

Cathryn M Lewis

List of Publications by Year in Descending Order

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

420
papers

47,719
citations

83
h-index

214
g-index

507
ext. papers

57,055
ext. citations

8.1
avg. IF

7.85
L-index

#	Paper	IF	Citations
4 ²⁰	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	4.9	3
4 ¹⁹	Anxiety disorders and age-related changes in physiology.. <i>British Journal of Psychiatry</i> , 2022 , 1-10	5.4	1
4 ¹⁸	Latent subtypes of manic and/or irritable episode symptoms in two population-based cohorts.. <i>British Journal of Psychiatry</i> , 2022 , 1-10	5.4	1
4 ¹⁷	A tool for translating polygenic scores onto the absolute scale using summary statistics.. <i>European Journal of Human Genetics</i> , 2022 ,	5.3	3
4 ¹⁶	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. <i>Biological Psychiatry</i> , 2022 , 91, 102-117	7.9	11
4 ¹⁵	Age-related changes in physiology in individuals with bipolar disorder. <i>Journal of Affective Disorders</i> , 2022 , 296, 157-168	6.6	2
4 ¹⁴	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	2.8	1
4 ¹³	Genetic architecture of 11 major psychiatric disorders at biobehavioral, functional genomic and molecular genetic levels of analysis.. <i>Nature Genetics</i> , 2022 , 54, 548-559	36.3	4
4 ¹²	Epidemiology of Kaposi's sarcoma in sub-Saharan Africa.. <i>Cancer Epidemiology</i> , 2022 , 78, 102167	2.8	3
4 ¹¹	Cardiovascular health and risk of hospitalization with COVID-19: A Mendelian Randomization study. <i>JRSM Cardiovascular Disease</i> , 2021 , 10, 20480040211059374	1.1	0
4 ¹⁰	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	3.7	0
4 ⁰⁹	Vitamin D and the risk of treatment-resistant and atypical depression: A Mendelian randomization study. <i>Translational Psychiatry</i> , 2021 , 11, 561	8.6	0
4 ⁰⁸	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	14.4	2
4 ⁰⁷	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	5	0
4 ⁰⁶	The consequences of a year of the COVID-19 pandemic for the mental health of young adult twins in England and Wales 2021 ,		2
4 ⁰⁵	Exploring health in the UK Biobank: associations with sociodemographic characteristics, psychosocial factors, lifestyle and environmental exposures. <i>BMC Medicine</i> , 2021 , 19, 240	11.4	7
4 ⁰⁴	Genetic copy number variants, cognition and psychosis: a meta-analysis and a family study. <i>Molecular Psychiatry</i> , 2021 , 26, 5307-5319	15.1	5

403	Genetic and clinical characteristics of treatment-resistant depression using primary care records in two UK cohorts. <i>Molecular Psychiatry</i> , 2021 , 26, 3363-3373	15.1	20
402	Associations and limited shared genetic aetiology between bipolar disorder and cardiometabolic traits in the UK Biobank. <i>Psychological Medicine</i> , 2021 , 1-10	6.9	4
401	Genome-wide association study of suicidal behaviour severity in mood disorders. <i>World Journal of Biological Psychiatry</i> , 2021 , 22, 722-731	3.8	1
400	Genetic basis of lacunar stroke: a pooled analysis of individual patient data and genome-wide association studies. <i>Lancet Neurology</i> , 2021 , 20, 351-361	24.1	21
399	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. <i>Nature Genetics</i> , 2021 , 53, 817-829	36.3	83
398	Lack of Support for the Genes by Early Environment Interaction Hypothesis in the Pathogenesis of Schizophrenia. <i>Schizophrenia Bulletin</i> , 2021 ,	1.3	6
397	Evaluation of polygenic prediction methodology within a reference-standardized framework. <i>PLoS Genetics</i> , 2021 , 17, e1009021	6	22
396	The association between genetically determined ABO blood types and major depressive disorder. <i>Psychiatry Research</i> , 2021 , 299, 113837	9.9	
395	Transcriptome-wide association study of treatment-resistant depression and depression subtypes for drug repurposing. <i>Neuropsychopharmacology</i> , 2021 , 46, 1821-1829	8.7	5
394	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	11.9	30
393	Age-related changes in physiology in individuals with lifetime bipolar disorder. <i>BJPsych Open</i> , 2021 , 7, S276-S276	5	
392	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	5.1	1
391	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	5	
390	Investigating Pleiotropy Between Depression and Autoimmune Diseases Using the UK Biobank. <i>Biological Psychiatry Global Open Science</i> , 2021 , 1, 48-58		4
389	Using major depression polygenic risk scores to explore the depressive symptom continuum. <i>Psychological Medicine</i> , 2021 , 1-10	6.9	2
388	Drug repositioning for treatment-resistant depression: Hypotheses from a pharmacogenomic study. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2021 , 104, 110050	5.5	6
387	Examining the association between family status and depression in the UK Biobank. <i>Journal of Affective Disorders</i> , 2021 , 279, 585-598	6.6	2
386	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	6.6	2

