

# Peter McGuffin

## List of Publications by Year in descending order

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Version: 2024-02-01

579  
papers

62,140  
citations

1294

109  
h-index

1250

226  
g-index

621  
all docs

621  
docs citations

621  
times ranked

46514  
citing authors

#	ARTICLE	IF	CITATIONS
1	Identifying the Common Genetic Basis of Antidepressant Response. <i>Biological Psychiatry Global Open Science</i> , 2022, 2, 115-126.	1.0	31
2	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
3	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
4	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
5	Genome-wide association study of suicidal behaviour severity in mood disorders. <i>World Journal of Biological Psychiatry</i> , 2021, 22, 1-19.	1.3	3
6	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
7	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. <i>JAMA Psychiatry</i> , 2021, 78, 1258.	6.0	88
8	Investigating rare pathogenic/likely pathogenic exonic variation in bipolar disorder. <i>Molecular Psychiatry</i> , 2021, 26, 5239-5250.	4.1	15
9	Dimensions of temperament and character as predictors of antidepressant discontinuation, response and adverse reactions during treatment with nortriptyline and escitalopram. <i>Psychological Medicine</i> , 2021, , 1-9.	2.7	3
10	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
11	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
12	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
13	Effect of antidepressant switching between nortriptyline and escitalopram after a failed first antidepressant treatment among patients with major depressive disorder. <i>British Journal of Psychiatry</i> , 2019, 215, 494-501.	1.7	10
14	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
15	Associations Between Attention-Deficit/Hyperactivity Disorder and Various Eating Disorders: A Swedish Nationwide Population Study Using Multiple Genetically Informative Approaches. <i>Biological Psychiatry</i> , 2019, 86, 577-586.	0.7	43
16	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019, 51, 793-803.	9.4	1,191
17	Genome-wide Burden of Rare Short Deletions Is Enriched in Major Depressive Disorder in Four Cohorts. <i>Biological Psychiatry</i> , 2019, 85, 1065-1073.	0.7	25
18	Trajectories of Suicidal Ideation During 12 Weeks of Escitalopram or Nortriptyline Antidepressant Treatment Among 811 Patients With Major Depressive Disorder. <i>Journal of Clinical Psychiatry</i> , 2019, 80,	1.1	7

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19	Association of Whole-Genome and NETRIN1 Signaling Pathwayâ€œDerived Polygenic Risk Scores for Major Depressive Disorder and White Matter Microstructure in the UK Biobank. <i>Biological Psychiatry: Cognitive Neuroscience and Neuroimaging</i> , 2019, 4, 91-100.	1.1	16
20	Family functioning, trauma exposure and PTSD: A cross sectional study. <i>Journal of Affective Disorders</i> , 2019, 245, 645-652.	2.0	10
21	Antidepressant drug-specific prediction of depression treatment outcomes from genetic and clinical variables. <i>Scientific Reports</i> , 2018, 8, 5530.	1.6	51
22	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
23	One year double blind study of high vs low frequency subcallosal cingulate stimulation for depression. <i>Journal of Psychiatric Research</i> , 2018, 96, 124-134.	1.5	39
24	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
25	Unravelling the GSK3 $\beta$ -related genotypic interaction network influencing hippocampal volume in recurrent major depressive disorder. <i>Psychiatric Genetics</i> , 2018, 28, 77-84.	0.6	27
26	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
27	Genes associated with anhedonia: a new analysis in a large clinical trial (GENDEP). <i>Translational Psychiatry</i> , 2018, 8, 150.	2.4	19
28	Genetic disposition to inflammation and response to antidepressants in major depressive disorder. <i>Journal of Psychiatric Research</i> , 2018, 105, 17-22.	1.5	18
29	Stressful life events and catechol-O-methyl-transferase (COMT) gene in bipolar disorder. <i>Depression and Anxiety</i> , 2017, 34, 419-426.	2.0	27
30	The DAOA gene is associated with schizophrenia in the Taiwanese population. <i>Psychiatry Research</i> , 2017, 252, 201-207.	1.7	6
31	Pharmacogenetics of antidepressant response: A polygenic approach. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2017, 75, 128-134.	2.5	71
32	Association between C-reactive protein (CRP) with depression symptom severity and specific depressive symptoms in major depression. <i>Brain, Behavior, and Immunity</i> , 2017, 62, 344-350.	2.0	202
33	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
34	Highly polygenic architecture of antidepressant treatment response: Comparative analysis of SSRI and NRI treatment in an animal model of depression. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 235-250.	1.1	10
35	Interaction between the FTO gene, body mass index and depression: meta-analysis of 13701 individuals. <i>British Journal of Psychiatry</i> , 2017, 211, 70-76.	1.7	49
36	Advancing psychiatric genetics through dissecting heterogeneity. <i>Human Molecular Genetics</i> , 2017, 26, R160-R165.	1.4	16

