral Cortex</i> , 2020, 30, 4899-4913.	1.6	7
683	HPA-axis multilocus genetic profile score moderates the association between maternal prenatal perceived stress and offspring depression in early adulthood. <i>Development and Psychopathology</i> , 2021, 33, 122-134.	1.4	10
684	Sex-differential DNA methylation and associated regulation networks in human brain implicated in the sex-biased risks of psychiatric disorders. <i>Molecular Psychiatry</i> , 2021, 26, 835-848.	4.1	47
685	Inflammatory modulation of the associations between prenatal maternal depression and neonatal brain. <i>Neuropsychopharmacology</i> , 2021, 46, 470-477.	2.8	13
686	An alternative approach to future diagnostic standards for major depressive disorder. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2021, 105, 110133.	2.5	10
687	Characterizing the effect of background selection on the polygenicity of brain-related traits. <i>Genomics</i> , 2021, 113, 111-119.	1.3	24
688	Shared transethnic genetic basis of panic disorder and psychiatric and related intermediate phenotypes. <i>European Neuropsychopharmacology</i> , 2021, 42, 87-96.	0.3	9
689	Genetic markers for depressive disorders with earlier age at onset. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2021, 108, 110176.	2.5	4
690	Sleep and high-risk behavior in military service members: a mega-analysis of four diverse U.S. Army units. <i>Sleep</i> , 2021, 44, .	0.6	21
691	Causal links between major depressive disorder and insomnia: A Mendelian randomisation study. <i>Gene</i> , 2021, 768, 145271.	1.0	25
692	Genetic analysis of endometriosis and depression identifies shared loci and implicates causal links with gastric mucosa abnormality. <i>Human Genetics</i> , 2021, 140, 529-552.	1.8	36

#	ARTICLE	IF	CITATIONS
693	Dissecting diagnostic heterogeneity in depression by integrating neuroimaging and genetics. <i>Neuropsychopharmacology</i> , 2021, 46, 156-175.	2.8	110
694	Ferroptosis-Related Genes in Neurodevelopment and Central Nervous System. <i>Biology</i> , 2021, 10, 35.	1.3	17
695	Challenges and Strategies for Current Classifications of Depressive Disorders: Proposal for Future Diagnostic Standards. <i>Advances in Experimental Medicine and Biology</i> , 2021, 1305, 103-116.	0.8	11
696	Genetic loci shared between major depression and intelligence with mixed directions of effect. <i>Nature Human Behaviour</i> , 2021, 5, 795-801.	6.2	23
697	Sirtuins, healthspan, and longevity in mammals. , 2021, , 77-149.		2
698	Medical epigenetics and twins. , 2021, , 103-116.		1
699	Genome-wide landscape of RNA-binding protein target site dysregulation reveals a major impact on psychiatric disorder risk. <i>Nature Genetics</i> , 2021, 53, 166-173.	9.4	49
700	From Package to Process: An Evidence-based Approach to Processes of Change in Psychotherapy. , 2021, , .		2
702	Interactions Between Glycogen Synthase Kinase-3 β Gene Polymorphisms, Negative Life Events, and Susceptibility to Major Depressive Disorder in a Chinese Population. <i>Frontiers in Psychiatry</i> , 2020, 11, 503477.	1.3	6
703	Major Depressive Disorder: Advances in Neuroscience Research and Translational Applications. <i>Neuroscience Bulletin</i> , 2021, 37, 863-880.	1.5	107
704	Genetics of eating disorders in the genome-wide era. <i>Psychological Medicine</i> , 2021, 51, 2287-2297.	2.7	17
706	Genome-wide analysis suggests the importance of vascular processes and neuroinflammation in late-life antidepressant response. <i>Translational Psychiatry</i> , 2021, 11, 127.	2.4	22
707	Can Molecular Biology Propose Reliable Biomarkers for Diagnosing Major Depression?. <i>Current Pharmaceutical Design</i> , 2021, 27, 305-318.	0.9	4
