

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
h-index

20307

116
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229
all docs

229
docs citations

229
times ranked

17277
citing authors

#	ARTICLE	IF	CITATIONS
1	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
2	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
3	Identification of common genetic risk variants for autism spectrum disorder. <i>Nature Genetics</i> , 2019, 51, 431-444.	9.4	1,538
4	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019, 51, 793-803.	9.4	1,191
5	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , 2019, 179, 1469-1482.e11.	13.5	935
6	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
7	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
8	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
9	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. <i>Cell</i> , 2018, 173, 1705-1715.e16.	13.5	623
10	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
11	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
12	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
13	Assessment of Bidirectional Relationships Between Physical Activity and Depression Among Adults. <i>JAMA Psychiatry</i> , 2019, 76, 399.	6.0	399
14	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
15	A major role for common genetic variation in anxiety disorders. <i>Molecular Psychiatry</i> , 2020, 25, 3292-3303.	4.1	243
16	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
17	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
18	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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19	An Examination of Polygenic Score Risk Prediction in Individuals With First-Episode Psychosis. <i>Biological Psychiatry</i> , 2017, 81, 470-477.	0.7	176
20	Genetic Association of Major Depression With Atypical Features and Obesity-Related Immunometabolic Dysregulations. <i>JAMA Psychiatry</i> , 2017, 74, 1214.	6.0	174
21	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
22	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
23	Improving genetic prediction by leveraging genetic correlations among human diseases and traits. <i>Nature Communications</i> , 2018, 9, 989.	5.8	136
24	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
25	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
26	A polygenic p factor for major psychiatric disorders. <i>Translational Psychiatry</i> , 2018, 8, 205.	2.4	117
27	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
28	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
29	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
30	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
31	Evaluation of polygenic prediction methodology within a reference-standardized framework. <i>PLoS Genetics</i> , 2021, 17, e1009021.	1.5	99
32	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
33	Integrated analysis of environmental and genetic influences on cord blood DNA methylation in new-borns. <i>Nature Communications</i> , 2019, 10, 2548.	5.8	94
34	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
35	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. <i>JAMA Psychiatry</i> , 2021, 78, 1258.	6.0	88
36	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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37	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
38	Genomics of body fat percentage may contribute to sex bias in anorexia nervosa. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019, 180, 428-438.	1.1	87
39	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
40	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
41	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
42	HPA AXIS RELATED GENES AND RESPONSE TO PSYCHOLOGICAL THERAPIES: GENETICS AND EPIGENETICS. <i>Depression and Anxiety</i> , 2015, 32, 861-870.	2.0	75
43	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
44	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
45	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
46	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
47	Genetic correlations of psychiatric traits with body composition and glycemic traits are sex- and age-dependent. <i>Nature Communications</i> , 2019, 10, 5765.	5.8	59
48	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
49	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
50	Examining Sex-Differentiated Genetic Effects Across Neuropsychiatric and Behavioral Traits. <i>Biological Psychiatry</i> , 2021, 89, 1127-1137.	0.7	48
51	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
52	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
53	Biological annotation of genetic loci associated with intelligence in a meta-analysis of 87,740 individuals. <i>Molecular Psychiatry</i> , 2019, 24, 182-197.	4.1	47
54	Genomic influences on self-reported childhood maltreatment. <i>Translational Psychiatry</i> , 2020, 10, 38.	2.4	47

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55	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
56	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
57	Genetic Overlap Between Alzheimer's Disease and Bipolar Disorder Implicates the MARK2 and VAC14 Genes. <i>Frontiers in Neuroscience</i> , 2019, 13, 220.	1.4	42
58	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
59	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
60	ukbtools: An R package to manage and query UK Biobank data. <i>PLoS ONE</i> , 2019, 14, e0214311.	1.1	37
61	A genome-wide association meta-analysis of prognostic outcomes following cognitive behavioural therapy in individuals with anxiety and depressive disorders. <i>Translational Psychiatry</i> , 2019, 9, 150.	2.4	35
62	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
63	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
64	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
65	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
66	Investigation of common, low-frequency and rare genome-wide variation in anorexia nervosa. <i>Molecular Psychiatry</i> , 2018, 23, 1169-1180.	4.1	32
67	Molecular genetic overlap between posttraumatic stress disorder and sleep phenotypes. <i>Sleep</i> , 2020, 43, .	0.6	32
68	Identifying the Common Genetic Basis of Antidepressant Response. <i>Biological Psychiatry Global Open Science</i> , 2022, 2, 115-126.	1.0	31
69	Genetics and neurobiology of eating disorders. <i>Nature Neuroscience</i> , 2022, 25, 543-554.	7.1	31
70	Multivariable G-E interplay in the prediction of educational achievement. <i>PLoS Genetics</i> , 2020, 16, e1009153.	1.5	30
71	Genome-Wide Association of Heroin Dependence in Han Chinese. <i>PLoS ONE</i> , 2016, 11, e0167388.	1.1	30
72	Cannabis use, depression and self-harm: phenotypic and genetic relationships. <i>Addiction</i> , 2020, 115, 482-492.	1.7	29