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37	Childhood maltreatment and the medical morbidity in bipolar disorder: a caseâ€“control study. <i>International Journal of Bipolar Disorders</i> , 2017, 5, 30.	0.8	12
38	Irving Gottesman. <i>BJPsych Bulletin</i> , 2017, 41, 124-125.	0.7	1
39	Gottesman, the Enemy of Genetic Determinism and His Role in a Curious Legal Case. <i>Clinical Psychological Science</i> , 2017, 5, 427-428.	2.4	0
40	A Method to Exploit the Structure of Genetic Ancestry Space to Enhance Case-Control Studies. <i>American Journal of Human Genetics</i> , 2016, 98, 857-868.	2.6	21
41	Combining clinical variables to optimize prediction of antidepressant treatment outcomes. <i>Journal of Psychiatric Research</i> , 2016, 78, 94-102.	1.5	149
42	Evaluation of the validity and utility of a transdiagnostic psychosis dimension encompassing schizophrenia and bipolar disorder. <i>British Journal of Psychiatry</i> , 2016, 209, 107-113.	1.7	67
43	Immune signatures and disorder-specific patterns in a cross-disorder gene expression analysis. <i>British Journal of Psychiatry</i> , 2016, 209, 202-208.	1.7	31
44	Transcriptomics and the mechanisms of antidepressant efficacy. <i>European Neuropsychopharmacology</i> , 2016, 26, 105-112.	0.3	19
45	Rare loss-of-function variants in SETD1A are associated with schizophrenia and developmental disorders. <i>Nature Neuroscience</i> , 2016, 19, 571-577.	7.1	388
46	Genome-wide association analysis identifies TXNRD2, ATXN2 and FOXC1 as susceptibility loci for primary open-angle glaucoma. <i>Nature Genetics</i> , 2016, 48, 189-194.	9.4	211
47	Phenotypic Association Analyses With Copy Number Variation in Recurrent Depressive Disorder. <i>Biological Psychiatry</i> , 2016, 79, 329-336.	0.7	21
48	The interaction between stress and genetic factors in the etiopathogenesis of depression. <i>World Psychiatry</i> , 2015, 14, 161-163.	4.8	51
49	The relationship between schizophrenia and rheumatoid arthritis revisited: Genetic and epidemiological analyses. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2015, 168, 81-88.	1.1	29
50	Exploring the role of drug-metabolising enzymes in antidepressant side effects. <i>Psychopharmacology</i> , 2015, 232, 2609-2617.	1.5	31
51	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
52	Modulatory effects of brain-derived neurotrophic factor Val66Met polymorphism on prefrontal regions in major depressive disorder. <i>British Journal of Psychiatry</i> , 2015, 206, 379-384.	1.7	56
53	Authors' reply. <i>British Journal of Psychiatry</i> , 2015, 207, 363-364.	1.7	1
54	Genetic Differences in the Immediate Transcriptome Response to Stress Predict Risk-Related Brain Function and Psychiatric Disorders. <i>Neuron</i> , 2015, 86, 1189-1202.	3.8	102

