

# Francis J McMahon

## List of Publications by Year in descending order

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254  
papers

27,956  
citations

7551

77  
h-index

7136

153  
g-index

285  
all docs

285  
docs citations

285  
times ranked

26587  
citing authors

#	ARTICLE	IF	CITATIONS
1	Using polygenic scores and clinical data for bipolar disorder patient stratification and lithium response prediction: machine learning approach. <i>British Journal of Psychiatry</i> , 2022, 220, 219-228.	1.7	11
2	Putting Genetics to Work in the Psychiatric Clinic. <i>American Journal of Psychiatry</i> , 2022, 179, 182-188.	4.0	2
3	Rare variants implicate NMDA receptor signaling and cerebellar gene networks in risk for bipolar disorder. <i>Molecular Psychiatry</i> , 2022, 27, 3842-3856.	4.1	5
4	Genome-wide association study of panic disorder reveals genetic overlap with neuroticism and depression. <i>Molecular Psychiatry</i> , 2021, 26, 4179-4190.	4.1	58
5	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
6	Gene-Based Association Testing of Dichotomous Traits With Generalized Functional Linear Mixed Models Using Extended Pedigrees: Applications to Age-Related Macular Degeneration. <i>Journal of the American Statistical Association</i> , 2021, 116, 531-545.	1.8	3
7	Exemplar scoring identifies genetically separable phenotypes of lithium responsive bipolar disorder. <i>Translational Psychiatry</i> , 2021, 11, 36.	2.4	16
8	Prediction of lithium response using genomic data. <i>Scientific Reports</i> , 2021, 11, 1155.	1.6	11
9	Deep transcriptome sequencing of subgenual anterior cingulate cortex reveals cross-diagnostic and diagnosis-specific RNA expression changes in major psychiatric disorders. <i>Neuropsychopharmacology</i> , 2021, 46, 1364-1372.	2.8	22
10	Genetic versus stress and mood determinants of sleep in the Amish. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2021, 186, 113-121.	1.1	2
11	Multiple dimensions of stress vs. genetic effects on depression. <i>Translational Psychiatry</i> , 2021, 11, 254.	2.4	4
12	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
13	Characterisation of age and polarity at onset in bipolar disorder. <i>British Journal of Psychiatry</i> , 2021, 219, 659-669.	1.7	20
14	Genetic Overlap Profiles of Cognitive Ability in Psychotic and Affective Illnesses: A Multisite Study of Multiplex Pedigrees. <i>Biological Psychiatry</i> , 2021, 90, 373-384.	0.7	5
15	HLA-DRB1 and HLA-DQB1 genetic diversity modulates response to lithium in bipolar affective disorders. <i>Scientific Reports</i> , 2021, 11, 17823.	1.6	10
16	Review and Consensus on Pharmacogenomic Testing in Psychiatry. <i>Pharmacopsychiatry</i> , 2021, 54, 5-17.	1.7	96
17	Combining schizophrenia and depression polygenic risk scores improves the genetic prediction of lithium response in bipolar disorder patients. <i>Translational Psychiatry</i> , 2021, 11, 606.	2.4	25
18	The genetics of bipolar disorder. <i>Molecular Psychiatry</i> , 2020, 25, 544-559.	4.1	161