385	Delineating the Genetic Component of Gene Expression in Major Depression. <i>Biological Psychiatry</i> , 2021 , 89, 627-636	7.9	16
384	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	3.5	
383	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	5.2	12
382	Imputed gene expression risk scores: a functionally informed component of polygenic risk. <i>Human Molecular Genetics</i> , 2021 , 30, 727-738	5.6	1
381	Multiple measures of depression to enhance validity of major depressive disorder in the UK Biobank. <i>BJPsych Open</i> , 2021 , 7, e44	5	10
380	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	5.6	5
379	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	8.6	1
378	Transcriptome-based polygenic score links depression-related corticolimbic gene expression changes to sex-specific brain morphology and depression risk. <i>Neuropsychopharmacology</i> , 2021 , 46, 2304-2311 ¹	8.7	1 ¹
377	Examining sex differences in neurodevelopmental and psychiatric genetic risk in anxiety and depression. <i>PLoS ONE</i> , 2021 , 16, e0248254	3.7	2
376	The Genetic Architecture of Depression in Individuals of East Asian Ancestry: A Genome-Wide Association Study. <i>JAMA Psychiatry</i> , 2021 , 78, 1258-1269	14.5	7
375	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	7.2	8
374	Investigating rare pathogenic/likely pathogenic exonic variation in bipolar disorder. <i>Molecular Psychiatry</i> , 2021 , 26, 5239-5250	15.1	3
373	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> , 2020 , 1-9	6.9	4
372	Genome-wide association study of MRI markers of cerebral small vessel disease in 42,310 participants. <i>Nature Communications</i> , 2020 , 11, 2175	17.4	21
371	Polygenic risk scores: from research tools to clinical instruments. <i>Genome Medicine</i> , 2020 , 12, 44	14.4	190
370	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	11.6	11
369	Genetic stratification of depression in UK Biobank. <i>Translational Psychiatry</i> , 2020 , 10, 163	8.6	8
368	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> , 2020 , 1-9	6.9	13