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55	The successful search for genetic loci associated with depression. <i>Genome Medicine</i> , 2015, 7, 92.	3.6	3
56	DNA Modification Study of Major Depressive Disorder: Beyond Locus-by-Locus Comparisons. <i>Biological Psychiatry</i> , 2015, 77, 246-255.	0.7	66
57	Joint Analysis of Psychiatric Disorders Increases Accuracy of Risk Prediction for Schizophrenia, Bipolar Disorder, and Major Depressive Disorder. <i>American Journal of Human Genetics</i> , 2015, 96, 283-294.	2.6	225
58	A genome-wide association study of suicide severity scores in bipolar disorder. <i>Journal of Psychiatric Research</i> , 2015, 65, 23-29.	1.5	36
59	A comparison of mental state examination documentation by junior clinicians in electronic health records before and after the introduction of a semi-structured assessment template (OPCRIT+). <i>International Journal of Medical Informatics</i> , 2015, 84, 675-682.	1.6	9
60	Putative Transcriptomic Biomarkers in the Inflammatory Cytokine Pathway Differentiate Major Depressive Disorder Patients from Control Subjects and Bipolar Disorder Patients. <i>PLoS ONE</i> , 2014, 9, e91076.	1.1	39
61	Interplay Between Childhood Physical Abuse and Familial Risk in the Onset of Psychotic Disorders. <i>Schizophrenia Bulletin</i> , 2014, 40, 1443-1451.	2.3	41
62	Genetic susceptibility for bipolar disorder and response to antidepressants in major depressive disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2014, 165, 77-83.	1.1	16
63	Mood stabilizers differentially affect housekeeping gene expression in human cells. <i>International Journal of Methods in Psychiatric Research</i> , 2014, 23, 279-288.	1.1	14
64	An Inflammatory Biomarker as a Differential Predictor of Outcome of Depression Treatment With Escitalopram and Nortriptyline. <i>American Journal of Psychiatry</i> , 2014, 171, 1278-1286.	4.0	336
65	Different genetic factors influence specific symptom dimensions of DSM-IV major depression. <i>Evidence-Based Mental Health</i> , 2014, 17, 18-18.	2.2	1
66	Moving from DSM-5 to ICD-11: A joint problem?. <i>Australian and New Zealand Journal of Psychiatry</i> , 2014, 48, 194-196.	1.3	4
67	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
68	Copy number variant study of bipolar disorder in Canadian and UK populations implicates synaptic genes. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2014, 165, 303-313.	1.1	76
69	Interaction between stress and the BDNF Val66Met polymorphism in depression: a systematic review and meta-analysis. <i>BMC Medicine</i> , 2014, 12, 7.	2.3	228
70	Genome-wide association study of bipolar disorder in Canadian and UK populations corroborates disease loci including SYNE1 and CSMD1. <i>BMC Medical Genetics</i> , 2014, 15, 2.	2.1	106
71	Genetic predictors of antidepressant side effects: A grouped candidate gene approach in the Genome-Based Therapeutic Drugs for Depression (GENDEP) study. <i>Journal of Psychopharmacology</i> , 2014, 28, 142-150.	2.0	18
72	Functional effects of polymorphisms on glucocorticoid receptor modulation of human anxiogenic substance-P gene promoter activity in primary amygdala neurones. <i>Psychoneuroendocrinology</i> , 2014, 47, 43-55.	1.3	13

