

Cathryn M Lewis

List of Publications by Year in descending order

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Version: 2024-02-01

418
papers

63,395
citations

3515

90
h-index

1044

234
g-index

507
all docs

507
docs citations

507
times ranked

55432
citing authors

#	ARTICLE	IF	CITATIONS
1	Depressive symptoms and neuroticism-related traits are the main factors associated with wellbeing independent of the history of lifetime depression in the UK Biobank. <i>Psychological Medicine</i> , 2023, 53, 3000-3008.	2.7	9
2	Depression with atypical neurovegetative symptoms shares genetic predisposition with immuno-metabolic traits and alcohol consumption. <i>Psychological Medicine</i> , 2022, 52, 726-736.	2.7	33
3	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. <i>Biological Psychiatry</i> , 2022, 91, 102-117.	0.7	61
4	Associations and limited shared genetic aetiology between bipolar disorder and cardiometabolic traits in the UK Biobank. <i>Psychological Medicine</i> , 2022, 52, 4039-4048.	2.7	10
5	Lack of Support for the Genes by Early Environment Interaction Hypothesis in the Pathogenesis of Schizophrenia. <i>Schizophrenia Bulletin</i> , 2022, 48, 20-26.	2.3	19
6	Using major depression polygenic risk scores to explore the depressive symptom continuum. <i>Psychological Medicine</i> , 2022, 52, 149-158.	2.7	9
7	Methylome-wide association study of early life stressors and adult mental health. <i>Human Molecular Genetics</i> , 2022, 31, 651-664.	1.4	7
8	Age-related changes in physiology in individuals with bipolar disorder. <i>Journal of Affective Disorders</i> , 2022, 296, 157-168.	2.0	17
9	Cross-classification between self-rated health and health status: longitudinal analyses of all-cause mortality and leading causes of death in the UK. <i>Scientific Reports</i> , 2022, 12, 459.	1.6	18
10	Anxiety disorders and age-related changes in physiology. <i>British Journal of Psychiatry</i> , 2022, 221, 528-537.	1.7	10
11	Latent subtypes of manic and/or irritable episode symptoms in two population-based cohorts. <i>British Journal of Psychiatry</i> , 2022, 221, 722-731.	1.7	4
12	A tool for translating polygenic scores onto the absolute scale using summary statistics. <i>European Journal of Human Genetics</i> , 2022, 30, 339-348.	1.4	18
13	Multi-phenotype analyses of hemostatic traits with cardiovascular events reveal novel genetic associations. <i>Journal of Thrombosis and Haemostasis</i> , 2022, 20, 1331-1349.	1.9	12
14	Polygenic scores in biomedical research. <i>Nature Reviews Genetics</i> , 2022, 23, 524-532.	7.7	69
15	Simulated distributions from negative experiments highlight the importance of the body mass index distribution in explaining depression-body mass index genetic risk score interactions. <i>International Journal of Epidemiology</i> , 2022, 51, 1581-1592.	0.9	2
16	Exploring polygenic-environment and residual-environment interactions for depressive symptoms within the UK Biobank. <i>Genetic Epidemiology</i> , 2022, 46, 219-233.	0.6	4
17	Lifestyle factors associated with sex differences in Kaposi sarcoma incidence among adult black South Africans: A case-control study. <i>Cancer Epidemiology</i> , 2022, 78, 102158.	0.8	1
18	Genetic architecture of 11 major psychiatric disorders at biobehavioral, functional genomic and molecular genetic levels of analysis. <i>Nature Genetics</i> , 2022, 54, 548-559.	9.4	101