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73	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
74	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
75	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
76	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
77	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
78	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
79	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
80	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
81	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
82	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
83	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
84	Genetic stratification of depression in UK Biobank. <i>Translational Psychiatry</i> , 2020, 10, 163.	2.4	19
85	Evidence for increased genetic risk load for major depression in patients assigned to electroconvulsive therapy. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019, 180, 35-45.	1.1	18
86	Exploring the genetic heterogeneity in major depression across diagnostic criteria. <i>Molecular Psychiatry</i> , 2021, 26, 7337-7345.	4.1	18
87	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
88	The utility of the SCAS-C/P to detect specific anxiety disorders among clinically anxious children. <i>Psychological Assessment</i> , 2019, 31, 1006-1018.	1.2	17
89	Genome-wide expression and response to exposure-based psychological therapy for anxiety disorders. <i>Translational Psychiatry</i> , 2017, 7, e1219-e1219.	2.4	16
90	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

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91	Association of Whole-Genome and NETRIN1 Signaling Pathwayâ€™Derived Polygenic Risk Scores for Major Depressive Disorder and White Matter Microstructure in the UK Biobank. <i>Biological Psychiatry: Cognitive Neuroscience and Neuroimaging</i> , 2019, 4, 91-100.	1.1	16
92	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
93	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
94	Imputed gene expression risk scores: a functionally informed component of polygenic risk. <i>Human Molecular Genetics</i> , 2021, 30, 727-738.	1.4	11
95	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
96	Genome-wide investigation of schizophrenia associated plasma Ndel1 enzyme activity. <i>Schizophrenia Research</i> , 2016, 172, 60-67.	1.1	10
97	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
98	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
99	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
100	Common Genetic Variation and Age of Onset of Anorexia Nervosa. <i>Biological Psychiatry Global Open Science</i> , 2022, 2, 368-378.	1.0	10
101	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
102	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
103	Familial Influences on Neuroticism and Education in the UK Biobank. <i>Behavior Genetics</i> , 2020, 50, 84-93.	1.4	9
104	Using major depression polygenic risk scores to explore the depressive symptom continuum. <i>Psychological Medicine</i> , 2022, 52, 149-158.	2.7	9
105	Genetic and early environmental predictors of adulthood self-reports of trauma. <i>British Journal of Psychiatry</i> , 2022, 221, 613-620.	1.7	9
106	SA16A MAJOR ROLE FOR COMMON GENETIC VARIATION IN ANXIETY DISORDERS. <i>European Neuropsychopharmacology</i> , 2019, 29, S1196.	0.3	8
107	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
108	Mental health in UK Biobank: development, implementation and results from an online questionnaire completed by 157 366 participants â€™ RETRACTED. <i>BJPsych Open</i> , 2019, 5, e56.	0.3	7

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109	Genetic influences on treatment-seeking for common mental health problems in the UK biobank. <i>Behaviour Research and Therapy</i> , 2019, 121, 103413.	1.6	7
110	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
111	158. Exploring the Common Genetic Architecture of PTSD Symptoms in the UK Biobank. <i>Biological Psychiatry</i> , 2018, 83, S64.	0.7	5
112	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
113	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
114	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
115	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
116	Trauma and posttraumatic stress disorder modulate polygenic predictors of hippocampal and amygdala volume. <i>Translational Psychiatry</i> , 2021, 11, 637.	2.4	4
117	Exploring polygenicâ€environment and residualâ€environment interactions for depressive symptoms within the UK Biobank. <i>Genetic Epidemiology</i> , 2022, 46, 219-233.	0.6	4
118	<scp>Selfâ€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
119	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
120	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
121	Evaluation of Genotype-Based Gene Expression Model Performance: A Cross-Framework and Cross-Dataset Study. <i>Genes</i> , 2021, 12, 1531.	1.0	2
122	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
123	1GENETIC COMORBIDITY BETWEEN DEPRESSION AND CARDIO-METABOLIC DISEASE, STRATIFIED BY AGE AT ONSET. <i>European Neuropsychopharmacology</i> , 2019, 29, S1066.	0.3	1
124	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
125	Shared Genetic Risk Between Psychiatric and Cognitive Symptoms in Huntingtonâ€™s Disease and in the General Population. <i>Biological Psychiatry</i> , 2020, 87, e25-e27.	0.7	1
126	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