708	Toward an Extended Definition of Major Depressive Disorder Symptomatology: Digital Assessment and Cross-validation Study. <i>JMIR Formative Research</i> , 2021, 5, e27908.	0.7	1
709	Multiple measures of depression to enhance validity of major depressive disorder in the UK Biobank. <i>BJPsych Open</i> , 2021, 7, e44.	0.3	27
710	The genetic basis of major depression. <i>Psychological Medicine</i> , 2021, 51, 2217-2230.	2.7	65
711	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. <i>Biological Psychiatry</i> , 2022, 91, 102-117.	0.7	61
712	Functional enhancer elements drive subclass-selective expression from mouse to primate neocortex. <i>Cell Reports</i> , 2021, 34, 108754.	2.9	88

#	ARTICLE	IF	CITATIONS
713	Locus of Control and Negative Cognitive Styles in Adolescence as Risk Factors for Depression Onset in Young Adulthood: Findings From a Prospective Birth Cohort Study. <i>Frontiers in Psychology</i> , 2021, 12, 599240.	1.1	8
714	Serotonin 2A receptor polymorphism rs3803189 mediated by dynamics of default mode network: a potential biomarker for antidepressant early response. <i>Journal of Affective Disorders</i> , 2021, 283, 130-138.	2.0	4
715	Effects of polygenic risk score, childhood trauma and resilience on depressive symptoms in Chinese adolescents in a three-year cohort study. <i>Journal of Affective Disorders</i> , 2021, 282, 627-636.	2.0	13
716	The Genetics of Major Depression: Perspectives on the State of Research and Opportunities for Precision Medicine. <i>Psychiatric Annals</i> , 2021, 51, 165-169.	0.1	2
717	Multiple dimensions of stress vs. genetic effects on depression. <i>Translational Psychiatry</i> , 2021, 11, 254.	2.4	4
719	Cell-Type-Specific Gene Modules Related to the Regional Homogeneity of Spontaneous Brain Activity and Their Associations With Common Brain Disorders. <i>Frontiers in Neuroscience</i> , 2021, 15, 639527.	1.4	6
720	A Comparison of Ten Polygenic Score Methods for Psychiatric Disorders Applied Across Multiple Cohorts. <i>Biological Psychiatry</i> , 2021, 90, 611-620.	0.7	103
721	Dissecting polygenic signals from genome-wide association studies on human behaviour. <i>Nature Human Behaviour</i> , 2021, 5, 686-694.	6.2	57
723	Infant inhibited temperament in primates predicts adult behavior, is heritable, and is associated with anxiety-relevant genetic variation. <i>Molecular Psychiatry</i> , 2021, 26, 6609-6618.	4.1	13
724	Therapy response prediction in major depressive disorder: current and novel genomic markers influencing pharmacokinetics and pharmacodynamics. <i>Pharmacogenomics</i> , 2021, 22, 485-503.	0.6	12
725	American psychiatry in the new millennium: a critical appraisal. <i>Psychological Medicine</i> , 2021, 51, 2762-2770.	2.7	28
727	Prediction of Probable Major Depressive Disorder in the Taiwan Biobank: An Integrated Machine Learning and Genome-Wide Analysis Approach. <i>Journal of Personalized Medicine</i> , 2021, 11, 597.	1.1	6
728	Genetic Basis of Dual Diagnosis: A Review of Genome-Wide Association Studies (GWAS) Focusing on Patients with Mood or Anxiety Disorders and Co-Occurring Alcohol-Use Disorders. <i>Diagnostics</i> , 2021, 11, 1055.	1.3	8
730	The "missing heritability" Problem in psychiatry: Is the interaction of genetics, epigenetics and transposable elements a potential solution?. <i>Neuroscience and Biobehavioral Reviews</i> , 2021, 126, 23-42.	2.9	11
733	Identification of pleiotropy at the gene level between psychiatric disorders and related traits. <i>Translational Psychiatry</i> , 2021, 11, 410.	2.4	7
