

# Jonathan Coleman

## List of Publications by Year in descending order

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Version: 2024-02-01

145  
papers

18,271  
citations

43973

48  
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20307

116  
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229  
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229  
docs citations

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times ranked

17277  
citing authors

#	ARTICLE	IF	CITATIONS
1	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
2	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
3	Psychological trauma and the genetic overlap between posttraumatic stress disorder and major depressive disorder. <i>Psychological Medicine</i> , 2022, 52, 3975-3984.	2.7	11
4	Using major depression polygenic risk scores to explore the depressive symptom continuum. <i>Psychological Medicine</i> , 2022, 52, 149-158.	2.7	9
5	Identifying the Common Genetic Basis of Antidepressant Response. <i>Biological Psychiatry Global Open Science</i> , 2022, 2, 115-126.	1.0	31
6	Common Genetic Variation and Age of Onset of Anorexia Nervosa. <i>Biological Psychiatry Global Open Science</i> , 2022, 2, 368-378.	1.0	10
7	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. <i>Biological Psychiatry</i> , 2022, 91, 313-327.	0.7	114
8	Enhancing Discovery of Genetic Variants for Posttraumatic Stress Disorder Through Integration of Quantitative Phenotypes and Trauma Exposure Information. <i>Biological Psychiatry</i> , 2022, 91, 626-636.	0.7	21
9	Genome-wide by Environment Interaction Study of Stressful Life Events and Hospital-Treated Depression in the iPSYCH2012 Sample. <i>Biological Psychiatry Global Open Science</i> , 2022, 2, 400-410.	1.0	2
10	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
11	Assessing the Evidence for Causal Associations Between Body Mass Index, C-Reactive Protein, Depression and Reported Trauma Using Mendelian Randomization. <i>Biological Psychiatry Global Open Science</i> , 2022, , .	1.0	4
12	Genetic and early environmental predictors of adulthood self-reports of trauma. <i>British Journal of Psychiatry</i> , 2022, 221, 613-620.	1.7	9
13	Latent subtypes of manic and/or irritable episode symptoms in two population-based cohorts – ERRATUM. <i>British Journal of Psychiatry</i> , 2022, , 1-2.	1.7	0
14	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
15	Editorial: Genome-wide Association Studies of Internalizing Symptoms: A Big Step on a Long Road. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2022, , .	0.3	0
16	Genetics and neurobiology of eating disorders. <i>Nature Neuroscience</i> , 2022, 25, 543-554.	7.1	31
17	Feasibility and application of polygenic score analysis to the morphology of human-induced pluripotent stem cells. <i>Molecular Genetics and Genomics</i> , 2022, 297, 1111-1122.	1.0	3
18	Association of polygenic score for major depression with response to lithium in patients with bipolar disorder. <i>Molecular Psychiatry</i> , 2021, 26, 2457-2470.	4.1	44