367	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	2.8	4
366	Minimal phenotyping yields genome-wide association signals of low specificity for major depression. <i>Nature Genetics</i> , 2020 , 52, 437-447	36.3	80
365	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	4.5	16
364	A polygenic predictor of treatment-resistant depression using whole exome sequencing and genome-wide genotyping. <i>Translational Psychiatry</i> , 2020 , 10, 50	8.6	16
363	The behavioral, cellular and immune mediators of HIV-1 acquisition: New insights from population genetics. <i>Scientific Reports</i> , 2020 , 10, 3304	4.9	4
362	Genome-wide gene-environment analyses of major depressive disorder and reported lifetime traumatic experiences in UK Biobank. <i>Molecular Psychiatry</i> , 2020 , 25, 1430-1446	15.1	47
361	Cannabis use, depression and self-harm: phenotypic and genetic relationships. <i>Addiction</i> , 2020 , 115, 482-492	4.2	16
360	S126. THE RELATION OF THE PSYCHOSIS CONTINUUM WITH SCHIZOPHRENIA POLYGENIC RISK SCORE AND CANNABIS USE. <i>Schizophrenia Bulletin</i> , 2020 , 46, S83-S83	1.3	78
359	Depression with atypical neurovegetative symptoms shares genetic predisposition with immuno-metabolic traits and alcohol consumption. <i>Psychological Medicine</i> , 2020 , 1-11	6.9	17
358	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	3.5	8
357	Genetics and major depressive disorder: clinical implications for disease risk, prognosis and treatment. <i>International Clinical Psychopharmacology</i> , 2020 , 35, 233-242	2.2	11
356	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	6.7	7
355	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	1.1	5
354	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	3.5	24
353	The Maudsley environmental risk score for psychosis. <i>Psychological Medicine</i> , 2020 , 50, 2213-2220	6.9	14
352	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	6.9	1
351	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	6.9	10
350	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. <i>Biological Psychiatry</i> , 2020 , 87, 419-430	7.9	9

349	Genetic Associations Between Childhood Psychopathology and Adult Depression and Associated Traits in 42 998 Individuals: A Meta-analysis. <i>JAMA Psychiatry</i> , 2020 , 77, 715-728	14.5	15
348	A PGC-MDD COLLABORATION ON THE GENETICS OF TREATMENT OUTCOMES IN DEPRESSION. <i>European Neuropsychopharmacology</i> , 2019 , 29, S1044	1.2	
347	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	3.5	16
346	ukbtools: An R package to manage and query UK Biobank data. <i>PLoS ONE</i> , 2019 , 14, e0214311	3.7	15
345	Prevalence of Advanced Liver Fibrosis in Patients With Severe Psoriasis. <i>JAMA Dermatology</i> , 2019 , 155, 1028-1032	5.1	8
344	GWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. <i>American Journal of Psychiatry</i> , 2019 , 176, 651-660	11.9	103
343	Genome-wide survey of copy number variants finds MAPT duplications in progressive supranuclear palsy. <i>Movement Disorders</i> , 2019 , 34, 1049-1059	7	15
342	Polygenic risk for circulating reproductive hormone levels and their influence on hippocampal volume and depression susceptibility. <i>Psychoneuroendocrinology</i> , 2019 , 106, 284-292	5	13
341	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019 , 51, 793-803	36.3	662
340	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	23.3	322
339	Genome-wide Burden of Rare Short Deletions Is Enriched in Major Depressive Disorder in Four Cohorts. <i>Biological Psychiatry</i> , 2019 , 85, 1065-1073	7.9	14
338	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	1.3	78
337	Evidence of causal effect of major depression on alcohol dependence: findings from the psychiatric genomics consortium. <i>Psychological Medicine</i> , 2019 , 49, 1218-1226	6.9	33
336	Uncovering the Genetic Architecture of Major Depression. <i>Neuron</i> , 2019 , 102, 91-103	13.9	66
335	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	8.6	30
334	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	8.5	8
333	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	3.7	8
332	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	25.5	639