734	Cohort profile for the STRatifying Resilience and Depression Longitudinally (STRADL) study: A depression-focused investigation of Generation Scotland, using detailed clinical, cognitive, and neuroimaging assessments. <i>Wellcome Open Research</i> , 0, 4, 185.	0.9	12
735	The role of rs242941, rs1876828, rs242939 and rs110402 polymorphisms of CRHR1 gene and the depression: systematic review and meta-analysis. <i>Genes and Genomics</i> , 2021, 43, 1339-1349.	0.5	2
736	Discriminating Heterogeneous Trajectories of Resilience and Depression After Major Life Stressors Using Polygenic Scores. <i>JAMA Psychiatry</i> , 2021, 78, 744.	6.0	33

#	ARTICLE	IF	CITATIONS
737	The associations of <i>CNR1</i> SNPs and haplotypes with vulnerability and treatment response phenotypes in Han Chinese with major depressive disorder: A case-control association study. <i>Molecular Genetics & Genomic Medicine</i> , 2021, 9, e1752.	0.6	6
738	Genetically Proxied Diurnal Preference, Sleep Timing, and Risk of Major Depressive Disorder. <i>JAMA Psychiatry</i> , 2021, 78, 903.	6.0	31
739	Association study of six candidate genes with major depressive disorder in the North-Western population of Pakistan. <i>PLoS ONE</i> , 2021, 16, e0248454.	1.1	1
740	Cardiovascular Disease Risk Factors Among Children and Adolescents With Depression. <i>Frontiers in Psychiatry</i> , 2021, 12, 702737.	1.3	6
741	Associations between major psychiatric disorder polygenic risk scores and blood-based markers in UK biobank. <i>Brain, Behavior, and Immunity</i> , 2021, 97, 32-41.	2.0	9
742	Differential and spatial expression meta-analysis of genes identified in genome-wide association studies of depression. <i>Translational Psychiatry</i> , 2021, 11, 8.	2.4	22
743	Genetic Architecture of Depression: Where Do We Stand Now?. <i>Advances in Experimental Medicine and Biology</i> , 2021, 1305, 203-230.	0.8	4
744	Teorias evolucionistas da depressão: panorama e perspectivas. <i>Psicologia USP</i> , 0, 32, .	0.1	1
745	Stress and Emotions. , 2018, , 319-340.		1
746	Epigenetics of Psychopathology. , 2014, , 283-309.		4
747	Genetik und Gen-Umwelt-Interaktionen bei psychischen Erkrankungen. , 2017, , 147-191.		2
748	Introduction. <i>Advances in Experimental Medicine and Biology</i> , 2019, 1180, 1-17.	0.8	3
749	Genetic Markers in Psychiatry. <i>Advances in Experimental Medicine and Biology</i> , 2019, 1192, 53-93.	0.8	2
750	Additive genetic risk from five serotonin system polymorphisms interacts with interpersonal stress to predict depression.. <i>Journal of Abnormal Psychology</i> , 2015, 124, 776-790.	2.0	45
751	Stability and change in etiological factors for alcohol use disorder and major depression.. <i>Journal of Abnormal Psychology</i> , 2017, 126, 812-822.	2.0	17
752	Inhibition of attention for affective material: Contributions by HOMER1 gene variation.. <i>Psychology and Neuroscience</i> , 2015, 8, 495-508.	0.5	3
753	Genome-wide meta-analysis of depression identifies 102 independent variants and highlights the importance of the prefrontal brain regions. <i>Nature Neuroscience</i> , 2019, 22, 343-352.	7.1	1,589
796	Tulsa 1000: a naturalistic study protocol for multilevel assessment and outcome prediction in a large psychiatric sample. <i>BMJ Open</i> , 2018, 8, e016620.	0.8	88

#	ARTICLE	IF	CITATIONS
797	Brexanolone, a neurosteroid antidepressant, vindicates the GABAergic deficit hypothesis of depression and may foster resilience. <i>F1000Research</i> , 2019, 8, 751.	0.8	56
798	Cohort profile for the STRatifying Resilience and Depression Longitudinally (STRADL) study: A depression-focused investigation of Generation Scotland, using detailed clinical, cognitive, and neuroimaging assessments. <i>Wellcome Open Research</i> , 2019, 4, 185.	0.9	27