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19	Polygenic risk for anxiety influences anxiety comorbidity and suicidal behavior in bipolar disorder. <i>Translational Psychiatry</i> , 2020, 10, 298.	2.4	16
20	Validity of the Mood Disorder Questionnaire (MDQ) as a screening tool for bipolar spectrum disorders in anabaptist populations. <i>Journal of Psychiatric Research</i> , 2020, 123, 159-163.	1.5	5
21	Linear mixed models for association analysis of quantitative traits with next-generation sequencing data. <i>Genetic Epidemiology</i> , 2019, 43, 189-206.	0.6	5
22	Investigating polygenic burden in age at disease onset in bipolar disorder: Findings from an international multicentric study. <i>Bipolar Disorders</i> , 2019, 21, 68-75.	1.1	20
23	Rediscovering the value of families for psychiatric genetics research. <i>Molecular Psychiatry</i> , 2019, 24, 523-535.	4.1	43
24	Clinical and genetic validity of quantitative bipolarity. <i>Translational Psychiatry</i> , 2019, 9, 228.	2.4	4
25	From genetics to biology: advancing mental health research in the Genomics ERA. <i>Molecular Psychiatry</i> , 2019, 24, 1576-1582.	4.1	7
26	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019, 51, 793-803.	9.4	1,191
27	Sodium valproate rescues expression of TRANK1 in iPSC-derived neural cells that carry a genetic variant associated with serious mental illness. <i>Molecular Psychiatry</i> , 2019, 24, 613-624.	4.1	34
28	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , 2019, 179, 1469-1482.e11.	13.5	935
29	Efficient region-based test strategy uncovers genetic risk factors for functional outcome in bipolar disorder. <i>European Neuropsychopharmacology</i> , 2019, 29, 156-170.	0.3	7
30	Association of Polygenic Score for Schizophrenia and HLA Antigen and Inflammation Genes With Response to Lithium in Bipolar Affective Disorder. <i>JAMA Psychiatry</i> , 2018, 75, 65-74.	6.0	102
31	Exome sequencing of a large family identifies potential candidate genes contributing risk to bipolar disorder. <i>Gene</i> , 2018, 645, 119-123.	1.0	29
32	Coherence Through Incongruence—Can Genetic Markers Inform Nosology After All?. <i>JAMA Psychiatry</i> , 2018, 75, 7.	6.0	0
33	Detecting significant genotype-phenotype association rules in bipolar disorder: market research meets complex genetics. <i>International Journal of Bipolar Disorders</i> , 2018, 6, 24.	0.8	8
34	Exploratory genome-wide association analysis of response to ketamine and a polygenic analysis of response to scopolamine in depression. <i>Translational Psychiatry</i> , 2018, 8, 280.	2.4	26
35	Population-Based Estimates of Heritability Shed New Light on Clinical Features of Major Depression. <i>American Journal of Psychiatry</i> , 2018, 175, 1058-1060.	4.0	4
36	Genetic pleiotropy between mood disorders, metabolic, and endocrine traits in a multigenerational pedigree. <i>Translational Psychiatry</i> , 2018, 8, 218.	2.4	17

