

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	Genome-wide association study of 14,000 cases of seven common diseases and 3,000 shared controls. <i>Nature</i> , 2007, 447, 661-678.	13.7	8,895
2	A strong candidate for the breast and ovarian cancer susceptibility gene BRCA1. <i>Science</i> , 1994, 266, 66-71.	6.0	5,747
3	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
4	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. <i>Nature Genetics</i> , 2013, 45, 984-994.	9.4	2,067
5	Localization of a breast cancer susceptibility gene, BRCA2, to chromosome 13q12-13. <i>Science</i> , 1994, 265, 2088-2090.	6.0	1,725
6	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
7	Genome-wide association analysis identifies 13 new risk loci for schizophrenia. <i>Nature Genetics</i> , 2013, 45, 1150-1159.	9.4	1,395
8	Association scan of 14,500 nonsynonymous SNPs in four diseases identifies autoimmunity variants. <i>Nature Genetics</i> , 2007, 39, 1329-1337.	9.4	1,298
9	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019, 51, 793-803.	9.4	1,191
10	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
11	PRSice: Polygenic Risk Score software. <i>Bioinformatics</i> , 2015, 31, 1466-1468.	1.8	1,109
12	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , 2018, 360, .	6.0	1,085
13	Genome Scan Meta-Analysis of Schizophrenia and Bipolar Disorder, Part II: Schizophrenia. <i>American Journal of Human Genetics</i> , 2003, 73, 34-48.	2.6	1,072
14	Association between insertion mutation in NOD2 gene and Crohn's disease in German and British populations. <i>Lancet</i> , The, 2001, 357, 1925-1928.	6.3	1,071
15	Sequence variants in the autophagy gene IRGM and multiple other replicating loci contribute to Crohn's disease susceptibility. <i>Nature Genetics</i> , 2007, 39, 830-832.	9.4	1,063
16	A mega-analysis of genome-wide association studies for major depressive disorder. <i>Molecular Psychiatry</i> , 2013, 18, 497-511.	4.1	1,002
17	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , 2019, 179, 1469-1482.e11.	13.5	935
18	Suberoylanilide hydroxamic acid, a histone deacetylase inhibitor, ameliorates motor deficits in a mouse model of Huntington's disease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 2041-2046.	3.3	805

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19	Psychiatric genome-wide association study analyses implicate neuronal, immune and histone pathways. <i>Nature Neuroscience</i> , 2015, 18, 199-209.	7.1	701
20	The contribution of NOD2 gene mutations to the risk and site of disease in inflammatory bowel disease. <i>Gastroenterology</i> , 2002, 122, 867-874.	0.6	670
21	Polygenic risk scores: from research tools to clinical instruments. <i>Genome Medicine</i> , 2020, 12, 44.	3.6	646
22	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
23	Meta-analysis of the Association Between the Level of Cannabis Use and Risk of Psychosis. <i>Schizophrenia Bulletin</i> , 2016, 42, 1262-1269.	2.3	615
24	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
25	Assignment of a locus for familial melanoma, MLM, to chromosome 9p13-p22. <i>Science</i> , 1992, 258, 1148-1152.	6.0	506
26	Genome-wide association analyses identify new risk variants and the genetic architecture of amyotrophic lateral sclerosis. <i>Nature Genetics</i> , 2016, 48, 1043-1048.	9.4	494
27	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
28	Genome-wide association study of major depressive disorder: new results, meta-analysis, and lessons learned. <i>Molecular Psychiatry</i> , 2012, 17, 36-48.	4.1	405
29	Genetic determinants of ulcerative colitis include the ECM1 locus and five loci implicated in Crohn's disease. <i>Nature Genetics</i> , 2008, 40, 710-712.	9.4	403