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19	Shared genetic risk between eating disorder and substance use-related phenotypes: Evidence from genome-wide association studies. <i>Addiction Biology</i> , 2021, 26, e12880.	1.4	28
20	No Evidence for Passive Gene-Environment Correlation or the Influence of Genetic Risk for Psychiatric Disorders on Adult Body Composition via the Adoption Design. <i>Behavior Genetics</i> , 2021, 51, 58-67.	1.4	2
21	Bipolar multiplex families have an increased burden of common risk variants for psychiatric disorders. <i>Molecular Psychiatry</i> , 2021, 26, 1286-1298.	4.1	33
22	Imputed gene expression risk scores: a functionally informed component of polygenic risk. <i>Human Molecular Genetics</i> , 2021, 30, 727-738.	1.4	11
23	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
24	PTSD Genome-Wide Association Study Identifies Novel Loci and Informs Future Expectations. <i>Biological Psychiatry</i> , 2021, 89, S69-S70.	0.7	0
25	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
26	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
27	Evaluation of polygenic prediction methodology within a reference-standardized framework. <i>PLoS Genetics</i> , 2021, 17, e1009021.	1.5	99
28	Sex differences in experiences of multiple traumas and mental health problems in the UK Biobank cohort. <i>Social Psychiatry and Psychiatric Epidemiology</i> , 2021, , 1.	1.6	8
29	Examining Sex-Differentiated Genetic Effects Across Neuropsychiatric and Behavioral Traits. <i>Biological Psychiatry</i> , 2021, 89, 1127-1137.	0.7	48
30	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
31	Genes in treatment: Polygenic risk scores for different psychopathologies, neuroticism, educational attainment and IQ and the outcome of two different exposure-based fear treatments. <i>World Journal of Biological Psychiatry</i> , 2021, 22, 699-712.	1.3	0
32	Examining Individual and Synergistic Contributions of PTSD and Genetics to Blood Pressure: A Trans-Ethnic Meta-Analysis. <i>Frontiers in Neuroscience</i> , 2021, 15, 678503.	1.4	10
33	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
34	Exploring the genetic heterogeneity in major depression across diagnostic criteria. <i>Molecular Psychiatry</i> , 2021, 26, 7337-7345.	4.1	18
35	Predicting clinical outcome to specialist multimodal inpatient treatment in patients with treatment resistant depression. <i>Journal of Affective Disorders</i> , 2021, 291, 188-197.	2.0	5
36	Evaluation of Genotype-Based Gene Expression Model Performance: A Cross-Framework and Cross-Dataset Study. <i>Genes</i> , 2021, 12, 1531.	1.0	2

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37	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. <i>JAMA Psychiatry</i> , 2021, 78, 1258.	6.0	88
38	Comparison of depression and anxiety symptom networks in reporters and non-reporters of lifetime trauma in two samples of differing severity. <i>Journal of Affective Disorders Reports</i> , 2021, 6, 100201.	0.9	4
39	The Validity of Brief Phenotyping in Population Biobanks for Psychiatric Genome-Wide Association Studies on the Biobank Scale. <i>Complex Psychiatry</i> , 2021, 7, 11-15.	1.3	10
40	<scp>Self-reported</scp> medication use as an alternate phenotyping method for anxiety and depression in the <scp>UK</scp> Biobank. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2021, 186, 389-398.	1.1	3
41	Association Between Genetic Risk for Psychiatric Disorders and the Probability of Living in Urban Settings. <i>JAMA Psychiatry</i> , 2021, 78, 1355.	6.0	20
42	Sociodemographic factors associated with treatment-seeking and treatment receipt: cross-sectional analysis of UK Biobank participants with lifetime generalised anxiety or major depressive disorder. <i>BJPsych Open</i> , 2021, 7, .	0.3	6
43	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
44	Trauma and posttraumatic stress disorder modulate polygenic predictors of hippocampal and amygdala volume. <i>Translational Psychiatry</i> , 2021, 11, 637.	2.4	4
45	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
46	The Genetics of the Mood Disorder Spectrum: Genome-wide Association Analyses of More Than 185,000 Cases and 439,000 Controls. <i>Biological Psychiatry</i> , 2020, 88, 169-184.	0.7	137
47	A major role for common genetic variation in anxiety disorders. <i>Molecular Psychiatry</i> , 2020, 25, 3292-3303.	4.1	243
48	Cannabis use, depression and self-harm: phenotypic and genetic relationships. <i>Addiction</i> , 2020, 115, 482-492.	1.7	29
49	Molecular genetic overlap between posttraumatic stress disorder and sleep phenotypes. <i>Sleep</i> , 2020, 43, .	0.6	32
50	Familial Influences on Neuroticism and Education in the UK Biobank. <i>Behavior Genetics</i> , 2020, 50, 84-93.	1.4	9
51	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
52	Genome-wide association study of intracranial aneurysms identifies 17 risk loci and genetic overlap with clinical risk factors. <i>Nature Genetics</i> , 2020, 52, 1303-1313.	9.4	163
53	An Exposure-Wide and Mendelian Randomization Approach to Identifying Modifiable Factors for the Prevention of Depression. <i>American Journal of Psychiatry</i> , 2020, 177, 944-954.	4.0	119
54	Genome-wide Meta-analysis Finds the ACSL5-ZDHHC6 Locus Is Associated with ALS and Links Weight Loss to the Disease Genetics. <i>Cell Reports</i> , 2020, 33, 108323.	2.9	41