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37	Convergent analysis of genome-wide genotyping and transcriptomic data suggests association of zinc finger genes with lithium response in bipolar disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2018, 177, 658-664.	1.1	10
38	Analysis of the Influence of microRNAs in Lithium Response in Bipolar Disorder. Frontiers in Psychiatry, 2018, 9, 207.	1.3	28
39	Evaluation of Recipients of Positive and Negative Secondary Findings Evaluations in a Hybrid CLIA-Research Sequencing Pilot. American Journal of Human Genetics, 2018, 103, 358-366.	2.6	29
40	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. Cell, 2018, 173, 1705-1715.e16.	13.5	623
41	Novel genetic loci associated with hippocampal volume. Nature Communications, 2017, 8, 13624.	5.8	250
42	200. Effects of Medications on RNA-Seq Gene Expression from Anterior Cingulate Cortex in Adult Major Psychiatric Disorders. Biological Psychiatry, 2017, 81, S83.	0.7	1
43	The antidepressant efficacy of subanesthetic-dose ketamine does not correlate with baseline subcortical volumes in a replication sample with major depressive disorder. Journal of Psychopharmacology, 2017, 31, 1570-1577.	2.0	17
44	691. RNA-sequencing of the Subgenual Anterior Cingulate Cortex in Adult Major Psychiatric Disorders. Biological Psychiatry, 2017, 81, S280.	0.7	0
45	A population-specific reference panel empowers genetic studies of Anabaptist populations. Scientific Reports, 2017, 7, 6079.	1.6	16
46	506. Cyclic Expression of Circadian Genes in Neural Progenitor Cells Derived from Human Induced Pluripotent Stem Cells. Biological Psychiatry, 2017, 81, S205-S206.	0.7	0
47	Circadian genes and lithium response in bipolar disorders: associations with <i>PPARGC1A</i> ( <i>PGC1<math>\alpha</math></i> ) and <i>RORA</i> . Genes, Brain and Behavior, 2016, 15, 660-668.	1.1	37
48	Novel genetic loci underlying human intracranial volume identified through genome-wide association. Nature Neuroscience, 2016, 19, 1569-1582.	7.1	213
49	Neurocognitive functioning in euthymic patients with bipolar disorder and unaffected relatives: A review of the literature. Neuroscience and Biobehavioral Reviews, 2016, 69, 193-215.	2.9	59
50	Constance E. Lieber, Theodore R. Stanley, and the Enduring Impact of Philanthropy on Psychiatry Research. Biological Psychiatry, 2016, 80, 84-86.	0.7	2
51	Symptom profiles and illness course among Anabaptist and Non-Anabaptist adults with major mood disorders. International Journal of Bipolar Disorders, 2016, 4, 21.	0.8	6
52	Genome-wide association study of 40,000 individuals identifies two novel loci associated with bipolar disorder. Human Molecular Genetics, 2016, 25, 3383-3394.	1.4	182
53	Genetic variants associated with response to lithium treatment in bipolar disorder: a genome-wide association study. Lancet, The, 2016, 387, 1085-1093.	6.3	306
54	Genetic influences on schizophrenia and subcortical brain volumes: large-scale proof of concept. Nature Neuroscience, 2016, 19, 420-431.	7.1	204

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55	Genetic association studies in psychiatry: time for pay-off. <i>Lancet Psychiatry</i> , 2016, 3, 309-310.	3.7	2
56	Risk factors for suicide in bipolar I disorder in two prospectively studied cohorts. <i>Journal of Affective Disorders</i> , 2016, 190, 1-5.	2.0	17
57	A genome-wide association study of bipolar disorder with comorbid eating disorder replicates the SOX2-OT region. <i>Journal of Affective Disorders</i> , 2016, 189, 141-149.	2.0	45
58	An Integrative Genomic Study Implicates the Postsynaptic Density in the Pathogenesis of Bipolar Disorder. <i>Neuropsychopharmacology</i> , 2016, 41, 886-895.	2.8	28
59	Finding Rare, Disease-Associated Variants in Isolated Groups: Potential Advantages of Mennonite Populations. <i>Human Biology</i> , 2016, 88, 109.	0.4	25
60	The Genetic Basis of Bipolar Disorder. <i>Milestones in Drug Therapy</i> , 2016, , 73-92.	0.1	0
61	A Novel Mixture Model to Estimate the Time to Drug Effect Onset and Its Association with Covariates. <i>Human Heredity</i> , 2015, 80, 90-99.	0.4	0
62	Test-retest reliability of a new questionnaire for the retrospective assessment of long-term lithium use in bipolar disorder. <i>Journal of Affective Disorders</i> , 2015, 174, 589-593.	2.0	8
63	Genome wide association study identifies variants in NBEA associated with migraine in bipolar disorder. <i>Journal of Affective Disorders</i> , 2015, 172, 453-461.	2.0	15
64	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
65	Psychiatric genome-wide association study analyses implicate neuronal, immune and histone pathways. <i>Nature Neuroscience</i> , 2015, 18, 199-209.	7.1	701
66	Common genetic variants influence human subcortical brain structures. <i>Nature</i> , 2015, 520, 224-229.	13.7	772
67	Rare variants in neuronal excitability genes influence risk for bipolar disorder. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 3576-3581.	3.3	152
68	Clinically Useful Genetic Markers of Antidepressant Response: How Do We Get There From Here?. <i>American Journal of Psychiatry</i> , 2015, 172, 697-699.	4.0	7
69	Variant <i>GADL1</i> and Response to Lithium in Bipolar I Disorder. <i>New England Journal of Medicine</i> , 2014, 370, 1855-1860.	13.9	36
70	Molecular genetic overlap in bipolar disorder, schizophrenia, and major depressive disorder. <i>World Journal of Biological Psychiatry</i> , 2014, 15, 200-208.	1.3	120
71	Identification of Pathways for Bipolar Disorder. <i>JAMA Psychiatry</i> , 2014, 71, 657.	6.0	204
72	The ENIGMA Consortium: large-scale collaborative analyses of neuroimaging and genetic data. <i>Brain Imaging and Behavior</i> , 2014, 8, 153-182.	1.1	696