30	Genome Scan Meta-Analysis of Schizophrenia and Bipolar Disorder, Part III: Bipolar Disorder. <i>American Journal of Human Genetics</i> , 2003, 73, 49-62.	2.6	400
31	Genetic association studies: Design, analysis and interpretation. <i>Briefings in Bioinformatics</i> , 2002, 3, 146-153.	3.2	392
32	EULAR recommendations for terminology and research in individuals at risk of rheumatoid arthritis: report from the Study Group for Risk Factors for Rheumatoid Arthritis. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 638-641.	0.5	354
33	Meta-Analysis of the Association of Urbanicity With Schizophrenia. <i>Schizophrenia Bulletin</i> , 2012, 38, 1118-1123.	2.3	349
34	Association analysis in over 329,000 individuals identifies 116 independent variants influencing neuroticism. <i>Nature Genetics</i> , 2018, 50, 6-11.	9.4	327
35	Genome-Wide Pharmacogenetics of Antidepressant Response in the GENDEP Project. <i>American Journal of Psychiatry</i> , 2010, 167, 555-564.	4.0	314
36	Adverse drug reactions to azathioprine therapy are associated with polymorphism in the gene encoding inosine triphosphate pyrophosphatase (ITPase). <i>Pharmacogenetics and Genomics</i> , 2004, 14, 181-187.	5.7	305

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37	Environmental enrichment slows disease progression in R6/2 Huntington's disease mice. <i>Annals of Neurology</i> , 2002, 51, 235-242.	2.8	303
38	Fecundity of Patients With Schizophrenia, Autism, Bipolar Disorder, Depression, Anorexia Nervosa, or Substance Abuse vs Their Unaffected Siblings. <i>JAMA Psychiatry</i> , 2013, 70, 22.	6.0	284
39	A genome-wide association study of anorexia nervosa. <i>Molecular Psychiatry</i> , 2014, 19, 1085-1094.	4.1	282
40	A Nonsynonymous SNP in ATG16L1 Predisposes to Ileal Crohn's Disease and Is Independent of CARD15 and IBD5. <i>Gastroenterology</i> , 2007, 132, 1665-1671.	0.6	268
41	An additional k-means clustering step improves the biological features of WGCNA gene co-expression networks. <i>BMC Systems Biology</i> , 2017, 11, 47.	3.0	253
42	Meta-analysis of 32 genome-wide linkage studies of schizophrenia. <i>Molecular Psychiatry</i> , 2009, 14, 774-785.	4.1	235
43	Meta-analysis of genome scans of age-related macular degeneration. <i>Human Molecular Genetics</i> , 2005, 14, 2257-2264.	1.4	224
44	Genome-Wide Association Study of Major Recurrent Depression in the U.K. Population. <i>American Journal of Psychiatry</i> , 2010, 167, 949-957.	4.0	221
45	Inflammatory bowel disease susceptibility loci defined by genome scan meta-analysis of 1952 affected relative pairs. <i>Human Molecular Genetics</i> , 2004, 13, 763-770.	1.4	219
46	Common Genetic Variation and Antidepressant Efficacy in Major Depressive Disorder: A Meta-Analysis of Three Genome-Wide Pharmacogenetic Studies. <i>American Journal of Psychiatry</i> , 2013, 170, 207-217.	4.0	216
47	Minimal phenotyping yields genome-wide association signals of low specificity for major depression. <i>Nature Genetics</i> , 2020, 52, 437-447.	9.4	207
48	Meta-analysis of genome searches. <i>Annals of Human Genetics</i> , 1999, 63, 263-272.	0.3	205
49	Chromosome 9p21 in sporadic amyotrophic lateral sclerosis in the UK and seven other countries: a genome-wide association study. <i>Lancet Neurology</i> , The, 2010, 9, 986-994.	4.9	205
50	Contribution of Common Genetic Variants to Antidepressant Response. <i>Biological Psychiatry</i> , 2013, 73, 679-682.	0.7	199
51	Investigation of Crohn's Disease Risk Loci in Ulcerative Colitis Further Defines Their Molecular Relationship. <i>Gastroenterology</i> , 2009, 136, 523-529.e3.	0.6	198
52	The Neuronal Transporter Gene SLC6A15 Confers Risk to Major Depression. <i>Neuron</i> , 2011, 70, 252-265.	3.8	189
53	Genetic predictors of response to antidepressants in the GENDEP project. <i>Pharmacogenomics Journal</i> , 2009, 9, 225-233.	0.9	188