799	A comparison of a single genetic factor, two stress factors, and one psychosocial coping factor as predictors of depression in an Australian community sample.. <i>Archives of Psychiatry and Psychotherapy</i> , 2014, 16, 15-26.	0.2	1
800	Widespread signatures of positive selection in common risk alleles associated to autism spectrum disorder. <i>PLoS Genetics</i> , 2017, 13, e1006618.	1.5	73
801	Investigating the Causal Relationship of C-Reactive Protein with 32 Complex Somatic and Psychiatric Outcomes: A Large-Scale Cross-Consortium Mendelian Randomization Study. <i>PLoS Medicine</i> , 2016, 13, e1001976.	3.9	150
802	Genetic and Environmental Risk for Chronic Pain and the Contribution of Risk Variants for Major Depressive Disorder: A Family-Based Mixed-Model Analysis. <i>PLoS Medicine</i> , 2016, 13, e1002090.	3.9	60
803	Modulatory Effects of the Piccolo Genotype on Emotional Memory in Health and Depression. <i>PLoS ONE</i> , 2013, 8, e61494.	1.1	48
804	Dopamine Genetic Risk Score Predicts Depressive Symptoms in Healthy Adults and Adults with Depression. <i>PLoS ONE</i> , 2014, 9, e93772.	1.1	71
805	Variability in the Effect of 5-HTTLPR on Depression in a Large European Population: The Role of Age, Symptom Profile, Type and Intensity of Life Stressors. <i>PLoS ONE</i> , 2015, 10, e0116316.	1.1	28
806	5-HTTLPR Genotype Moderates the Effects of Past Ecstasy Use on Verbal Memory Performance in Adolescent and Emerging Adults: A Pilot Study. <i>PLoS ONE</i> , 2015, 10, e0134708.	1.1	4
807	Substance Use and Depression Symptomatology: Measurement Invariance of the Beck Depression Inventory (BDI-II) among Non-Users and Frequent-Users of Alcohol, Nicotine and Cannabis. <i>PLoS ONE</i> , 2016, 11, e0152118.	1.1	10
808	Genome-Wide Association of Heroin Dependence in Han Chinese. <i>PLoS ONE</i> , 2016, 11, e0167388.	1.1	30
811	Genetic overlap between type 2 diabetes and major depressive disorder identified by bioinformatics analysis. <i>Oncotarget</i> , 2016, 7, 17410-17414.	0.8	21
812	Genetic basis of depressive disorders. <i>Vavilovskii Zhurnal Genetiki i Seleksii</i> , 2019, 23, 465-472.	0.4	7
813	The Landscape of Pervasive Horizontal Pleiotropy in Human Genetic Variation is Driven by Extreme Polygenicity of Human Traits and Diseases. <i>SSRN Electronic Journal</i> , 0, , .	0.4	3
814	Mitochondria: A Connecting Link in the Major Depressive Disorder Jigsaw. <i>Current Neuropharmacology</i> , 2019, 17, 550-562.	1.4	29
815	Family history as an important factor for stratifying participants in genetic studies of major depression. <i>Balkan Journal of Medical Genetics</i> , 2018, 21, 5-12.	0.5	10
816	Epigenetic alterations in depression and antidepressant treatment. <i>Dialogues in Clinical Neuroscience</i> , 2014, 16, 395-404.	1.8	129

#	ARTICLE	IF	CITATIONS
817	Intermediate phenotypes and biomarkers of treatment outcome in major depressive disorder. Dialogues in Clinical Neuroscience, 2014, 16, 525-537.	1.8	32
818	Neuropsychiatric genomics in precision medicine: diagnostics, gene discovery, and translation. Dialogues in Clinical Neuroscience, 2016, 18, 237-252.	1.8	6
819	Seasonality Shows Evidence for Polygenic Architecture and Genetic Correlation With Schizophrenia and Bipolar Disorder. Journal of Clinical Psychiatry, 2015, 76, 128-134.	1.1	25
820	Novel association between TGFA, TGFB1, IRF1, PTGS2 and IKBKB single-nucleotide polymorphisms and occurrence, severity and treatment response of major depressive disorder. PeerJ, 2020, 8, e8676.	0.9	16
821	Geographic Clustering of Polygenic Scores at Different Stages of the Life Course. Rsf, 2018, 4, 137-149.	0.6	10