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73	Genetic differences in cytochrome P450 enzymes and antidepressant treatment response. <i>Journal of Psychopharmacology</i> , 2014, 28, 133-141.	2.0	75
74	Comorbid medical illness in bipolar disorder. <i>British Journal of Psychiatry</i> , 2014, 205, 465-472.	1.7	113
75	Chipping away at major depressive disorder. <i>Genome Biology</i> , 2014, 15, 421.	3.8	4
76	Common variants near ABCA1, AFAP1 and GMDS confer risk of primary open-angle glaucoma. <i>Nature Genetics</i> , 2014, 46, 1120-1125.	9.4	186
77	The endogenous and reactive depression subtypes revisited: integrative animal and human studies implicate multiple distinct molecular mechanisms underlying major depressive disorder. <i>BMC Medicine</i> , 2014, 12, 73.	2.3	52
78	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
79	Investigating the genetic variation underlying episodicity in major depressive disorder: Suggestive evidence for a bipolar contribution. <i>Journal of Affective Disorders</i> , 2014, 155, 81-89.	2.0	15
80	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
81	Trauma, post-traumatic stress disorder and psychiatric disorders in a middle-income setting: prevalence and comorbidity. <i>British Journal of Psychiatry</i> , 2014, 205, 383-389.	1.7	53
82	Genome-wide association analysis of copy number variation in recurrent depressive disorder. <i>Molecular Psychiatry</i> , 2013, 18, 183-189.	4.1	45
83	Contribution of Common Genetic Variants to Antidepressant Response. <i>Biological Psychiatry</i> , 2013, 73, 679-682.	0.7	199
84	Allele-specific expression of the serotonin transporter and its transcription factors following lamotrigine treatment in vitro. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2013, 162, 474-483.	1.1	7
85	Residual association at C9orf72 suggests an alternative amyotrophic lateral sclerosis-causing hexanucleotide repeat. <i>Neurobiology of Aging</i> , 2013, 34, 2234.e1-2234.e7.	1.5	22
86	Meta-analysis of genome-wide association studies in five cohorts reveals common variants in RBFox1, a regulator of tissue-specific splicing, associated with refractive error. <i>Human Molecular Genetics</i> , 2013, 22, 2754-2764.	1.4	60
87	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. <i>Nature Genetics</i> , 2013, 45, 984-994.	9.4	2,067
88	Genome-wide association analysis accounting for environmental factors through propensity score matching: Application to stressful life events in major depressive disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2013, 162, 521-529.	1.1	16
89	Resistance to antidepressant treatment is associated with polymorphisms in the leptin gene, decreased leptin mRNA expression, and decreased leptin serum levels. <i>European Neuropsychopharmacology</i> , 2013, 23, 653-662.	0.3	32
90	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

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91	The interaction between child maltreatment, adult stressful life events and the 5-HTTLPR in major depression. <i>Journal of Psychiatric Research</i> , 2013, 47, 1032-1035.	1.5	21
92	Interaction between specific forms of childhood maltreatment and the serotonin transporter gene (5-HTT) in recurrent depressive disorder. <i>Journal of Affective Disorders</i> , 2013, 145, 136-141.	2.0	39
93	Modulation of amygdala response and connectivity in depression by serotonin transporter polymorphism and diagnosis. <i>Journal of Affective Disorders</i> , 2013, 150, 96-103.	2.0	70
94	A mega-analysis of genome-wide association studies for major depressive disorder. <i>Molecular Psychiatry</i> , 2013, 18, 497-511.	4.1	1,002
95	A genome-wide association study of a sustained pattern of antidepressant response. <i>Journal of Psychiatric Research</i> , 2013, 47, 1157-1165.	1.5	52
96	Association at SYNE1 in both bipolar disorder and recurrent major depression. <i>Molecular Psychiatry</i> , 2013, 18, 614-617.	4.1	80
97	Schizophrenia as a Human Leukocyte Antigen-Associated Disease Revisited. <i>American Journal of Psychiatry</i> , 2013, 170, 821-823.	4.0	8
98	Genome-wide association study of intraocular pressure identifies the GLCC1/ICA1 region as a glaucoma susceptibility locus. <i>Human Molecular Genetics</i> , 2013, 22, 4653-4660.	1.4	29
99	Genome-wide association study of co-occurring anxiety in major depression. <i>World Journal of Biological Psychiatry</i> , 2013, 14, 611-621.	1.3	17
100	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
101	The Sri Lankan Twin Registry: 2012 Update. <i>Twin Research and Human Genetics</i> , 2013, 16, 307-312.	0.3	14
102	Integrative mouse and human mRNA studies using WGCNA nominates novel candidate genes involved in the pathogenesis of major depressive disorder. <i>Pharmacogenomics</i> , 2013, 14, 1979-1990.	0.6	55
103	Whole-exome sequencing identifies a polymorphism in the BMP5 gene associated with SSRI treatment response in major depression. <i>Journal of Psychopharmacology</i> , 2013, 27, 915-920.	2.0	31
104	Antidepressant effects of nortriptyline and escitalopram in the GENDEP study: Is one better than the other?. <i>Acta Psychiatrica Scandinavica</i> , 2013, 127, 330-330.	2.2	1
105	ATP-binding cassette sub-family F member 1 (ABCF1) is identified as a putative therapeutic target of escitalopram in the inflammatory cytokine pathway. <i>Journal of Psychopharmacology</i> , 2013, 27, 609-615.	2.0	20
106	Candidate Genes Expression Profile Associated with Antidepressants Response in the GENDEP Study: Differentiating between Baseline "Predictors"™ and Longitudinal "Targets"™. <i>Neuropsychopharmacology</i> , 2013, 38, 377-385.	2.8	372
107	The current state of play on the molecular genetics of depression. <i>Psychological Medicine</i> , 2013, 43, 673-687.	2.7	73
108	Estimating the heritability of reporting stressful life events captured by common genetic variants. <i>Psychological Medicine</i> , 2013, 43, 1965-1971.	2.7	46