54	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

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55	Polygenic interactions with environmental adversity in the aetiology of major depressive disorder. <i>Psychological Medicine</i> , 2016, 46, 759-770.	2.7	176
56	An Examination of Polygenic Score Risk Prediction in Individuals With First-Episode Psychosis. <i>Biological Psychiatry</i> , 2017, 81, 470-477.	0.7	176
57	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
58	Genetic Association of Major Depression With Atypical Features and Obesity-Related Immunometabolic Dysregulations. <i>JAMA Psychiatry</i> , 2017, 74, 1214.	6.0	174
59	Cost-effectiveness of pharmacogenetic-guided treatment: are we there yet?. <i>Pharmacogenomics Journal</i> , 2017, 17, 395-402.	0.9	173
60	IL23R Variation Determines Susceptibility But Not Disease Phenotype in Inflammatory Bowel Disease. <i>Gastroenterology</i> , 2007, 132, 1657-1664.	0.6	170
61	Meta-analysis of genome-wide linkage scans of attention deficit hyperactivity disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008, 147B, 1392-1398.	1.1	160
62	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
63	Prospects for using risk scores in polygenic medicine. <i>Genome Medicine</i> , 2017, 9, 96.	3.6	153
64	Prospective evaluation of the pharmacogenetics of azathioprine in the treatment of inflammatory bowel disease. <i>Alimentary Pharmacology and Therapeutics</i> , 2008, 28, 973-983.	1.9	152
65	Combining clinical variables to optimize prediction of antidepressant treatment outcomes. <i>Journal of Psychiatric Research</i> , 2016, 78, 94-102.	1.5	149
66	Genome Scan Meta-Analysis of Schizophrenia and Bipolar Disorder, Part I: Methods and Power Analysis. <i>American Journal of Human Genetics</i> , 2003, 73, 17-33.	2.6	147
67	Meta-Analysis of Genome-Wide Scans Provides Evidence for Sex- and Site-Specific Regulation of Bone Mass. <i>Journal of Bone and Mineral Research</i> , 2006, 22, 173-183.	3.1	144
68	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.	5.8	144
69	Introduction to Genetic Association Studies. <i>Cold Spring Harbor Protocols</i> , 2012, 2012, pdb.top068163.	0.2	140
70	Pharmacogenetic variants in the DPYD, TYMS, CDA and MTHFR genes are clinically significant predictors of fluoropyrimidine toxicity. <i>British Journal of Cancer</i> , 2013, 108, 2505-2515.	2.9	139
71	Meta-analysis of Genome-wide Linkage Studies in BMI and Obesity. <i>Obesity</i> , 2007, 15, 2263-2275.	1.5	138
72	Therapygenetics: the 5HTTLPR and response to psychological therapy. <i>Molecular Psychiatry</i> , 2012, 17, 236-237.	4.1	135

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73	Penetrance for copy number variants associated with schizophrenia. <i>Human Molecular Genetics</i> , 2010, 19, 3477-3481.	1.4	132
74	Standardization and statistical approaches to therapeutic trials in the R6/2 mouse. <i>Brain Research Bulletin</i> , 2003, 61, 469-479.	1.4	129
75	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
76	The ?174G allele of the interleukin-6 gene confers susceptibility to systemic arthritis in children: A multicenter study using simplex and multiplex juvenile idiopathic arthritis families. <i>Arthritis and Rheumatism</i> , 2003, 48, 3202-3206.	6.7	123
77	A genome-wide association meta-analysis identifies a novel locus at 17q11.2 associated with sporadic amyotrophic lateral sclerosis. <i>Human Molecular Genetics</i> , 2014, 23, 2220-2231.	1.4	123
78	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
79	Response to radiation therapy and prognosis in breast cancer patients with BRCA1 and BRCA2 mutations. <i>Radiotherapy and Oncology</i> , 1998, 47, 129-136.	0.3	114
80	Comorbid medical illness in bipolar disorder. <i>British Journal of Psychiatry</i> , 2014, 205, 465-472.	1.7	113