822	Impact of ACE2 genetic variant on antidepressant efficacy of SSRIs. Acta Neuropsychiatrica, 2022, 34, 30-36.	1.0	13
823	Molecular Link between Circadian Rhythmicity and Mood Disorders. Current Medicinal Chemistry, 2022, 29, 5692-5709.	1.2	5
824	The Candidate Schizophrenia Risk Gene Tmem108 Regulates Glucose Metabolism Homeostasis. Frontiers in Endocrinology, 2021, 12, 770145.	1.5	6
825	Precursors and correlates of transient and persistent longitudinal profiles of psychotic experiences from late childhood through early adulthood. British Journal of Psychiatry, 2021, , 1-9.	1.7	5
826	Future Directions in Genetics of Psychiatric Disorders. , 2014, , 311-337.		1
827	Influence of Serotonin Transporter Gene Polymorphisms and Adverse Life Events on Depressive Symptoms in the Elderly: A Population-Based Study. PLoS ONE, 2015, 10, e0143395.	1.1	3
830	Bipolar Illness. , 2016, , 53-78.		0
832	Genetik und Gen-Umwelt-Interaktionen bei psychischen Erkrankungen. , 2016, , 1-45.		0
841	Genetik und Gen-Umwelt-Interaktionen bei psychischen Erkrankungen. , 2017, , 1-45.		0
846	Uncoupling protein 2 haplotype does not affect human brain structure and function in a sample of community-dwelling older adults. PLoS ONE, 2017, 12, e0181392.	1.1	4
864	GENETICS OF MAJOR DEPRESSIVE DISORDER. Bulletin of Problems Biology and Medicine, 2019, 1, 40.	0.0	0
865	Genetic Advance in Depressive Disorder. Advances in Experimental Medicine and Biology, 2019, 1180, 19-57.	0.8	2
869	Common (Genetic) Links Between Clinics and the Community: New Evidence From a Tourette Syndrome Polygenic Score. Biological Psychiatry, 2019, 85, 281-282.	0.7	1

#	ARTICLE	IF	CITATIONS
871	The Multifaceted Legacy of the Human Genome Program for Evolutionary Biology: An Epistemological Perspective. <i>Perspectives on Science</i> , 2019, 27, 117-152.	0.3	3
876	Polymorphism and expression of the DVL3 gene in the etiology of depressive disorder. <i>Psychiatria Polska</i> , 2020, 54, 509-523.	0.2	1
878	Genetic liability in individuals at ultra-high risk of psychosis: A comparison study of 9 psychiatric traits. <i>PLoS ONE</i> , 2020, 15, e0243104.	1.1	3
879	Characterization of genome-wide association study data reveals spatiotemporal heterogeneity of mental disorders. <i>BMC Medical Genomics</i> , 2020, 13, 192.	0.7	8
880	Childhood Trauma in Depressive Disorders. , 2020, , 161-184.		2
883	Insight into the genetic architecture of back pain and its risk factors from a study of 509.000 individuals. <i>BMC Medical Genomics</i> , 2020, 20, 1-14.	0.1	0
884	What Have We Learned About the Genetics of Obsessive-Compulsive and Related Disorders in Recent Years?. <i>Focus (American Psychiatric Publishing)</i> , 2021, 19, 384-391.	0.4	2
885	Genome-wide analysis of 53,400 people with irritable bowel syndrome highlights shared genetic pathways with mood and anxiety disorders. <i>Nature Genetics</i> , 2021, 53, 1543-1552.	9.4	96
887	Association Analysis Between Common Variants of the TRPM1 Gene and Three Mental Disorders in the Han Chinese Population. <i>Genetic Testing and Molecular Biomarkers</i> , 2020, 24, 649-657.	0.3	1
889	Nanoparticle Based Gene Therapy Approach: A Pioneering Rebellion in the Management of Psychiatric Disorders. <i>Current Gene Therapy</i> , 2020, 20, 164-173.	0.9	6
890	Horizons of psychiatric genetics and epigenetics: where are we and where are we heading?. <i>Iranian Journal of Psychiatry and Behavioral Sciences</i> , 2014, 8, 1-10.	0.1	4
891	Why is Diagnosing MDD Challenging?. <i>Shanghai Archives of Psychiatry</i> , 2016, 28, 343-345.	0.7	2