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55	A phenome-wide association and Mendelian Randomisation study of polygenic risk for depression in UK Biobank. <i>Nature Communications</i> , 2020, 11, 2301.	5.8	81
56	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
57	Genetic stratification of depression in UK Biobank. <i>Translational Psychiatry</i> , 2020, 10, 163.	2.4	19
58	Genomic influences on self-reported childhood maltreatment. <i>Translational Psychiatry</i> , 2020, 10, 38.	2.4	47
59	Mental health in UK Biobank – development, implementation and results from an online questionnaire completed by 157 366 participants: a reanalysis. <i>BJPsych Open</i> , 2020, 6, e18.	0.3	210
60	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
61	The genetic and environmental hierarchical structure of anxiety and depression in the UK Biobank. <i>Depression and Anxiety</i> , 2020, 37, 512-520.	2.0	25
62	Shared Genetic Risk Between Psychiatric and Cognitive Symptoms in Huntington’s Disease and in the General Population. <i>Biological Psychiatry</i> , 2020, 87, e25-e27.	0.7	1
63	Genetic identification of cell types underlying brain complex traits yields insights into the etiology of Parkinson’s disease. <i>Nature Genetics</i> , 2020, 52, 482-493.	9.4	216
64	Comparison of Adopted and Nonadopted Individuals Reveals Gene-Environment Interplay for Education in the UK Biobank. <i>Psychological Science</i> , 2020, 31, 582-591.	1.8	71
65	Multivariable G-E interplay in the prediction of educational achievement. <i>PLoS Genetics</i> , 2020, 16, e1009153.	1.5	30
66	Multivariable G-E interplay in the prediction of educational achievement. , 2020, 16, e1009153.		0
67	Multivariable G-E interplay in the prediction of educational achievement. , 2020, 16, e1009153.		0
68	Multivariable G-E interplay in the prediction of educational achievement. , 2020, 16, e1009153.		0
69	Multivariable G-E interplay in the prediction of educational achievement. , 2020, 16, e1009153.		0
70	Genome-wide association study identifies eight risk loci and implicates metabo-psychiatric origins for anorexia nervosa. <i>Nature Genetics</i> , 2019, 51, 1207-1214.	9.4	641
71	FUNCTIONAL CONSEQUENCES OF GENETIC LOCI ASSOCIATED WITH IQ IN A META-ANALYSIS OF 87,740 INDIVIDUALS. <i>European Neuropsychopharmacology</i> , 2019, 29, S809-S810.	0.3	0
72	M17 EVALUATING PREDICTIVE ABILITY OF FUNCTIONALLY INFORMED GENETIC RISK SCORES. <i>European Neuropsychopharmacology</i> , 2019, 29, S175.	0.3	0