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73	RNA-sequencing of the brain transcriptome implicates dysregulation of neuroplasticity, circadian rhythms and GTPase binding in bipolar disorder. <i>Molecular Psychiatry</i> , 2014, 19, 1179-1185.	4.1	100
74	The genetic interacting landscape of 63 candidate genes in Major Depressive Disorder: an explorative study. <i>BioData Mining</i> , 2014, 7, 19.	2.2	7
75	Corpus Callosum Size Is Highly Heritable in Humans, and May Reflect Distinct Genetic Influences on Ventral and Rostral Regions. <i>PLoS ONE</i> , 2014, 9, e99980.	1.1	14
76	Do Participants in Genome Sequencing Studies of Psychiatric Disorders Wish to Be Informed of Their Results? A Survey Study. <i>PLoS ONE</i> , 2014, 9, e101111.	1.1	30
77	Common and Rare Variant Analysis in Early-Onset Bipolar Disorder Vulnerability. <i>PLoS ONE</i> , 2014, 9, e104326.	1.1	34
78	Prediction of treatment outcomes in psychiatry—where do we stand ?. <i>Dialogues in Clinical Neuroscience</i> , 2014, 16, 455-464.	1.8	47
79	Association study of 83 candidate genes for bipolar disorder in chromosome 6q selected using an evidence-based prioritization algorithm. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2013, 162, 898-906.	1.1	9
80	Race, Genetic Ancestry and Response to Antidepressant Treatment for Major Depression. <i>Neuropsychopharmacology</i> , 2013, 38, 2598-2606.	2.8	39
81	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. <i>Nature Genetics</i> , 2013, 45, 984-994.	9.4	2,067
82	Two gene co-expression modules differentiate psychotics and controls. <i>Molecular Psychiatry</i> , 2013, 18, 1308-1314.	4.1	160
83	RETENTION AND ATTRITION AMONG AFRICAN AMERICANS IN THE STAR*D STUDY: WHAT CAUSES RESEARCH VOLUNTEERS TO STAY OR STRAY?. <i>Depression and Anxiety</i> , 2013, 30, 1137-1144.	2.0	17
84	Amish revisited: next-generation sequencing studies of psychiatric disorders among the Plain people. <i>Trends in Genetics</i> , 2013, 29, 412-418.	2.9	24
85	Enrichment of cis-regulatory gene expression SNPs and methylation quantitative trait loci among bipolar disorder susceptibility variants. <i>Molecular Psychiatry</i> , 2013, 18, 340-346.	4.1	153
86	Genome-wide association study meta-analysis of European and Asian-ancestry samples identifies three novel loci associated with bipolar disorder. <i>Molecular Psychiatry</i> , 2013, 18, 195-205.	4.1	180
87	In vivo radioligand binding to translocator protein correlates with severity of Alzheimer's disease. <i>Brain</i> , 2013, 136, 2228-2238.	3.7	280
88	A Genetic Polymorphism for Translocator Protein 18 Kda Affects both <i>in Vitro</i> and <i>in Vivo</i> Radioligand Binding in Human Brain to this Putative Biomarker of Neuroinflammation. <i>Journal of Cerebral Blood Flow and Metabolism</i> , 2013, 33, 53-58.	2.4	207
89	Cancer Regression and Neurological Toxicity Following Anti-MAGE-A3 TCR Gene Therapy. <i>Journal of Immunotherapy</i> , 2013, 36, 133-151.	1.2	953
90	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