331	Association of genetic variants in CHEK2 with oesophageal squamous cell carcinoma in the South African Black population. <i>Carcinogenesis</i> , 2019 , 40, 513-520	4.6	8
330	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , 2019 , 179, 1469-1482.e11	56.2	402
329	Association of Whole-Genome and NETRIN1 Signaling Pathway-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	3.4	12
328	The polygenic nature of telomere length and the anti-ageing properties of lithium. <i>Neuropsychopharmacology</i> , 2019 , 44, 757-765	8.7	25
327	Using genetics to understand the causal influence of higher BMI on depression. <i>International Journal of Epidemiology</i> , 2019 , 48, 834-848	7.8	81
326	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	5.4	24
325	Telomere length as a predictor of emotional processing in the brain. <i>Human Brain Mapping</i> , 2019 , 40, 1750-1759	5.9	13
324	Transdiagnostic dimensions of psychopathology at first episode psychosis: findings from the multinational EU-GEI study. <i>Psychological Medicine</i> , 2019 , 49, 1378-1391	6.9	42
323	Genome-wide association study of depression phenotypes in UK Biobank identifies variants in excitatory synaptic pathways. <i>Nature Communications</i> , 2018 , 9, 1470	17.4	226
322	Antidepressant drug-specific prediction of depression treatment outcomes from genetic and clinical variables. <i>Scientific Reports</i> , 2018 , 8, 5530	4.9	31
321	Association analysis in over 329,000 individuals identifies 116 independent variants influencing neuroticism. <i>Nature Genetics</i> , 2018 , 50, 6-11	36.3	202
320	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. <i>Nature Genetics</i> , 2018 , 50, 668-681	36.3	1301
319	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	7.9	48
318	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	3.5	39
317	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	16.6	26
316	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , 2018 , 360,	33.3	666
315	Use of schizophrenia and bipolar disorder polygenic risk scores to identify psychotic disorders. <i>British Journal of Psychiatry</i> , 2018 , 213, 535-541	5.4	21
314	Genes associated with anhedonia: a new analysis in a large clinical trial (GENDEP). <i>Translational Psychiatry</i> , 2018 , 8, 150	8.6	11

313	Genetic disposition to inflammation and response to antidepressants in major depressive disorder. <i>Journal of Psychiatric Research</i> , 2018 , 105, 17-22	5.2	12
312	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	19	41
311	Genetic fine mapping of systemic lupus erythematosus MHC associations in Europeans and African Americans. <i>Human Molecular Genetics</i> , 2018 , 27, 3813-3824	5.6	23
310	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	11.9	1
309	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	36.3	536
308	Genome-wide gene-environment interaction in depression: A systematic evaluation of candidate genes: The childhood trauma working-group of PGC-MDD. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018 , 177, 40-49	3.5	43
307	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	3.5	26
306	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	1.1	9
305	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	1.3	1
304	Genetic Risk for Psychiatric Disorders and Telomere Length. <i>Frontiers in Genetics</i> , 2018 , 9, 468	4.5	13
303	Expression quantitative trait loci in the developing human brain and their enrichment in neuropsychiatric disorders. <i>Genome Biology</i> , 2018 , 19, 194	18.3	69
302	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	3.7	25
301	Exome array analysis of adverse reactions to fluoropyrimidine-based therapy for gastrointestinal cancer. <i>PLoS ONE</i> , 2018 , 13, e0188911	3.7	1
300	Addendum: Genome-wide association study of depression phenotypes in UK Biobank identifies variants in excitatory synaptic pathways. <i>Nature Communications</i> , 2018 , 9, 3578	17.4	3
299	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	1.3	1
298	DNA methylation and inflammation marker profiles associated with a history of depression. <i>Human Molecular Genetics</i> , 2018 , 27, 2840-2850	5.6	23
297	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	1.2	43
296	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	7.9	129