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73	International meta-analysis of PTSD genome-wide association studies identifies sex- and ancestry-specific genetic risk loci. <i>Nature Communications</i> , 2019, 10, 4558.	5.8	363
74	Indicators of mental disorders in UK Biobank – A comparison of approaches. <i>International Journal of Methods in Psychiatric Research</i> , 2019, 28, e1796.	1.1	77
75	T30AN INVESTIGATION OF THE SHARED GENETIC ETIOLOGY OF DEPRESSION AND AUTOIMMUNE DISEASES: PLEIOTROPY AND SUB-GROUP HETEROGENEITY. <i>European Neuropsychopharmacology</i> , 2019, 29, S233-S234.	0.3	0
76	64GENOME-WIDE ASSOCIATION STUDY OF SCHOOL GRADES INFORMS COGNITIVE GENETIC ARCHITECTURE OF SIX MAJOR PSYCHIATRIC DISORDERS. <i>European Neuropsychopharmacology</i> , 2019, 29, S1103-S1104.	0.3	0
77	49 EXPLORING THE RELATIONSHIP BETWEEN POST TRAUMATIC STRESS DISORDER AND MAJOR DEPRESSIVE DISORDER IN THE CONTEXT OF DIFFERENT TRAUMA TYPES. <i>European Neuropsychopharmacology</i> , 2019, 29, S87.	0.3	0
78	S9GENETIC INFLUENCES ON ANXIETY AND DEPRESSION OUTCOMES FOLLOWING TREATMENTS IN A CLINICALLY RELEVANT POPULATION-BASED COHORT, THE GLAD STUDY. <i>European Neuropsychopharmacology</i> , 2019, 29, S118.	0.3	0
79	The Genetic Links to Anxiety and Depression (GLAD) Study: Online recruitment into the largest recontactable study of depression and anxiety. <i>Behaviour Research and Therapy</i> , 2019, 123, 103503.	1.6	47
80	M39 CHILDHOOD ADOPTION AND BODY COMPOSITION IN ADULTHOOD: THE ROLE OF GENETIC CONFOUNDING IN CONTEXT OF ANOREXIA NERVOSA. <i>European Neuropsychopharmacology</i> , 2019, 29, S187.	0.3	0
81	1GENETIC COMORBIDITY BETWEEN DEPRESSION AND CARDIO-METABOLIC DISEASE, STRATIFIED BY AGE AT ONSET. <i>European Neuropsychopharmacology</i> , 2019, 29, S1066.	0.3	1
82	4DETERMINING THE RELATIONSHIP BETWEEN CANNABIS USE AND MAJOR DEPRESSION IN UK BIOBANK. <i>European Neuropsychopharmacology</i> , 2019, 29, S1067-S1068.	0.3	0
83	Mental health in UK Biobank: development, implementation and results from an online questionnaire completed by 157 366 participants – RETRACTED. <i>BJPsych Open</i> , 2019, 5, e56.	0.3	7
84	SA16A MAJOR ROLE FOR COMMON GENETIC VARIATION IN ANXIETY DISORDERS. <i>European Neuropsychopharmacology</i> , 2019, 29, S1196.	0.3	8
85	FEMALE-SPECIFIC GENETIC VARIATION ASSOCIATED WITH BODY FAT PERCENTAGE MAY CONTRIBUTE TO RISK FOR ANOREXIA NERVOSA. <i>European Neuropsychopharmacology</i> , 2019, 29, S1048.	0.3	0
86	SA41GENOME-WIDE GENE-ENVIRONMENT ANALYSES OF DEPRESSION AND REPORTED LIFETIME TRAUMATIC EXPERIENCES IN UK BIOBANK. <i>European Neuropsychopharmacology</i> , 2019, 29, S1210-S1211.	0.3	1
87	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
88	Assessment of Bidirectional Relationships Between Physical Activity and Depression Among Adults. <i>JAMA Psychiatry</i> , 2019, 76, 399.	6.0	399
89	DNA methylation of FKBP5 and response to exposure-based psychological therapy. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019, 180, 150-158.	1.1	44
90	Genome-wide by environment interaction studies of depressive symptoms and psychosocial stress in UK Biobank and Generation Scotland. <i>Translational Psychiatry</i> , 2019, 9, 14.	2.4	87