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19	Epidemiology of Kaposi's sarcoma in sub-Saharan Africa. <i>Cancer Epidemiology</i> , 2022, 78, 102167.	0.8	14
20	Clarifying the causes of consistent and inconsistent findings in genetics. <i>Genetic Epidemiology</i> , 2022, 46, 372-389.	0.6	4
21	Clinical, demographic, and genetic risk factors of treatment-attributed suicidality in >10,000 Australian adults taking antidepressants. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2022, 189, 196-206.	1.1	2
22	Genetic copy number variants, cognition and psychosis: a meta-analysis and a family study. <i>Molecular Psychiatry</i> , 2021, 26, 5307-5319.	4.1	18
23	Association of extent of cannabis use and psychotic like intoxication experiences in a multi-national sample of first episode psychosis patients and controls. <i>Psychological Medicine</i> , 2021, 51, 2074-2082.	2.7	7
24	Daily use of high-potency cannabis is associated with more positive symptoms in first-episode psychosis patients: the EU-GEI case-control study. <i>Psychological Medicine</i> , 2021, 51, 1329-1337.	2.7	38
25	Drug repositioning for treatment-resistant depression: Hypotheses from a pharmacogenomic study. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2021, 104, 110050.	2.5	21
26	Examining the association between family status and depression in the UK Biobank. <i>Journal of Affective Disorders</i> , 2021, 279, 585-598.	2.0	22
27	Cost-effectiveness of genetic and clinical predictors for choosing combined psychotherapy and pharmacotherapy in major depression. <i>Journal of Affective Disorders</i> , 2021, 279, 722-729.	2.0	7
28	Delineating the Genetic Component of Gene Expression in Major Depression. <i>Biological Psychiatry</i> , 2021, 89, 627-636.	0.7	63
29	Investigating an in silico approach for prioritizing antidepressant drug prescription based on drug-induced expression profiles and predicted gene expression. <i>Pharmacogenomics Journal</i> , 2021, 21, 85-93.	0.9	1
30	Genetic risk for Alzheimer's disease, cognition, and mild behavioral impairment in healthy older adults. <i>Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring</i> , 2021, 13, e12164.	1.2	32
31	Imputed gene expression risk scores: a functionally informed component of polygenic risk. <i>Human Molecular Genetics</i> , 2021, 30, 727-738.	1.4	11
32	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
33	Genetic and clinical characteristics of treatment-resistant depression using primary care records in two UK cohorts. <i>Molecular Psychiatry</i> , 2021, 26, 3363-3373.	4.1	66
34	Genome-wide association study of suicidal behaviour severity in mood disorders. <i>World Journal of Biological Psychiatry</i> , 2021, 22, 1-19.	1.3	3
35	Genetic basis of lacunar stroke: a pooled analysis of individual patient data and genome-wide association studies. <i>Lancet Neurology</i> , The, 2021, 20, 351-361.	4.9	95
36	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. <i>Nature Genetics</i> , 2021, 53, 817-829.	9.4	629

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37	Evaluation of polygenic prediction methodology within a reference-standardized framework. <i>PLoS Genetics</i> , 2021, 17, e1009021.	1.5	99
38	The association between genetically determined ABO blood types and major depressive disorder. <i>Psychiatry Research</i> , 2021, 299, 113837.	1.7	4
39	Transcriptome-wide association study of treatment-resistant depression and depression subtypes for drug repurposing. <i>Neuropsychopharmacology</i> , 2021, 46, 1821-1829.	2.8	27
40	Elevated C-Reactive Protein in Patients With Depression, Independent of Genetic, Health, and Psychosocial Factors: Results From the UK Biobank. <i>American Journal of Psychiatry</i> , 2021, 178, 522-529.	4.0	110
41	Age-related changes in physiology in individuals with lifetime bipolar disorder. <i>BJPsych Open</i> , 2021, 7, S276-S276.	0.3	0
42	A Genome-Wide Association Study and Polygenic Risk Score Analysis of Posttraumatic Stress Disorder and Metabolic Syndrome in a South African Population. <i>Frontiers in Neuroscience</i> , 2021, 15, 677800.	1.4	6
43	Lifetime depression and age-related changes in body composition, cardiovascular function, grip strength and lung function: sex-specific analyses in the UK Biobank. <i>BJPsych Open</i> , 2021, 7, S44-S44.	0.3	0
44	Investigating Pleiotropy Between Depression and Autoimmune Diseases Using the UK Biobank. <i>Biological Psychiatry Global Open Science</i> , 2021, 1, 48-58.	1.0	29
45	Exploring the genetic heterogeneity in major depression across diagnostic criteria. <i>Molecular Psychiatry</i> , 2021, 26, 7337-7345.	4.1	18
46	Lifetime depression and age-related changes in body composition, cardiovascular function, grip strength and lung function: sex-specific analyses in the UK Biobank. <i>Aging</i> , 2021, 13, 17038-17079.	1.4	20
47	The continuity of effect of schizophrenia polygenic risk score and patterns of cannabis use on transdiagnostic symptom dimensions at first-episode psychosis: findings from the EU-GEI study. <i>Translational Psychiatry</i> , 2021, 11, 423.	2.4	12
48	242Exploring health in the UK Biobank: cross-sectional analyses of sociodemographic, psychosocial, lifestyle and environmental factors. <i>International Journal of Epidemiology</i> , 2021, 50, .	0.9	0
49	Transcriptome-based polygenic score links depression-related corticolimbic gene expression changes to sex-specific brain morphology and depression risk. <i>Neuropsychopharmacology</i> , 2021, 46, 2304-2311.	2.8	5
50	Examining sex differences in neurodevelopmental and psychiatric genetic risk in anxiety and depression. <i>PLoS ONE</i> , 2021, 16, e0248254.	1.1	4
51	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. <i>JAMA Psychiatry</i> , 2021, 78, 1258.	6.0	88
52	Polygenic Risk and the Course of Attention-Deficit/Hyperactivity Disorder From Childhood to Young Adulthood: Findings From a Nationally Representative Cohort. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2021, 60, 1147-1156.	0.3	28
53	Investigating rare pathogenic/likely pathogenic exonic variation in bipolar disorder. <i>Molecular Psychiatry</i> , 2021, 26, 5239-5250.	4.1	15
54	Exploring health in the UK Biobank: associations with sociodemographic characteristics, psychosocial factors, lifestyle and environmental exposures. <i>BMC Medicine</i> , 2021, 19, 240.	2.3	36