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127	M17 EVALUATING PREDICTIVE ABILITY OF FUNCTIONALLY INFORMED GENETIC RISK SCORES. European Neuropsychopharmacology, 2019, 29, S175.	0.3	0
128	T30AN INVESTIGATION OF THE SHARED GENETIC ETIOLOGY OF DEPRESSION AND AUTOIMMUNE DISEASES: PLEIOTROPY AND SUB-GROUP HETEROGENEITY. European Neuropsychopharmacology, 2019, 29, S233-S234.	0.3	0
129	64GENOME-WIDE ASSOCIATION STUDY OF SCHOOL GRADES INFORMS COGNITIVE GENETIC ARCHITECTURE OF SIX MAJOR PSYCHIATRIC DISORDERS. European Neuropsychopharmacology, 2019, 29, S1103-S1104.	0.3	0
130	49 EXPLORING THE RELATIONSHIP BETWEEN POST TRAUMATIC STRESS DISORDER AND MAJOR DEPRESSIVE DISORDER IN THE CONTEXT OF DIFFERENT TRAUMA TYPES. European Neuropsychopharmacology, 2019, 29, S87.	0.3	0
131	S9GENETIC INFLUENCES ON ANXIETY AND DEPRESSION OUTCOMES FOLLOWING TREATMENTS IN A CLINICALLY RELEVANT POPULATION-BASED COHORT, THE GLAD STUDY. European Neuropsychopharmacology, 2019, 29, S118.	0.3	0
132	M39 CHILDHOOD ADOPTION AND BODY COMPOSITION IN ADULTHOOD: THE ROLE OF GENETIC CONFOUNDING IN CONTEXT OF ANOREXIA NERVOSA. European Neuropsychopharmacology, 2019, 29, S187.	0.3	0
133	4DETERMINING THE RELATIONSHIP BETWEEN CANNABIS USE AND MAJOR DEPRESSION IN UK BIOBANK. European Neuropsychopharmacology, 2019, 29, S1067-S1068.	0.3	0
134	FEMALE-SPECIFIC GENETIC VARIATION ASSOCIATED WITH BODY FAT PERCENTAGE MAY CONTRIBUTE TO RISK FOR ANOREXIA NERVOSA. European Neuropsychopharmacology, 2019, 29, S1048.	0.3	0
135	152. Taking a Closer Look at PTSD Genomics: Rare Copy Number Variants and Extended Phenotyping. Biological Psychiatry, 2019, 85, S63.	0.7	0
136	F30GENETIC STRUCTURE WITHIN THE UK BIOBANK MENTAL HEALTH QUESTIONNAIRE. European Neuropsychopharmacology, 2019, 29, S1125-S1126.	0.3	0
137	SU39GENETIC VARIATION IN THE MAJOR HISTOCOMPATIBILITY COMPLEX AND ASSOCIATION WITH DEPRESSION. European Neuropsychopharmacology, 2019, 29, S1288-S1289.	0.3	0
138	PTSD Genome-Wide Association Study Identifies Novel Loci and Informs Future Expectations. Biological Psychiatry, 2021, 89, S69-S70.	0.7	0
139	Genes in treatment: Polygenic risk scores for different psychopathologies, neuroticism, educational attainment and IQ and the outcome of two different exposure-based fear treatments. World Journal of Biological Psychiatry, 2021, 22, 699-712.	1.3	0
140	Latent subtypes of manic and/or irritable episode symptoms in two population-based cohorts “ERRATUM. British Journal of Psychiatry, 2022, , 1-2.	1.7	0
141	Multivariable G-E interplay in the prediction of educational achievement. , 2020, 16, e1009153.		0
142	Multivariable G-E interplay in the prediction of educational achievement. , 2020, 16, e1009153.		0
143	Multivariable G-E interplay in the prediction of educational achievement. , 2020, 16, e1009153.		0
144	Multivariable G-E interplay in the prediction of educational achievement. , 2020, 16, e1009153.		0

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145	Editorial: Genome-wide Association Studies of Internalizing Symptoms: A Big Step on a Long Road. Journal of the American Academy of Child and Adolescent Psychiatry, 2022, , .	0.3	0