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109	Harnessing Clinical Psychiatric Data with an Electronic Assessment Tool (OPCRIT+): The Utility of Symptom Dimensions. <i>PLoS ONE</i> , 2013, 8, e58790.	1.1	10
110	Male-Biased Autosomal Effect of 16p13.11 Copy Number Variation in Neurodevelopmental Disorders. <i>PLoS ONE</i> , 2013, 8, e61365.	1.1	101
111	Common Genetic Determinants of Intraocular Pressure and Primary Open-Angle Glaucoma. <i>PLoS Genetics</i> , 2012, 8, e1002611.	1.5	164
112	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
113	<i>CYP2C19</i> genotype predicts steady state escitalopram concentration in GENDEP. <i>Journal of Psychopharmacology</i> , 2012, 26, 398-407.	2.0	69
114	Allele-specific Differences in Activity of a Novel Cannabinoid Receptor 1 (CNR1) Gene Intronic Enhancer in Hypothalamus, Dorsal Root Ganglia, and Hippocampus. <i>Journal of Biological Chemistry</i> , 2012, 287, 12828-12834.	1.6	14
115	The Genetic Basis of Depression. <i>Current Topics in Behavioral Neurosciences</i> , 2012, 14, 81-99.	0.8	8
116	Non-steroidal anti-inflammatory drugs and efficacy of antidepressants in major depressive disorder. <i>Psychological Medicine</i> , 2012, 42, 2027-2035.	2.7	30
117	Depression symptom dimensions as predictors of antidepressant treatment outcome: replicable evidence for interest-activity symptoms. <i>Psychological Medicine</i> , 2012, 42, 967-980.	2.7	298
118	Genomic structural variation in psychiatric disorders. <i>Development and Psychopathology</i> , 2012, 24, 1335-1344.	1.4	14
119	Life-event specificity: bipolar disorder compared with unipolar depression. <i>British Journal of Psychiatry</i> , 2012, 201, 458-465.	1.7	32
120	Genetic and Environmental Etiology of Nicotine Use in Sri Lankan Male Twins. <i>Behavior Genetics</i> , 2012, 42, 798-807.	1.4	6
121	Non-random dropout and the relative efficacy of escitalopram and nortriptyline in treating major depressive disorder. <i>Journal of Psychiatric Research</i> , 2012, 46, 1333-1338.	1.5	12
122	The role of loss and danger events in symptom exacerbation in bipolar disorder. <i>Journal of Psychiatric Research</i> , 2012, 46, 1584-1589.	1.5	32
123	Genome-wide approaches to antidepressant treatment: working towards understanding and predicting response. <i>Genome Medicine</i> , 2012, 4, 52.	3.6	16
124	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
125	Dissecting the Genetic Heterogeneity of Depression Through Age at Onset. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2012, 159B, 859-868.	1.1	31
126	SELF-REPORT AND CLINICIAN-RATED MEASURES OF DEPRESSION SEVERITY: CAN ONE REPLACE THE OTHER?. <i>Depression and Anxiety</i> , 2012, 29, 1043-1049.	2.0	182