892	Polygenic Risk Scales for the Development of Depression Using GWAS with Clinical Validation: Methodology and Study Design in the Russian Population. <i>Neuroscience and Behavioral Physiology</i> , 2021, 51, 1040-1049.	0.2	0
893	Potential Genetic Overlap Between Insomnia and Sleep Symptoms in Major Depressive Disorder: A Polygenic Risk Score Analysis. <i>Frontiers in Psychiatry</i> , 2021, 12, 734077.	1.3	2
894	Gene-gene interaction and new onset of major depressive disorder: Findings from a Chinese freshmen nested case-control study. <i>Journal of Affective Disorders</i> , 2022, 300, 505-510.	2.0	5
895	Chapitre 6. Génétique de la dépression et de l'épilepsie. , 2015, , 65-67.		0
896	Insights into the genomics of affective disorders. <i>Medizinische Genetik</i> , 2020, 32, 9-18.	0.1	2
898	Brain differential gene expression and blood cross-validation of a molecular signature of patients with major depressive disorder. <i>Psychiatric Genetics</i> , 2022, Publish Ahead of Print, .	0.6	3

#	ARTICLE	IF	CITATIONS
900	Body mass index interacts with a genetic-risk score for depression increasing the risk of the disease in high-susceptibility individuals. <i>Translational Psychiatry</i> , 2022, 12, 30.	2.4	4
901	Neurobiological Highlights of Cognitive Impairment in Psychiatric Disorders. <i>International Journal of Molecular Sciences</i> , 2022, 23, 1217.	1.8	26
902	The Gender-Specific Interaction of DVL3 and GSK3 β Polymorphisms on Major Depressive Disorder Susceptibility in a Chinese Han Population: A Case-Control Study. <i>Oxidative Medicine and Cellular Longevity</i> , 2022, 2022, 1-10.	1.9	3
904	Use of multiple polygenic risk scores for distinguishing schizophrenia-spectrum disorder and affective psychosis categories in a first-episode sample; the EU-GEI study. <i>Psychological Medicine</i> , 2023, 53, 3396-3405.	2.7	9
905	Emerging roles of long non-coding RNA in depression. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2022, 115, 110515.	2.5	16
906	Facial Emotion Recognition in Psychosis and Associations With Polygenic Risk for Schizophrenia: Findings From the Multi-Center EU-GEI Case-Control Study. <i>Schizophrenia Bulletin</i> , 2022, 48, 1104-1114.	2.3	9
907	The Potential of Polygenic Risk Scores to Predict Antidepressant Treatment Response in Major Depression: A Systematic Review. <i>Journal of Affective Disorders</i> , 2022, 304, 1-11.	2.0	16
908	Can epigenetics shine a light on the biological pathways underlying major mental disorders?. <i>Psychological Medicine</i> , 2022, 52, 1645-1665.	2.7	16
909	Sex Difference in Global Burden of Major Depressive Disorder: Findings From the Global Burden of Disease Study 2019. <i>Frontiers in Psychiatry</i> , 2022, 13, 789305.	1.3	14
910	The Involvement of Long Non-coding RNA and Messenger RNA Based Molecular Networks and Pathways in the Subacute Phase of Traumatic Brain Injury in Adult Mice. <i>Frontiers in Neuroinformatics</i> , 2022, 16, 794342.	1.3	3
911	The Molecular Basis of Spinocerebellar Ataxia Type 7. <i>Frontiers in Neuroscience</i> , 2022, 16, 818757.	1.4	5
912	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.	2.4	6
913	New Insights on Gene by Environmental Effects of Drugs of Abuse in Animal Models Using GeneNetwork. <i>Genes</i> , 2022, 13, 614.	1.0	4
914	Multi-Omics Characterization of Early- and Adult-Onset Major Depressive Disorder. <i>Journal of Personalized Medicine</i> , 2022, 12, 412.	1.1	7
915	Major Depressive Disorder: Existing Hypotheses about Pathophysiological Mechanisms and New Genetic Findings. <i>Genes</i> , 2022, 13, 646.	1.0	16