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91	Common and rare alleles of the serotonin transporter gene, <i>SLC6A4</i> , associated with Tourette's disorder. <i>Movement Disorders</i> , 2013, 28, 1263-1270.	2.2	44
92	Genome-Wide Association Study of Irritable vs. Elated Mania Suggests Genetic Differences between Clinical Subtypes of Bipolar Disorder. <i>PLoS ONE</i> , 2013, 8, e53804.	1.1	22
93	Assessment of Response to Lithium Maintenance Treatment in Bipolar Disorder: A Consortium on Lithium Genetics (ConLiGen) Report. <i>PLoS ONE</i> , 2013, 8, e65636.	1.1	156
94	Pharmacogenetics of antidepressants, mood stabilizers, and antipsychotics in diverse human populations. <i>Discovery Medicine</i> , 2013, 16, 113-22.	0.5	19
95	Genome-wide significant association between a "negative mood delusions" dimension in bipolar disorder and genetic variation on chromosome 3q26.1. <i>Translational Psychiatry</i> , 2012, 2, e165-e165.	2.4	14
96	Common genetic variation in the indoleamine-2,3-dioxygenase genes and antidepressant treatment outcome in major depressive disorder. <i>Journal of Psychopharmacology</i> , 2012, 26, 360-367.	2.0	36
97	Replication and meta-analysis of TMEM132D gene variants in panic disorder. <i>Translational Psychiatry</i> , 2012, 2, e156-e156.	2.4	74
98	Genome-wide linkage analysis of 972 bipolar pedigrees using single-nucleotide polymorphisms. <i>Molecular Psychiatry</i> , 2012, 17, 818-826.	4.1	31
99	Evidence for association of bipolar disorder to haplotypes in the 22q12.3 region near the genes stargazin, <i>IFT27</i> and parvalbumin. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2012, 159B, 941-950.	1.1	10
100	Identification of common variants associated with human hippocampal and intracranial volumes. <i>Nature Genetics</i> , 2012, 44, 552-561.	9.4	594
101	Pharmacogenomics and Personalized Medicine in Neuropsychiatry. <i>Neuron</i> , 2012, 74, 773-776.	3.8	64
102	Interaction networks of lithium and valproate molecular targets reveal a striking enrichment of apoptosis functional clusters and neurotrophin signaling. <i>Pharmacogenomics Journal</i> , 2012, 12, 328-341.	0.9	36
103	Brain-Derived Neurotrophic Factor Val66Met Polymorphism and Antidepressant Efficacy of Ketamine in Depressed Patients. <i>Biological Psychiatry</i> , 2012, 72, e27-e28.	0.7	187
104	A genome-wide association study of attempted suicide. <i>Molecular Psychiatry</i> , 2012, 17, 433-444.	4.1	141
105	Association study of serotonin pathway genes in attempted suicide. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2012, 159B, 112-119.	1.1	15
106	Gene Expression and Genetic Variation Data Implicate PCLO in Bipolar Disorder. <i>Biological Psychiatry</i> , 2011, 69, 353-359.	0.7	53
107	The Bcl-2 Gene Polymorphism rs956572AA Increases Inositol 1,4,5-Trisphosphate Receptor-Mediated Endoplasmic Reticulum Calcium Release in Subjects with Bipolar Disorder. <i>Biological Psychiatry</i> , 2011, 69, 344-352.	0.7	65
108	The genetics of panic disorder. <i>Journal of Medical Genetics</i> , 2011, 48, 361-368.	1.5	46