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55	Vitamin D and the risk of treatment-resistant and atypical depression: A Mendelian randomization study. <i>Translational Psychiatry</i> , 2021, 11, 561.	2.4	9
56	The genetic case for cardiorespiratory fitness as a clinical vital sign and the routine prescription of physical activity in healthcare. <i>Genome Medicine</i> , 2021, 13, 180.	3.6	16
57	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
58	Cardiovascular health and risk of hospitalization with COVID-19: A Mendelian Randomization study. <i>JRSM Cardiovascular Disease</i> , 2021, 10, 204800402110593.	0.4	5
59	Ranking lifestyle risk factors for cervical cancer among Black women: A case-control study from Johannesburg, South Africa. <i>PLoS ONE</i> , 2021, 16, e0260319.	1.1	5
60	Genome-wide association study of antidepressant treatment resistance in a population-based cohort using health service prescription data and meta-analysis with GENDEP. <i>Pharmacogenomics Journal</i> , 2020, 20, 329-341.	0.9	45
61	The Maudsley environmental risk score for psychosis. <i>Psychological Medicine</i> , 2020, 50, 2213-2220.	2.7	42
62	Letter to the editor: Is polygenic risk for Parkinson's disease associated with less risk of first episode psychosis?. <i>Psychological Medicine</i> , 2020, 50, 173-176.	2.7	1
63	Genetic and Inflammatory Biomarkers Classify Small Intestine Inflammation in Asymptomatic First-degree Relatives of Patients With Crohn's Disease. <i>Clinical Gastroenterology and Hepatology</i> , 2020, 18, 908-916.e13.	2.4	18
64	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
65	Cannabis use, depression and self-harm: phenotypic and genetic relationships. <i>Addiction</i> , 2020, 115, 482-492.	1.7	29
66	S126. THE RELATION OF THE PSYCHOSIS CONTINUUM WITH SCHIZOPHRENIA POLYGENIC RISK SCORE AND CANNABIS USE. <i>Schizophrenia Bulletin</i> , 2020, 46, S83-S83.	2.3	0
67	Genetic comorbidity between major depression and cardio-metabolic traits, stratified by age at onset of major depression. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2020, 183, 309-330.	1.1	33
68	Genetics and major depressive disorder: clinical implications for disease risk, prognosis and treatment. <i>International Clinical Psychopharmacology</i> , 2020, 35, 233-242.	0.9	22
69	Genome-Wide Association Study Meta-Analysis of Stroke in 22 000 Individuals of African Descent Identifies Novel Associations With Stroke. <i>Stroke</i> , 2020, 51, 2454-2463.	1.0	26
70	Multifactorial disorders and polygenic risk scores: predicting common diseases and the possibility of adverse selection in life and protection insurance. <i>Annals of Actuarial Science</i> , 2020, , 1-16.	1.0	8
71	Genome-wide association study of MRI markers of cerebral small vessel disease in 42,310 participants. <i>Nature Communications</i> , 2020, 11, 2175.	5.8	93
72	Polygenic risk scores: from research tools to clinical instruments. <i>Genome Medicine</i> , 2020, 12, 44.	3.6	646