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127	Pharmacoproteomic investigation into antidepressant response in two mouse inbred strains. <i>Proteomics</i> , 2012, 12, 2355-2365.	1.3	18
128	A Polymorphism Associated with Depressive Disorders Differentially Regulates Brain Derived Neurotrophic Factor Promoter IV Activity. <i>Biological Psychiatry</i> , 2012, 71, 618-626.	0.7	51
129	Common Genetic Variants and Gene-Expression Changes Associated with Bipolar Disorder Are Over-Represented in Brain Signaling Pathway Genes. <i>Biological Psychiatry</i> , 2012, 72, 311-317.	0.7	56
130	Replication Study and Meta-Analysis in European Samples Supports Association of the 3p21.1 Locus with Bipolar Disorder. <i>Biological Psychiatry</i> , 2012, 72, 645-650.	0.7	15
131	Meta-analyses of genome-wide linkage scans of anxiety-related phenotypes. <i>European Journal of Human Genetics</i> , 2012, 20, 1078-1084.	1.4	28
132	White matter abnormalities and illness severity in major depressive disorder. <i>British Journal of Psychiatry</i> , 2012, 201, 33-39.	1.7	126
133	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
134	Depressive disorder moderates the effect of the FTO gene on body mass index. <i>Molecular Psychiatry</i> , 2012, 17, 604-611.	4.1	72
135	Stressful life events and the serotonin transporter gene ( 5-HTT ) in recurrent clinical depression. <i>Journal of Affective Disorders</i> , 2012, 136, 189-193.	2.0	22
136	A twin study of schizoaffective mania, schizoaffective depression, and other psychotic syndromes. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2012, 159B, 172-182.	1.1	32
137	Antidepressant-dependent mRNA changes in mouse associated with hippocampal neurogenesis in a mouse model of depression. <i>Pharmacogenetics and Genomics</i> , 2012, 22, 765-776.	0.7	28
138	Convergent Animal and Human Evidence Suggests a Role of PPM1A Gene in Response to Antidepressants. <i>Biological Psychiatry</i> , 2011, 69, 360-365.	0.7	30
139	No association between the Catechol-O-Methyltransferase (COMT) val158met polymorphism and cognitive improvement following cognitive remediation therapy (CRT) in schizophrenia. <i>Neuroscience Letters</i> , 2011, 496, 65-69.	1.0	40
140	Admixture analysis of age at onset in bipolar disorder. <i>Psychiatry Research</i> , 2011, 185, 27-32.	1.7	51
141	Genomewide Association Scan of Suicidal Thoughts and Behaviour in Major Depression. <i>PLoS ONE</i> , 2011, 6, e20690.	1.1	98
142	Childhood adversity and psychosis. <i>International Clinical Psychopharmacology</i> , 2011, 26, e96.	0.9	0
143	OPCRIT+: an electronic system for psychiatric diagnosis and data collection in clinical and research settings. <i>British Journal of Psychiatry</i> , 2011, 199, 151-155.	1.7	48
144	No effect of 5HTTLPR or BDNF Val66Met polymorphism on hippocampal morphology in major depression. <i>Genes, Brain and Behavior</i> , 2011, 10, 756-764.	1.1	78

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145	Replication of association of 3p21.1 with susceptibility to bipolar disorder but not major depression. <i>Nature Genetics</i> , 2011, 43, 3-5.	9.4	29
146	Meta-analysis of genome-wide association data of bipolar disorder and major depressive disorder. <i>Molecular Psychiatry</i> , 2011, 16, 2-4.	4.1	150
147	Melancholic, atypical and anxious depression subtypes and outcome of treatment with escitalopram and nortriptyline. <i>Journal of Affective Disorders</i> , 2011, 132, 112-120.	2.0	93
148	Hippocampal atrophy in first episode depression: A meta-analysis of magnetic resonance imaging studies. <i>Journal of Affective Disorders</i> , 2011, 134, 483-487.	2.0	262
149	Genome-Wide Searches for Bipolar Disorder Genes. <i>Current Psychiatry Reports</i> , 2011, 13, 522-527.	2.1	12
150	Methylenetetrahydrofolate Reductase Gene Variant (MTHFR C677T) and Migraine: A Case Control Study and Meta-analysis. <i>BMC Neurology</i> , 2011, 11, 66.	0.8	45
151	Heritability estimates for psychotic symptom dimensions in twins with psychotic disorders. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2011, 156, 89-98.	1.1	26
152	Response to the letter from Dr. Maher and colleagues Re. Linkage on suicidality. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2011, 156, 864-865.	1.1	0
153	Phenotype evaluation and genomewide linkage study of clinical variables in schizophrenia. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2011, 156, 929-940.	1.1	14
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