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109	Genome-wide association studies of antidepressant outcome: A brief review. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2011, 35, 1553-1557.	2.5	41
110	High Frequencies of De Novo CNVs in Bipolar Disorder and Schizophrenia. <i>Neuron</i> , 2011, 72, 951-963.	3.8	290
111	A Network-Based Approach to Prioritize Results from Genome-Wide Association Studies. <i>PLoS ONE</i> , 2011, 6, e24220.	1.1	64
112	Genetic association of bipolar disorder with the $\alpha 3$ nicotinic receptor subunit gene. <i>Psychiatric Genetics</i> , 2011, 21, 77-84.	0.6	10
113	Reply to "Replication of association of 3p21.1 with susceptibility to bipolar disorder but not major depression": <i>Nature Genetics</i> , 2011, 43, 5-5.	9.4	13
114	Genetic variation in cholinergic muscarinic-2 receptor gene modulates M2 receptor binding in vivo and accounts for reduced binding in bipolar disorder. <i>Molecular Psychiatry</i> , 2011, 16, 407-418.	4.1	52
115	A functional alternative splicing mutation in human tryptophan hydroxylase-2. <i>Molecular Psychiatry</i> , 2011, 16, 1169-1176.	4.1	21
116	A non-synonymous polymorphism in galactose mutarotase (GALM) is associated with serotonin transporter binding potential in the human thalamus: results of a genome-wide association study. <i>Molecular Psychiatry</i> , 2011, 16, 584-585.	4.1	19
117	Identity-by-descent filtering as a tool for the identification of disease alleles in exome sequence data from distant relatives. <i>BMC Proceedings</i> , 2011, 5, S76.	1.8	6
118	Genome-wide association analysis of age at onset and psychotic symptoms in bipolar disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2011, 156, 370-378.	1.1	42
119	Large-scale genome-wide association analysis of bipolar disorder identifies a new susceptibility locus near ODZ4. <i>Nature Genetics</i> , 2011, 43, 977-983.	9.4	1,283
120	Genome-Wide Association of Bipolar Disorder Suggests an Enrichment of Replicable Associations in Regions near Genes. <i>PLoS Genetics</i> , 2011, 7, e1002134.	1.5	59
121	Genetic variation in HTR2A influences serotonin transporter binding potential as measured using PET and [ <sup>11</sup> C]DASB. <i>International Journal of Neuropsychopharmacology</i> , 2010, 13, 715-724.	1.0	35
122	Increased gene expression of diacylglycerol kinase eta in bipolar disorder. <i>International Journal of Neuropsychopharmacology</i> , 2010, 13, 1127-1128.	1.0	38
123	A Genome-Wide Association Study of Amygdala Activation in Youths With and Without Bipolar Disorder. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2010, 49, 33-41.	0.3	10
124	Sex-specific association of the reelin gene with bipolar disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2010, 153B, 549-553.	1.1	55
125	Case-control association study of <i>TGOLN2</i> in attempted suicide. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2010, 153B, 1016-1023.	1.1	4
126	Coming to grips with complex disorders: Genetic risk prediction in bipolar disorder using panels of genes identified through convergent functional genomics. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2010, 153B, 850-877.	1.1	50