295	Pharmacogenetics of antidepressant response: A polygenic approach. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2017 , 75, 128-134	5.5	54
294	Genetic and environmental risk factors for rheumatoid arthritis in a UK African ancestry population: the GENRA case-control study. <i>Rheumatology</i> , 2017 , 56, 1282-1292	3.9	15
293	Validating a two-dimensional bipolar spectrum model integrating DSM-5's mixed features specifier for Major Depressive Disorder. <i>Comprehensive Psychiatry</i> , 2017 , 77, 89-99	7.3	7
292	MicroRNAs 146a/b-5 and 425-3p and 24-3p are markers of antidepressant response and regulate MAPK/Wnt-system genes. <i>Nature Communications</i> , 2017 , 8, 15497	17.4	93
291	Genetics of Depression: Progress at Last. <i>Current Psychiatry Reports</i> , 2017 , 19, 43	9.1	65
290	Reproductive fitness and genetic risk of psychiatric disorders in the general population. <i>Nature Communications</i> , 2017 , 8, 15833	17.4	19
289	Cost-effectiveness of pharmacogenetic-guided treatment: are we there yet?. <i>Pharmacogenomics Journal</i> , 2017 , 17, 395-402	3.5	118
288	The Relationship Between Mental Health, Disease Severity, and Genetic Risk for Depression in Early Rheumatoid Arthritis. <i>Psychosomatic Medicine</i> , 2017 , 79, 638-645	3.7	21
287	Genetic effects influencing risk for major depressive disorder in China and Europe. <i>Translational Psychiatry</i> , 2017 , 7, e1074	8.6	48
286	Genetics of stroke in a UK African ancestry case-control study: South London Ethnicity and Stroke Study. <i>Neurology: Genetics</i> , 2017 , 3, e142	3.8	15
285	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. <i>Biological Psychiatry</i> , 2017 , 82, 322-329	7.9	68
284	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	7.9	17
283	Genetic variation at 16q24.2 is associated with small vessel stroke. <i>Annals of Neurology</i> , 2017 , 81, 383-394	9.4	51
282	Genetic Association of Major Depression With Atypical Features and Obesity-Related Immunometabolic Dysregulations. <i>JAMA Psychiatry</i> , 2017 , 74, 1214-1225	14.5	109
281	A bidirectional relationship between depression and the autoimmune disorders - New perspectives from the National Child Development Study. <i>PLoS ONE</i> , 2017 , 12, e0173015	3.7	46
280	Prospects for using risk scores in polygenic medicine. <i>Genome Medicine</i> , 2017 , 9, 96	14.4	103
279	Interaction between the gene, body mass index and depression: meta-analysis of 13701 individuals. <i>British Journal of Psychiatry</i> , 2017 , 211, 70-76	5.4	33
278	Advancing psychiatric genetics through dissecting heterogeneity. <i>Human Molecular Genetics</i> , 2017 , 26, R160-R165	5.6	7

277	An additional k-means clustering step improves the biological features of WGCNA gene co-expression networks. <i>BMC Systems Biology</i> , 2017 , 11, 47	3.5	134
276	Pharmacogenetic testing through the direct-to-consumer genetic testing company 23andMe. <i>BMC Medical Genomics</i> , 2017 , 10, 47	3.7	18
275	An Examination of Polygenic Score Risk Prediction in Individuals With First-Episode Psychosis. <i>Biological Psychiatry</i> , 2017 , 81, 470-477	7.9	126
274	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	7.9	26
273	Genetic discovery in multi-ethnic populations. <i>European Journal of Human Genetics</i> , 2016 , 24, 1097-8	5.3	4
272	Rare genetic variation in UNC13A may modify survival in amyotrophic lateral sclerosis. <i>Amyotrophic Lateral Sclerosis and Frontotemporal Degeneration</i> , 2016 , 17, 593-599	3.6	16
271	Genome-wide association analyses identify new risk variants and the genetic architecture of amyotrophic lateral sclerosis. <i>Nature Genetics</i> , 2016 , 48, 1043-8	36.3	328
270	Association of a Locus in the CAMTA1 Gene With Survival in Patients With Sporadic Amyotrophic Lateral Sclerosis. <i>JAMA Neurology</i> , 2016 , 73, 812-20	17.2	40
269	Meta-analysis of the Association Between the Level of Cannabis Use and Risk of Psychosis. <i>Schizophrenia Bulletin</i> , 2016 , 42, 1262-9	1.3	414
268	Genome-wide meta-analysis of cerebral white matter hyperintensities in patients with stroke. <i>Neurology</i> , 2016 , 86, 146-53	6.5	67
267	Impact of intensive treatment and remission on health-related quality of life in early and established rheumatoid arthritis. <i>RMD Open</i> , 2016 , 2, e000270	5.9	12
266	Polygenic interactions with environmental adversity in the aetiology of major depressive disorder. <i>Psychological Medicine</i> , 2016 , 46, 759-70	6.9	125
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33	Genetic variation in the Major Histocompatibility Complex and association with depression		1
32	Identifying Potential Causal Risk Factors for Self-Harm: A Polygenic Risk Scoring and Mendelian Randomisation Approach		1
31	What's in a diagnosis? A genetic decomposition of major depression		1
30	Pharmacogenetic testing through the direct-to-consumer genetic testing company 23andMe		1
29	Genetic stratification of depression in UK Biobank suggests a subgroup linked to age of natural menopause		6
28	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depressive disorder		21
27	Genome-wide association study of depression phenotypes in UK Biobank (n = 322,580) identifies the enrichment of variants in excitatory synaptic pathways		8
26	116 independent genetic variants influence the neuroticism personality trait in over 329,000 UK Biobank individuals		4