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91	Genetic influences on treatment-seeking for common mental health problems in the UK biobank. Behaviour Research and Therapy, 2019, 121, 103413.	1.6	7
92	A genome-wide association meta-analysis of prognostic outcomes following cognitive behavioural therapy in individuals with anxiety and depressive disorders. Translational Psychiatry, 2019, 9, 150.	2.4	35
93	ukbtools: An R package to manage and query UK Biobank data. PLoS ONE, 2019, 14, e0214311.	1.1	37
94	GWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. American Journal of Psychiatry, 2019, 176, 651-660.	4.0	186
95	Integrated analysis of environmental and genetic influences on cord blood DNA methylation in new-borns. Nature Communications, 2019, 10, 2548.	5.8	94
96	152. Taking a Closer Look at PTSD Genomics: Rare Copy Number Variants and Extended Phenotyping. Biological Psychiatry, 2019, 85, S63.	0.7	0
97	Genome-wide association study identifies 30 loci associated with bipolar disorder. Nature Genetics, 2019, 51, 793-803.	9.4	1,191
98	Genetic Overlap Between Alzheimer's Disease and Bipolar Disorder Implicates the MARK2 and VAC14 Genes. Frontiers in Neuroscience, 2019, 13, 220.	1.4	42
99	Identification of common genetic risk variants for autism spectrum disorder. Nature Genetics, 2019, 51, 431-444.	9.4	1,538
100	F30GENETIC STRUCTURE WITHIN THE UK BIOBANK MENTAL HEALTH QUESTIONNAIRE. European Neuropsychopharmacology, 2019, 29, S1125-S1126.	0.3	0
101	Genetic correlations of psychiatric traits with body composition and glycemic traits are sex- and age-dependent. Nature Communications, 2019, 10, 5765.	5.8	59
102	SU39GENETIC VARIATION IN THE MAJOR HISTOCOMPATIBILITY COMPLEX AND ASSOCIATION WITH DEPRESSION. European Neuropsychopharmacology, 2019, 29, S1288-S1289.	0.3	0
103	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. Cell, 2019, 179, 1469-1482.e11.	13.5	935
104	Association of Whole-Genome and NETRIN1 Signaling Pathway-Derived Polygenic Risk Scores for Major Depressive Disorder and White Matter Microstructure in the UK Biobank. Biological Psychiatry: Cognitive Neuroscience and Neuroimaging, 2019, 4, 91-100.	1.1	16
105	Genomics of body fat percentage may contribute to sex bias in anorexia nervosa. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, 428-438.	1.1	87
106	Evidence for increased genetic risk load for major depression in patients assigned to electroconvulsive therapy. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, 35-45.	1.1	18
107	Biological annotation of genetic loci associated with intelligence in a meta-analysis of 87,740 individuals. Molecular Psychiatry, 2019, 24, 182-197.	4.1	47
108	The utility of the SCAS-C/P to detect specific anxiety disorders among clinically anxious children.. Psychological Assessment, 2019, 31, 1006-1018.	1.2	17



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109	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
110	Improving genetic prediction by leveraging genetic correlations among human diseases and traits. <i>Nature Communications</i> , 2018, 9, 989.	5.8	136
111	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
112	Genetic influence on social outcomes during and after the Soviet era in Estonia. <i>Nature Human Behaviour</i> , 2018, 2, 269-275.	6.2	74
113	Association of Polygenic Risk for Attention-Deficit/Hyperactivity Disorder With Co-occurring Traits and Disorders. <i>Biological Psychiatry: Cognitive Neuroscience and Neuroimaging</i> , 2018, 3, 635-643.	1.1	57
114	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
115	The impact of treatment delivery format on response to cognitive behaviour therapy for preadolescent children with anxiety disorders. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2018, 59, 763-772.	3.1	25
116	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
117	Investigation of common, low-frequency and rare genome-wide variation in anorexia nervosa. <i>Molecular Psychiatry</i> , 2018, 23, 1169-1180.	4.1	32
118	Evidence for gene-environment correlation in child feeding: Links between common genetic variation for BMI in children and parental feeding practices. <i>PLoS Genetics</i> , 2018, 14, e1007757.	1.5	67
119	Genome-wide interaction study of a proxy for stress-sensitivity and its prediction of major depressive disorder. <i>PLoS ONE</i> , 2018, 13, e0209160.	1.1	14
120	A polygenic p factor for major psychiatric disorders. <i>Translational Psychiatry</i> , 2018, 8, 205.	2.4	117
121	Applying polygenic risk scoring for psychiatric disorders to a large family with bipolar disorder and major depressive disorder. <i>Communications Biology</i> , 2018, 1, 163.	2.0	17
122	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
123	Genome-wide association meta-analysis in 269,867 individuals identifies new genetic and functional links to intelligence. <i>Nature Genetics</i> , 2018, 50, 912-919.	9.4	893
124	Individual and shared effects of social environment and polygenic risk scores on adolescent body mass index. <i>Scientific Reports</i> , 2018, 8, 6344.	1.6	10
125	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. <i>Cell</i> , 2018, 173, 1705-1715.e16.	13.5	623
126	158. Exploring the Common Genetic Architecture of PTSD Symptoms in the UK Biobank. <i>Biological Psychiatry</i> , 2018, 83, S64.	0.7	5