81	Uncovering the Genetic Architecture of Major Depression. <i>Neuron</i> , 2019, 102, 91-103.	3.8	113
82	Genetic Evidence for Interaction of the 5q31 Cytokine Locus and the CARD15 Gene in Crohn Disease. <i>American Journal of Human Genetics</i> , 2003, 72, 1018-1022.	2.6	111
83	Genetic Predictors of Response to Serotonergic and Noradrenergic Antidepressants in Major Depressive Disorder: A Genome-Wide Analysis of Individual-Level Data and a Meta-Analysis. <i>PLoS Medicine</i> , 2012, 9, e1001326.	3.9	110
84	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
85	A multiobserver, population-based analysis of histologic dysplasia in melanocytic nevi. <i>Journal of the American Academy of Dermatology</i> , 1994, 30, 707-714.	0.6	108
86	Correlation of SMNt and SMNc gene copy number with age of onset and survival in spinal muscular atrophy. <i>European Journal of Human Genetics</i> , 1998, 6, 467-474.	1.4	108
87	Genetics of inflammatory bowel disease: progress and prospects. <i>Human Molecular Genetics</i> , 2004, 13, 161R-168.	1.4	106
88	Genetic Predictors of Increase in Suicidal Ideation During Antidepressant Treatment in the GENDEP Project. <i>Neuropsychopharmacology</i> , 2009, 34, 2517-2528.	2.8	105
89	Polymorphisms in Folate, Pyrimidine, and Purine Metabolism Are Associated with Efficacy and Toxicity of Methotrexate in Psoriasis. <i>Journal of Investigative Dermatology</i> , 2007, 127, 1860-1867.	0.3	104
90	Genetics of Depression: Progress at Last. <i>Current Psychiatry Reports</i> , 2017, 19, 43.	2.1	101

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91	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
92	Incidence of multiple primary cancers in a cohort of women diagnosed with breast cancer in southeast England. <i>British Journal of Cancer</i> , 2001, 84, 435-440.	2.9	99
93	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
94	Evaluation of polygenic prediction methodology within a reference-standardized framework. <i>PLoS Genetics</i> , 2021, 17, e1009021.	1.5	99
95	Genomewide Association Scan of Suicidal Thoughts and Behaviour in Major Depression. <i>PLoS ONE</i> , 2011, 6, e20690.	1.1	98
96	The risk of subsequent primary cancers after colorectal cancer in southeast England. <i>Gut</i> , 2002, 50, 647-652.	6.1	97
97	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
98	Independent and population-specific association of risk variants at the IRGM locus with Crohn's disease. <i>Human Molecular Genetics</i> , 2010, 19, 1828-1839.	1.4	93
99	Modelling the Effects of Penetrance and Family Size on Rates of Sporadic and Familial Disease. <i>Human Heredity</i> , 2011, 71, 281-288.	0.4	93
100	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
101	Genome-wide association study of increasing suicidal ideation during antidepressant treatment in the GENDEP project. <i>Pharmacogenomics Journal</i> , 2012, 12, 68-77.	0.9	92
102	Genome-wide meta-analysis of cerebral white matter hyperintensities in patients with stroke. <i>Neurology</i> , 2016, 86, 146-153.	1.5	91
103	The protective effect of alcohol on developing rheumatoid arthritis: a systematic review and meta-analysis. <i>Rheumatology</i> , 2013, 52, 856-867.	0.9	89
104	A Genome-Wide Test of the Differential Susceptibility Hypothesis Reveals a Genetic Predictor of Differential Response to Psychological Treatments for Child Anxiety Disorders. <i>Psychotherapy and Psychosomatics</i> , 2016, 85, 146-158.	4.0	89
105	Molecular Characterization of CTNS Deletions in Nephropathic Cystinosis: Development of a PCR-Based Detection Assay. <i>American Journal of Human Genetics</i> , 1999, 65, 353-359.	2.6	88
106	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. <i>JAMA Psychiatry</i> , 2021, 78, 1258.	6.0	88