25	Genome-wide association study identifies 30 Loci Associated with Bipolar Disorder	28
24	Atypical depression shares genetic predisposition with immuno-metabolic traits: a population-based study	2
23	Using Major Depression Polygenic Risk Scores to Explore the Depressive Symptom Continuum	2
22	Phenotypic Associations and Shared Genetic Etiology between Bipolar Disorder and Cardiometabolic Traits	1
21	Exploring health in the UK Biobank: associations with sociodemographic characteristics, psychosocial factors, lifestyle and environmental exposures	3
20	Evaluation of Polygenic Prediction Methodology within a Reference-Standardized Framework	1
19	Genetic and clinical characteristics of treatment-resistant depression using primary care records in two UK cohorts	3
18	Genome-wide association study of over 40,000 bipolar disorder cases provides new insights into the underlying biology	11
17	Multiple measures of depression to enhance validity of Major Depressive Disorder in the UK Biobank	4
16	Genetic Architecture of 11 Major Psychiatric Disorders at Biobehavioral, Functional Genomic, and Molecular Genetic Levels of Analysis	4
15	Dissecting the shared genetic architecture of suicide attempt, psychiatric disorders and known risk factors	2
14	Genome-wide gene-environment analyses of depression and reported lifetime traumatic experiences in UK Biobank	9
13	Transforming summary statistics from logistic regression to the liability scale: application to genetic and environmental risk scores	1
12	Evidence of causal effect of major depression on alcohol dependence: Findings from the Psychiatric Genomics Consortium	4
11	Genome-wide meta-analysis of depression identifies 102 independent variants and highlights the importance of the prefrontal brain regions	8
10	Minimal phenotyping yields GWAS hits of reduced specificity for major depression	12
9	Genome wide meta-analysis identifies genomic relationships, novel loci, and pleiotropic mechanisms across eight psychiatric disorders	10
8	The independent and combined influence of schizophrenia polygenic risk score and heavy cannabis use on risk for psychotic disorder: A case-control analysis from the EUGEI study.	2

7	Impute.me: an open source, non-profit tool for using data from DTC genetic testing to calculate and interpret polygenic risk scores	2
6	Predictors of intestinal inflammation in asymptomatic first-degree relatives of patients with Crohn's disease	1
5	Age-related changes in physiology in individuals with bipolar disorder	2
4	Cross-classification between self-rated health and health status: longitudinal analyses of all-cause mortality and leading causes of death in the UK	1
3	Lifetime depression and age-related changes in body composition, cardiovascular function, grip strength and lung function: sex-specific analyses in the UK Biobank	1
2	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> ,1-9	6.9
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