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73	Studying individual risk factors for self-harm in the UK Biobank: A polygenic scoring and Mendelian randomisation study. <i>PLoS Medicine</i> , 2020, 17, e1003137.	3.9	34
74	Genetic stratification of depression in UK Biobank. <i>Translational Psychiatry</i> , 2020, 10, 163.	2.4	19
75	Johannesburg Cancer Study (JCS): contribution to knowledge and opportunities arising from 20 years of data collection in an African setting. <i>Cancer Epidemiology</i> , 2020, 65, 101701.	0.8	11
76	Minimal phenotyping yields genome-wide association signals of low specificity for major depression. <i>Nature Genetics</i> , 2020, 52, 437-447.	9.4	207
77	Impute.me: An Open-Source, Non-profit Tool for Using Data From Direct-to-Consumer Genetic Testing to Calculate and Interpret Polygenic Risk Scores. <i>Frontiers in Genetics</i> , 2020, 11, 578.	1.1	47
78	CYP2D6 Revisited in GENDEP: Inter-Platform Concordance. <i>Biological Psychiatry</i> , 2020, 87, S148.	0.7	0
79	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
80	The behavioral, cellular and immune mediators of HIV-1 acquisition: New insights from population genetics. <i>Scientific Reports</i> , 2020, 10, 3304.	1.6	8
81	Genome-wide gene-environment analyses of major depressive disorder and reported lifetime traumatic experiences in UK Biobank. <i>Molecular Psychiatry</i> , 2020, 25, 1430-1446.	4.1	116
82	Genetic Associations Between Childhood Psychopathology and Adult Depression and Associated Traits in 42,998 Individuals. <i>JAMA Psychiatry</i> , 2020, 77, 715.	6.0	56
83	Treatment response classes in major depressive disorder identified by model-based clustering and validated by clinical prediction models. <i>Translational Psychiatry</i> , 2019, 9, 187.	2.4	51
84	Gene Variants at Loci Related to Blood Pressure Account for Variation in Response to Antihypertensive Drugs Between Black and White Individuals. <i>Hypertension</i> , 2019, 74, 614-622.	1.3	14
85	F105AN EXOME SEQUENCING STUDY IN TREATMENT-RESISTANT DEPRESSION. <i>European Neuropsychopharmacology</i> , 2019, 29, S1166-S1167.	0.3	0
86	Genetic associations with radiological damage in rheumatoid arthritis: Meta-analysis of seven genome-wide association studies of 2,775 cases. <i>PLoS ONE</i> , 2019, 14, e0223246.	1.1	11
87	SU106IS GENETIC LIABILITY TO SCHIZOPHRENIA MAPPING TO THE PSYCHOSIS CONTINUUM MODEL?. <i>European Neuropsychopharmacology</i> , 2019, 29, S1322-S1323.	0.3	0
88	META-ANALYSIS OF CYP2C19 ASSOCIATION WITH EFFICACY AND SIDE EFFECTS OF CITALOPRAM AND ESCITALOPRAM USING DATA FROM GENOME-WIDE ASSOCIATION STUDIES. <i>European Neuropsychopharmacology</i> , 2019, 29, S808.	0.3	0
89	1GENETIC COMORBIDITY BETWEEN DEPRESSION AND CARDIO-METABOLIC DISEASE, STRATIFIED BY AGE AT ONSET. <i>European Neuropsychopharmacology</i> , 2019, 29, S1066.	0.3	1
90	4DETERMINING THE RELATIONSHIP BETWEEN CANNABIS USE AND MAJOR DEPRESSION IN UK BIOBANK. <i>European Neuropsychopharmacology</i> , 2019, 29, S1067-S1068.	0.3	0