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127	Separate and combined effects of genetic variants and pre-treatment whole blood gene expression on response to exposure-based cognitive behavioural therapy for anxiety disorders. <i>World Journal of Biological Psychiatry</i> , 2017, 18, 215-226.	1.3	9
128	Significant Locus and Metabolic Genetic Correlations Revealed in Genome-Wide Association Study of Anorexia Nervosa. <i>American Journal of Psychiatry</i> , 2017, 174, 850-858.	4.0	410
129	Genome-wide association meta-analysis of 78,308 individuals identifies new loci and genes influencing human intelligence. <i>Nature Genetics</i> , 2017, 49, 1107-1112.	9.4	425
130	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
131	Genetic Association of Major Depression With Atypical Features and Obesity-Related Immunometabolic Dysregulations. <i>JAMA Psychiatry</i> , 2017, 74, 1214.	6.0	174
132	Genome-wide expression and response to exposure-based psychological therapy for anxiety disorders. <i>Translational Psychiatry</i> , 2017, 7, e1219-e1219.	2.4	16
133	Hair Cortisol in Twins: Heritability and Genetic Overlap with Psychological Variables and Stress-System Genes. <i>Scientific Reports</i> , 2017, 7, 15351.	1.6	50
134	Genetic variation in the endocannabinoid system and response to Cognitive Behavior Therapy for child anxiety disorders. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 144-155.	1.1	23
135	An Examination of Polygenic Score Risk Prediction in Individuals With First-Episode Psychosis. <i>Biological Psychiatry</i> , 2017, 81, 470-477.	0.7	176
136	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
137	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
138	Quality control, imputation and analysis of genome-wide genotyping data from the Illumina HumanCoreExome microarray. <i>Briefings in Functional Genomics</i> , 2016, 15, 298-304.	1.3	65
139	Non-replication of the association between 5HTTLPR and response to psychological therapy for child anxiety disorders. <i>British Journal of Psychiatry</i> , 2016, 208, 182-188.	1.7	25
140	Genome-wide investigation of schizophrenia associated plasma Nde1 enzyme activity. <i>Schizophrenia Research</i> , 2016, 172, 60-67.	1.1	10
141	Genome-Wide Association of Heroin Dependence in Han Chinese. <i>PLoS ONE</i> , 2016, 11, e0167388.	1.1	30
142	High definition versus standard definition white light endoscopy for detecting dysplasia in patients with Barrett's esophagus. <i>Ecological Management and Restoration</i> , 2015, 28, 742-749.	0.2	47
143	Clinical Predictors of Response to Cognitive-Behavioral Therapy in Pediatric Anxiety Disorders: The Genes for Treatment (GxT) Study. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2015, 54, 454-463.	0.3	118
144	HPA AXIS RELATED GENES AND RESPONSE TO PSYCHOLOGICAL THERAPIES: GENETICS AND EPIGENETICS. <i>Depression and Anxiety</i> , 2015, 32, 861-870.	2.0	75

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145	Serotonin transporter methylation and response to cognitive behaviour therapy in children with anxiety disorders. <i>Translational Psychiatry</i> , 2014, 4, e444-e444.	2.4	97