107	Associations of allelic variants of the multidrug resistance gene (ABCB1 or MDR1) and Inflammatory Bowel Disease and their effects on disease behavior: A case-control and meta-analysis study. <i>Inflammatory Bowel Diseases</i> , 2006, 12, 263-271.	0.9	87
108	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

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109	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
110	Relationship Between p53 Codon 72 Polymorphism and Susceptibility to Sunburn and Skin Cancer. <i>Journal of Investigative Dermatology</i> , 2002, 119, 84-90.	0.3	83
111	Controlling misdiagnosis errors in preimplantation genetic diagnosis: a comprehensive model encompassing extrinsic and intrinsic sources of error. <i>Human Reproduction</i> , 2001, 16, 43-50.	0.4	81
112	Novel pharmacogenetic markers for treatment outcome in azathioprine-treated inflammatory bowel disease. <i>Alimentary Pharmacology and Therapeutics</i> , 2009, 30, 375-384.	1.9	78
113	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
114	Two Families with Familial Amyotrophic Lateral Sclerosis Are Linked to a Novel Locus on Chromosome 16q. <i>American Journal of Human Genetics</i> , 2003, 73, 390-396.	2.6	76
115	Meta-Analysis of 4 Coronary Heart Disease Genome-Wide Linkage Studies Confirms a Susceptibility Locus on Chromosome 3q. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2003, 23, 1863-1868.	1.1	76
116	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
117	A Novel MMP12 Locus Is Associated with Large Artery Atherosclerotic Stroke Using a Genome-Wide Age-at-Onset Informed Approach. <i>PLoS Genetics</i> , 2014, 10, e1004469.	1.5	75
118	EPIBLASTER-fast exhaustive two-locus epistasis detection strategy using graphical processing units. <i>European Journal of Human Genetics</i> , 2011, 19, 465-471.	1.4	74
119	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
120	Novel IL10 gene family associations with systemic juvenile idiopathic arthritis. <i>Arthritis Research and Therapy</i> , 2006, 8, R148.	1.6	73
121	Genetic variation at 16q24.2 is associated with small vessel stroke. <i>Annals of Neurology</i> , 2017, 81, 383-394.	2.8	73
122	Meta-analysis of four rheumatoid arthritis genome-wide linkage studies: Confirmation of a susceptibility locus on chromosome 16. <i>Arthritis and Rheumatism</i> , 2003, 48, 1200-1206.	6.7	72
123	Depressive disorder moderates the effect of the FTO gene on body mass index. <i>Molecular Psychiatry</i> , 2012, 17, 604-611.	4.1	72
124	ACPA-positive and ACPA-negative rheumatoid arthritis differ in their requirements for combination DMARDs and corticosteroids: secondary analysis of a randomized controlled trial. <i>Arthritis Research and Therapy</i> , 2014, 16, R13.	1.6	72
125	The correlation between reading and mathematics ability at age twelve has a substantial genetic component. <i>Nature Communications</i> , 2014, 5, 4204.	5.8	72
126	Pharmacogenetics of antidepressant response: A polygenic approach. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2017, 75, 128-134.	2.5	71

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127	Association of DLG5 R30Q variant with inflammatory bowel disease. <i>European Journal of Human Genetics</i> , 2005, 13, 835-839.	1.4	70
128	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
129	Polygenic scores in biomedical research. <i>Nature Reviews Genetics</i> , 2022, 23, 524-532.	7.7	69
130	Predicting outcomes following cognitive behaviour therapy in child anxiety disorders: the influence of genetic, demographic and clinical information. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2013, 54, 1086-1094.	3.1	68
131	Tumor necrosis factor and its targets in the inflammatory cytokine pathway are identified as putative transcriptomic biomarkers for escitalopram response. <i>European Neuropsychopharmacology</i> , 2013, 23, 1105-1114.	0.3	68