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91	A PGC-MDD COLLABORATION ON THE GENETICS OF TREATMENT OUTCOMES IN DEPRESSION. <i>European Neuropsychopharmacology</i> , 2019, 29, S1044.	0.3	0
92	GENETIC TESTING TO PREDICT ONSET AND TREATMENT OF PSYCHIATRIC DISORDERS—2018 UPDATE. <i>European Neuropsychopharmacology</i> , 2019, 29, S1063.	0.3	0
93	Progressing Polygenic Medicine in Psychiatry Through Electronic Health Records. <i>JAMA Psychiatry</i> , 2019, 76, 470.	6.0	6
94	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
95	Secondary Analysis of PGC Data: Examples And Inspiration. <i>European Neuropsychopharmacology</i> , 2019, 29, S714-S715.	0.3	0
96	ukbtools: An R package to manage and query UK Biobank data. <i>PLoS ONE</i> , 2019, 14, e0214311.	1.1	37
97	Prevalence of Advanced Liver Fibrosis in Patients With Severe Psoriasis. <i>JAMA Dermatology</i> , 2019, 155, 1028.	2.0	17
98	GWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. <i>American Journal of Psychiatry</i> , 2019, 176, 651-660.	4.0	186
99	GENOME-WIDE ASSOCIATION STUDY OF SUICIDE ATTEMPT IN MAJOR PSYCHIATRIC DISORDERS. <i>European Neuropsychopharmacology</i> , 2019, 29, S820-S821.	0.3	0
100	Genome-wide survey of copy number variants finds MAPT duplications in progressive supranuclear palsy. <i>Movement Disorders</i> , 2019, 34, 1049-1059.	2.2	24
101	Polygenic risk for circulating reproductive hormone levels and their influence on hippocampal volume and depression susceptibility. <i>Psychoneuroendocrinology</i> , 2019, 106, 284-292.	1.3	18
102	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019, 51, 793-803.	9.4	1,191
103	The contribution of cannabis use to variation in the incidence of psychotic disorder across Europe (EU-GEI): a multicentre case-control study. <i>Lancet Psychiatry</i> , 2019, 6, 427-436.	3.7	528
104	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
105	O3.1. ASSOCIATION OF EXTENT OF CANNABIS USE AND ACUTE INTOXICATION EXPERIENCES IN A MULTI-NATIONAL SAMPLE OF FIRST EPISODE PSYCHOSIS PATIENTS AND CONTROLS. <i>Schizophrenia Bulletin</i> , 2019, 45, S165-S166.	2.3	0
106	Evidence of causal effect of major depression on alcohol dependence: findings from the psychiatric genomics consortium. <i>Psychological Medicine</i> , 2019, 49, 1218-1226.	2.7	74
107	Uncovering the Genetic Architecture of Major Depression. <i>Neuron</i> , 2019, 102, 91-103.	3.8	113
108	Association of genetic variants in CHEK2 with oesophageal squamous cell carcinoma in the South African Black population. <i>Carcinogenesis</i> , 2019, 40, 513-520.	1.3	13

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109	T76FEASIBILITY OF DEFINING TREATMENT RESISTANT DEPRESSION IN PRIMARY CARE RECORDS FOR PHARMACOGENETIC STUDIES. <i>European Neuropsychopharmacology</i> , 2019, 29, S256.	0.3	0
110	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , 2019, 179, 1469-1482.e11.	13.5	935
111	Association of Whole-Genome and NETRIN1 Signaling Pathwayâ€œDerived 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
112	The polygenic nature of telomere length and the anti-ageing properties of lithium. <i>Neuropsychopharmacology</i> , 2019, 44, 757-765.	2.8	45
113	Using genetics to understand the causal influence of higher BMI on depression. <i>International Journal of Epidemiology</i> , 2019, 48, 834-848.	0.9	156
114	Genome-wide association study of treatment-resistance in depression and meta-analysis of three independent samples. <i>British Journal of Psychiatry</i> , 2019, 214, 36-41.	1.7	44
115	Telomere length as a predictor of emotional processing in the brain. <i>Human Brain Mapping</i> , 2019, 40, 1750-1759.	1.9	16
116	Transdiagnostic dimensions of psychopathology at first episode psychosis: findings from the multinational EU-GEI study. <i>Psychological Medicine</i> , 2019, 49, 1378-1391.	2.7	69
117	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
118	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
119	Antidepressant drug-specific prediction of depression treatment outcomes from genetic and clinical variables. <i>Scientific Reports</i> , 2018, 8, 5530.	1.6	51
120	Association analysis in over 329,000 individuals identifies 116 independent variants influencing neuroticism. <i>Nature Genetics</i> , 2018, 50, 6-11.	9.4	327
121	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
122	Does Childhood Trauma Moderate Polygenic Risk for Depression? A Meta-analysis of 5765 Subjects From the Psychiatric Genomics Consortium. <i>Biological Psychiatry</i> , 2018, 84, 138-147.	0.7	87
123	A polygenic risk score analysis of psychosis endophenotypes across brain functional, structural, and cognitive domains. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018, 177, 21-34.	1.1	57
124	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
125	Genome-wide gene-environment interaction in depression: A systematic evaluation of candidate genes. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018, 177, 40-49.	1.1	55
126	New insights into the pharmacogenomics of antidepressant response from the GENDEP and STAR*D studies: rare variant analysis and high-density imputation. <i>Pharmacogenomics Journal</i> , 2018, 18, 413-421.	0.9	40