916	Transethnic meta-analysis of exome-wide variants identifies new loci associated with male-specific metabolic syndrome. <i>Genes and Genomics</i> , 2022, , 1.	0.5	0
917	Exome-wide screening identifies novel rare risk variants for major depression disorder. <i>Molecular Psychiatry</i> , 2022, 27, 3069-3074.	4.1	15
918	Incarceration, polygenic risk, and depressive symptoms among males in late adulthood. <i>Social Science Research</i> , 2022, 104, 102683.	1.1	1

#	ARTICLE	IF	CITATIONS
919	Reviewing the Role of the Endocannabinoid System in the Pathophysiology of Depression. <i>Frontiers in Pharmacology</i> , 2021, 12, 762738.	1.6	30
920	Sex differences in the genetic regulation of the blood transcriptome response to glucocorticoid receptor activation. <i>Translational Psychiatry</i> , 2021, 11, 632.	2.4	8
921	An Investigation into the Association Between Dopamine Receptor $D1$ Multilocus Genetic Variation, Multiparametric Magnetic Resonance Imaging, and Antidepressant Treatment. <i>Journal of Magnetic Resonance Imaging</i> , 2022, 56, 282-290.	1.9	1
939	Sleep disturbance and psychiatric disorders: a bidirectional Mendelian randomisation study. <i>Epidemiology and Psychiatric Sciences</i> , 2022, 31, e26.	1.8	24
940	Research Status of Related Influencing Factors of Depressive Disorder. <i>Advances in Clinical Medicine</i> , 2022, 12, 4246-4256.	0.0	0
941	Digital tools for the assessment of pharmacological treatment for depressive disorder: State of the art. <i>European Neuropsychopharmacology</i> , 2022, 60, 100-116.	0.3	8
942	The Immune System and Depression: From Epidemiological to Clinical Evidence. <i>Current Topics in Behavioral Neurosciences</i> , 2022, , 15-34.	0.8	3
943	Genome-wide association studies in non-anxiety individuals identified novel risk loci for depression. <i>European Psychiatry</i> , 2022, 65, .	0.1	1
944	Depression and anxiety in inflammatory bowel disease: epidemiology, mechanisms and treatment. <i>Nature Reviews Gastroenterology and Hepatology</i> , 2022, 19, 717-726.	8.2	114
946	L1 Retrotransposons: A Potential Endogenous Regulator for Schizophrenia. <i>Frontiers in Genetics</i> , 0, 13, .	1.1	4
947	Pursuing the Epidemiology and Familial Risks of Depression and Developing an Evidence Based Psychotherapy. <i>Psychiatry Research</i> , 2022, , 114754.	1.7	0
950	Index of multiple deprivation contributed to common psychiatric disorders: A systematic review and comprehensive analysis. <i>Neuroscience and Biobehavioral Reviews</i> , 2022, 140, 104806.	2.9	8
951	Insomnia and post-traumatic stress disorder: A meta-analysis on interrelated association ($n=57,618$) and prevalence ($n=573,665$). <i>Neuroscience and Biobehavioral Reviews</i> , 2022, 141, 104850.	2.9	15
952	Attempting to Increase the Effectiveness of the Antidepressant Trazodone Hydrochloride Drug Using $5\text{-HT}_2\text{A}$ -Receptors. <i>International Journal of Environmental Research and Public Health</i> , 2022, 19, 11281.	1.2	0
953	Regional gene expression signatures are associated with sex-specific functional connectivity changes in depression. <i>Nature Communications</i> , 2022, 13, .	5.8	16
954	Polygenic risk score and peer victimisation independently predict depressive symptoms in adolescence: results from the Quebec Longitudinal Study of Children Development. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2023, 64, 388-396.	3.1	2
955	Meals, Microbiota and Mental Health in Children and Adolescents (MMM-Study): A protocol for an observational longitudinal case-control study. <i>PLoS ONE</i> , 2022, 17, e0273855.	1.1	1
956	Genetic correlation between female infertility and mental health and lifestyle factors: A linkage disequilibrium score regression study. <i>Health Science Reports</i> , 2022, 5, .	0.6	1