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127	An investigation of candidate regions for association with bipolar disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 1292-1297.	1.1	6
128	Meta-analysis of genome-wide association data identifies a risk locus for major mood disorders on 3p21.1. Nature Genetics, 2010, 42, 128-131.	9.4	152
129	The International Consortium on Lithium Genetics (ConLiGen): An Initiative by the NIMH and IGSLI to Study the Genetic Basis of Response to Lithium Treatment. Neuropsychobiology, 2010, 62, 72-78.	0.9	134
130	Brain-derived neurotrophic factor ( BDNF) gene: no major impact on antidepressant treatment response. International Journal of Neuropsychopharmacology, 2010, 13, 93.	1.0	104
131	Genome-Wide Association Study of Suicide Attempts in Mood Disorder Patients. American Journal of Psychiatry, 2010, 167, 1499-1507.	4.0	140
132	Genetics of Bipolar Disorder. Current Topics in Behavioral Neurosciences, 2010, 5, 19-30.	0.8	2
133	Pioneering First Steps and Cautious Conclusions. Biological Psychiatry, 2010, 67, 99-100.	0.7	5
134	A Genome-Wide Association Study of Amygdala Activation in Youths With and Without Bipolar Disorder. Journal of the American Academy of Child and Adolescent Psychiatry, 2010, 49, 33-41.	0.3	30
135	Genome-Wide Linkage and Follow-Up Association Study of Postpartum Mood Symptoms. American Journal of Psychiatry, 2009, 166, 1229-1237.	4.0	85
136	Pharmacogenetics Studies in STAR*D: Strengths, Limitations, and Results. Psychiatric Services, 2009, 60, 1446-1457.	1.1	69
137	Genetic and Clinical Predictors of Sexual Dysfunction in Citalopram-Treated Depressed Patients. Neuropsychopharmacology, 2009, 34, 1819-1828.	2.8	88
138	The DISC locus and schizophrenia: evidence from an association study in a central European sample and from a meta-analysis across different European populations. Human Molecular Genetics, 2009, 18, 2719-2727.	1.4	78
139	Convergent functional genomics of genome-wide association data for bipolar disorder: Comprehensive identification of candidate genes, pathways and mechanisms. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2009, 150B, 155-181.	1.1	178
140	Family-based association study of Neuregulin 1 with psychotic bipolar disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2009, 150B, 693-702.	1.1	31
141	Convergent genome wide association results for bipolar disorder and substance dependence. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2009, 150B, 182-190.	1.1	45
142	Common and rare variants of <i>DAOA</i> in bipolar disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2009, 150B, 960-966.	1.1	11
143	Family-based association of <i>YWHAH</i> in psychotic bipolar disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2009, 150B, 977-983.	1.1	49
144	Premenstrual mood symptoms: study of familiarity and personality correlates in mood disorder pedigrees. Archives of Women's Mental Health, 2009, 12, 27-34.	1.2	14

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145	Family-based association of FKBP5 in bipolar disorder. <i>Molecular Psychiatry</i> , 2009, 14, 261-268.	4.1	140
146	Two variants in Ankyrin 3 (ANK3) are independent genetic risk factors for bipolar disorder. <i>Molecular Psychiatry</i> , 2009, 14, 487-491.	4.1	171
147	Singleton deletions throughout the genome increase risk of bipolar disorder. <i>Molecular Psychiatry</i> , 2009, 14, 376-380.	4.1	137
148	Genome-wide association study of bipolar disorder in European American and African American individuals. <i>Molecular Psychiatry</i> , 2009, 14, 755-763.	4.1	326
149	Microduplications of 16p11.2 are associated with schizophrenia. <i>Nature Genetics</i> , 2009, 41, 1223-1227.	9.4	646
150	Bcl-2 Polymorphism Influences Gray Matter Volume in the Ventral Striatum in Healthy Humans. <i>Biological Psychiatry</i> , 2009, 66, 804-807.	0.7	24
151	SSRI response in depression may be influenced by SNPs in HTR1B and HTR1A. <i>Psychiatric Genetics</i> , 2009, 19, 281-291.	0.6	62
152	Genome-wide association study of suicidal ideation emerging during citalopram treatment of depressed outpatients. <i>Pharmacogenetics and Genomics</i> , 2009, 19, 666-674.	0.7	103
153	The genetic basis of bipolar disorder. , 2009, , 59-76.		2
154	Association study of phosphodiesterase genes in the Sequenced Treatment Alternatives to Relieve Depression sample. <i>Pharmacogenetics and Genomics</i> , 2009, 19, 235-238.	0.7	29
155	Pharmacogenetics Studies in STAR*D: Strengths, Limitations, and Results. <i>Psychiatric Services</i> , 2009, 60, 1446-57.	1.1	50
156	Gene-based SNP mapping of a psychotic bipolar affective disorder linkage region on 22q12.3: Association with <i>HMG2L1</i> and <i>TOM1</i> . <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008, 147B, 59-67.	1.1	16
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