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127	Transforming Summary Statistics from Logistic Regression to the Liability Scale: Application to Genetic and Environmental Risk Scores. <i>Human Heredity</i> , 2018, 83, 210-224.	0.4	21
128	5.4 BIOLOGICAL AND EPIDEMIOLOGICAL EXAMINATION OF TRANSDIAGNOSTIC AND SPECIFIC SYMPTOM DIMENSIONS AT PSYCHOSIS ONSET: FINDINGS FROM THE EUGEI STUDY. <i>Schizophrenia Bulletin</i> , 2018, 44, S7-S7.	2.3	2
129	Genetic Risk for Psychiatric Disorders and Telomere Length. <i>Frontiers in Genetics</i> , 2018, 9, 468.	1.1	20
130	Expression quantitative trait loci in the developing human brain and their enrichment in neuropsychiatric disorders. <i>Genome Biology</i> , 2018, 19, 194.	3.8	126
131	Polygenic risk scores for major depressive disorder and neuroticism as predictors of antidepressant response: Meta-analysis of three treatment cohorts. <i>PLoS ONE</i> , 2018, 13, e0203896.	1.1	37
132	F132. The Effect of Telomere Length and Their Polygenic Risk Scores on Emotional Brain Function and Connectivity. <i>Biological Psychiatry</i> , 2018, 83, S289.	0.7	0
133	Exome array analysis of adverse reactions to fluoropyrimidine-based therapy for gastrointestinal cancer. <i>PLoS ONE</i> , 2018, 13, e0188911.	1.1	3
134	Addendum: Genome-wide association study of depression phenotypes in UK Biobank identifies variants in excitatory synaptic pathways. <i>Nature Communications</i> , 2018, 9, 3578.	5.8	16
135	T110. FIRST EPISODE PSYCHOTIC PATIENTS WITH A HISTORY OF FREQUENT CANNABIS USE EXPRESS MORE POSITIVE SYMPTOMS AT ILLNESS ONSET THAN THOSE WHO NEVER USED CANNABIS. <i>Schizophrenia Bulletin</i> , 2018, 44, S158-S159.	2.3	1
136	DNA methylation and inflammation marker profiles associated with a history of depression. <i>Human Molecular Genetics</i> , 2018, 27, 2840-2850.	1.4	46
137	Effect of cytochrome CYP2C19 metabolizing activity on antidepressant response and side effects: Meta-analysis of data from genome-wide association studies. <i>European Neuropsychopharmacology</i> , 2018, 28, 945-954.	0.3	64
138	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , 2018, 360, .	6.0	1,085
139	Use of schizophrenia and bipolar disorder polygenic risk scores to identify psychotic disorders. <i>British Journal of Psychiatry</i> , 2018, 213, 535-541.	1.7	37
140	Genes associated with anhedonia: a new analysis in a large clinical trial (GENDEP). <i>Translational Psychiatry</i> , 2018, 8, 150.	2.4	19
141	Genetic disposition to inflammation and response to antidepressants in major depressive disorder. <i>Journal of Psychiatric Research</i> , 2018, 105, 17-22.	1.5	18
142	Joint genome-wide association study of progressive supranuclear palsy identifies novel susceptibility loci and genetic correlation to neurodegenerative diseases. <i>Molecular Neurodegeneration</i> , 2018, 13, 41.	4.4	77
143	Genetic fine mapping of systemic lupus erythematosus MHC associations in Europeans and African Americans. <i>Human Molecular Genetics</i> , 2018, 27, 3813-3824.	1.4	43
144	No Association Between Antidepressant Efficacy and rs28365143 in Corticotropin-Releasing Hormone Binding Protein in a Large Meta-Analysis. <i>American Journal of Psychiatry</i> , 2018, 175, 575-576.	4.0	1

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145	Multiancestry genome-wide association study of 520,000 subjects identifies 32 loci associated with stroke and stroke subtypes. <i>Nature Genetics</i> , 2018, 50, 524-537.	9.4	1,124
146	Abstract 237: The genetic architecture of African esophageal cancer. , 2018, , .		0
147	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
148	Effect of waist-to-hip ratio on the association between type 2 diabetes and depression: an exploratory study using the polygenic scores approach in the UK Biobank. <i>Lancet, The</i> , 2017, 389, S53.	6.3	0
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