#	ARTICLE	IF	CITATIONS
957	Predictive validity in drug discovery: what it is, why it matters and how to improve it. <i>Nature Reviews Drug Discovery</i> , 2022, 21, 915-931.	21.5	28
958	Molecular pathways of major depressive disorder converge on the synapse. <i>Molecular Psychiatry</i> , 2023, 28, 284-297.	4.1	84
959	Identify novel, shared and disorder-specific genetic architecture of major depressive disorder, insomnia and chronic pain. <i>Journal of Psychiatric Research</i> , 2022, 155, 511-517.	1.5	5
960	Polygenic risk scores for schizophrenia and major depression are associated with socio-economic indicators of adversity in two British community samples. <i>Translational Psychiatry</i> , 2022, 12, .	2.4	5
961	Understanding the genetics of peripartum depression: Research challenges, strategies, and opportunities. <i>Frontiers in Genetics</i> , 0, 13, .	1.1	0
963	Cerebral blood flow changes and their genetic mechanisms in major depressive disorder: a combined neuroimaging and transcriptome study. <i>Psychological Medicine</i> , 2023, 53, 6468-6480.	2.7	3
964	Transcriptional substrates of brain structural and functional impairments in drug-naïve first-episode patients with major depressive disorder. <i>Journal of Affective Disorders</i> , 2023, 325, 522-533.	2.0	6
965	The genetic basis of major depressive disorder. <i>Molecular Psychiatry</i> , 2023, 28, 2254-2265.	4.1	28
966	Deep phenotyping as a contribution to personalized depression therapy: the GEPaD and DaCFail protocols. <i>Journal of Neural Transmission</i> , 2023, 130, 707-722.	1.4	2
967	Identification of novel biomarkers linking depressive disorder and Alzheimer's disease based on an integrative bioinformatics analysis. <i>BMC Genomic Data</i> , 2023, 24, .	0.7	3
968	Association Between Single Nucleotide Polymorphisms Within lncRNA NONHSAT102891 and Depression Susceptibility in a Chinese Population. <i>Neuropsychiatric Disease and Treatment</i> , 0, Volume 19, 293-302.	1.0	2
969	Identification and investigation of depression-related molecular subtypes in inflammatory bowel disease and the anti-inflammatory mechanisms of paroxetine. <i>Frontiers in Immunology</i> , 0, 14, .	2.2	7
970	Genetic risk, muscle strength and risk of incident major depressive disorder: results from the UK Biobank. <i>Age and Ageing</i> , 2023, 52, .	0.7	0
971	Genomic analysis of firework fear and noise reactivity in standard poodles. <i>Canine Medicine and Genetics</i> , 2023, 10, .	1.4	1
972	Pathways link environmental and genetic factors with structural brain networks and psychopathology in youth. <i>Neuropsychopharmacology</i> , 0, , .	2.8	1
973	The Melanocortin System: A Promising Target for the Development of New Antidepressant Drugs. <i>International Journal of Molecular Sciences</i> , 2023, 24, 6664.	1.8	2
974	Gender differences in the association of polygenic risk and divergent depression trajectories from mid to late life: a national longitudinal study. <i>Biodemography and Social Biology</i> , 2023, 68, 32-53.	0.4	1
976	Leisure Activities, Genetic Risk, and Frailty: Evidence from the Chinese Adults Aged 80 Years or Older. <i>Gerontology</i> , 2023, 69, 961-971.	1.4	0

#	ARTICLE	IF	CITATIONS
993	Depression and depressive disorders. , 2023, , .		0
997	Genetic evidence for causal effects of immune dysfunction in psychiatric disorders: where are we?. Translational Psychiatry, 2024, 14, .	2.4	0
1000	Effects of sex and gender on the etiologies and presentation of select internalizing psychopathologies. Translational Psychiatry, 2024, 14, .	2.4	0