132	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
133	A candidate gene analysis of three related photosensitivity disorders: cutaneous lupus erythematosus, polymorphic light eruption and actinic prurigo. <i>British Journal of Dermatology</i> , 2001, 145, 229-236.	1.4	65
134	Predicting treatment response in psoriasis using serum levels of adalimumab and etanercept: a single-centre, cohort study. <i>British Journal of Dermatology</i> , 2013, 169, 306-313.	1.4	65
135	Genetic effects influencing risk for major depressive disorder in China and Europe. <i>Translational Psychiatry</i> , 2017, 7, e1074-e1074.	2.4	64
136	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
137	Delineating the Genetic Component of Gene Expression in Major Depression. <i>Biological Psychiatry</i> , 2021, 89, 627-636.	0.7	63
138	Relationship between obesity and the risk of clinically significant depression: Mendelian randomisation study. <i>British Journal of Psychiatry</i> , 2014, 205, 24-28.	1.7	62
139	From SNPs to Genes: Disease Association at the Gene Level. <i>PLoS ONE</i> , 2011, 6, e20133.	1.1	61
140	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
141	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. <i>Biological Psychiatry</i> , 2022, 91, 102-117.	0.7	61
142	Genetic modelling of dizygotic twinning in pedigrees of spontaneous dizygotic twins. , 1996, 61, 258-263.		60
143	Genetic risk score analysis indicates migraine with and without comorbid depression are genetically different disorders. <i>Human Genetics</i> , 2014, 133, 173-186.	1.8	60
144	The association between lower educational attainment and depression owing to shared genetic effects? Results in ~25% subjects. <i>Molecular Psychiatry</i> , 2015, 20, 735-743.	4.1	59

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145	Pooled Sequencing of 531 Genes in Inflammatory Bowel Disease Identifies an Associated Rare Variant in BTNL2 and Implicates Other Immune Related Genes. <i>PLoS Genetics</i> , 2015, 11, e1004955.	1.5	59
146	Meta-analysis of genome-wide scans for hypertension and blood pressure in Caucasians shows evidence of susceptibility regions on chromosomes 2 and 3. <i>Human Molecular Genetics</i> , 2004, 13, 2325-2332.	1.4	58
147	Comprehensive association study of genetic variants in the IL-1 gene family in systemic juvenile idiopathic arthritis. <i>Genes and Immunity</i> , 2008, 9, 349-357.	2.2	58
148	Meta-analysis of linkage studies for Alzheimer's disease—A web resource. <i>Neurobiology of Aging</i> , 2009, 30, 1037-1047.	1.5	58
149	Mutation in the TPA Gene Predicts Intolerance to Azathioprine. <i>Nucleosides, Nucleotides and Nucleic Acids</i> , 2004, 23, 1393-1397.	0.4	57
150	Association of a Locus in the CAMTA1 Gene With Survival in Patients With Sporadic Amyotrophic Lateral Sclerosis. <i>JAMA Neurology</i> , 2016, 73, 812.	4.5	57
151	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
152	Meta-analysis of genome-wide linkage studies of systemic lupus erythematosus. <i>Genes and Immunity</i> , 2006, 7, 609-614.	2.2	56
153	A genetic risk score combining 32 SNPs is associated with body mass index and improves obesity prediction in people with major depressive disorder. <i>BMC Medicine</i> , 2015, 13, 86.	2.3	56
154	Shared genetic contribution to ischemic stroke and Alzheimer's disease. <i>Annals of Neurology</i> , 2016, 79, 739-747.	2.8	56
155	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
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326	Data Acquisition for Meta-Analysis of Genome-Wide Linkage Studies Using the Genome Search Meta-Analysis Method. <i>Human Heredity</i> , 2007, 64, 74-81.	0.4	11
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