

Francis J McMahon

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

249
papers

21,539
citations

75
h-index

143
g-index

282
ext. papers

25,287
ext. citations

9.3
avg, IF

6.72
L-index

#	Paper	IF	Citations
249	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. <i>Nature Genetics</i> , 2013 , 45, 984-94	36.3	1628
248	Large-scale genome-wide association analysis of bipolar disorder identifies a new susceptibility locus near ODZ4. <i>Nature Genetics</i> , 2011 , 43, 977-83	36.3	1094
247	Cancer regression and neurological toxicity following anti-MAGE-A3 TCR gene therapy. <i>Journal of Immunotherapy</i> , 2013 , 36, 133-51	5	758
246	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019 , 51, 793-803	36.3	662
245	Common genetic variants influence human subcortical brain structures. <i>Nature</i> , 2015 , 520, 224-9	50.4	601
244	Psychiatric genome-wide association study analyses implicate neuronal, immune and histone pathways. <i>Nature Neuroscience</i> , 2015 , 18, 199-209	25.5	572
243	Microduplications of 16p11.2 are associated with schizophrenia. <i>Nature Genetics</i> , 2009 , 41, 1223-7	36.3	550
242	A genome-wide association study implicates diacylglycerol kinase eta (DGKH) and several other genes in the etiology of bipolar disorder. <i>Molecular Psychiatry</i> , 2008 , 13, 197-207	15.1	548
241	The ENIGMA Consortium: large-scale collaborative analyses of neuroimaging and genetic data. <i>Brain Imaging and Behavior</i> , 2014 , 8, 153-82	4.1	539
240	Identification of common variants associated with human hippocampal and intracranial volumes. <i>Nature Genetics</i> , 2012 , 44, 552-61	36.3	498
239	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , 2019 , 179, 1469-1482.e11	56.2	402
238	Variation in the gene encoding the serotonin 2A receptor is associated with outcome of antidepressant treatment. <i>American Journal of Human Genetics</i> , 2006 , 78, 804-814	11	396
237	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. <i>Cell</i> , 2018 , 173, 1705-1715.e16	56.2	360
236	Genome scan meta-analysis of schizophrenia and bipolar disorder, part III: Bipolar disorder. <i>American Journal of Human Genetics</i> , 2003 , 73, 49-62	11	353
235	Genome-wide association study of bipolar disorder in European American and African American individuals. <i>Molecular Psychiatry</i> , 2009 , 14, 755-63	15.1	287
234	High frequencies of de novo CNVs in bipolar disorder and schizophrenia. <i>Neuron</i> , 2011 , 72, 951-63	13.9	240
233	In vivo radioligand binding to translocator protein correlates with severity of Alzheimer's disease. <i>Brain</i> , 2013 , 136, 2228-38	11.2	232

232	G72/G30 in schizophrenia and bipolar disorder: review and meta-analysis. <i>Biological Psychiatry</i> , 2006 , 60, 106-14	7.9	226
231	The FKBP5-gene in depression and treatment response--an association study in the Sequenced Treatment Alternatives to Relieve Depression (STAR*D) Cohort. <i>Biological Psychiatry</i> , 2008 , 63, 1103-10	7.9	218
230	Genetic variants associated with response to lithium treatment in bipolar disorder: a genome-wide association study. <i>Lancet, The</i> , 2016 , 387, 1085-1093	40	216
229	Combined analysis from eleven linkage studies of bipolar disorder provides strong evidence of susceptibility loci on chromosomes 6q and 8q. <i>American Journal of Human Genetics</i> , 2005 , 77, 582-95	11	192
228	Genomewide linkage analyses of bipolar disorder: a new sample of 250 pedigrees from the National Institute of Mental Health Genetics Initiative. <i>American Journal of Human Genetics</i> , 2003 , 73, 107-14	11	186
227	Association between a functional serotonin transporter promoter polymorphism and citalopram treatment in adult outpatients with major depression. <i>Archives of General Psychiatry</i> , 2007 , 64, 783-92		182
226	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, 207-17	11.9	181
225	A genetic polymorphism for translocator protein 18 kDa affects both in vitro and in vivo radioligand binding in human brain to this putative biomarker of neuroinflammation. <i>Journal of Cerebral Blood Flow and Metabolism</i> , 2013 , 33, 53-8	7.3	175
224	Novel genetic loci associated with hippocampal volume. <i>Nature Communications</i> , 2017 , 8, 13624	17.4	173
223	Identification of pathways for bipolar disorder: a meta-analysis. <i>JAMA Psychiatry</i> , 2014 , 71, 657-64	14.5	172
222	Genetic markers of suicidal ideation emerging during citalopram treatment of major depression. <i>American Journal of Psychiatry</i> , 2007 , 164, 1530-8	11.9	171
221	Association of GRIK4 with outcome of antidepressant treatment in the STAR*D cohort. <i>American Journal of Psychiatry</i> , 2007 , 164, 1181-8	11.9	169
220	Genetic influences on schizophrenia and subcortical brain volumes: large-scale proof of concept. <i>Nature Neuroscience</i> , 2016 , 19, 420-431	25.5	163
219	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, 283-94	11	161
218	Clock genes may influence bipolar disorder susceptibility and dysfunctional circadian rhythm. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008 , 147B, 1047-55	3.5	159
217	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	15.1	155
216	Suggestive linkage to chromosomal regions 13q31 and 22q12 in families with psychotic bipolar disorder. <i>American Journal of Psychiatry</i> , 2003 , 160, 680-6	11.9	153
215	Convergent functional genomics of genome-wide association data for bipolar disorder: comprehensive identification of candidate genes, pathways and mechanisms. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2009 , 150B, 155-81	3.5	152

214	Brain-derived neurotrophic factor Val66Met polymorphism and antidepressant efficacy of ketamine in depressed patients. <i>Biological Psychiatry</i> , 2012 , 72, e27-8	7.9	150
213	Two variants in Ankyrin 3 (ANK3) are independent genetic risk factors for bipolar disorder. <i>Molecular Psychiatry</i> , 2009 , 14, 487-91	15.1	147
212	Novel genetic loci underlying human intracranial volume identified through genome-wide association. <i>Nature Neuroscience</i> , 2016 , 19, 1569-1582	25.5	147
211	Meta-analysis of genome-wide association data identifies a risk locus for major mood disorders on 3p21.1. <i>Nature Genetics</i> , 2010 , 42, 128-31	36.3	135
210	Enrichment of cis-regulatory gene expression SNPs and methylation quantitative trait loci among bipolar disorder susceptibility variants. <i>Molecular Psychiatry</i> , 2013 , 18, 340-6	15.1	134
209	Initial genomic scan of the NIMH genetics initiative bipolar pedigrees: Chromosomes 3, 5, 15, 16, 17, and 22 1997 , 74, 238-246		133
208	Initial genome scan of the NIMH genetics initiative bipolar pedigrees: chromosomes 4, 7, 9, 18, 19, 20, and 21q. <i>American Journal of Medical Genetics Part A</i> , 1997 , 74, 254-62		132
207	Genome-wide scan of bipolar disorder in 65 pedigrees: supportive evidence for linkage at 8q24, 18q22, 4q32, 2p12, and 13q12. <i>Molecular Psychiatry</i> , 2003 , 8, 288-98	15.1	127
206	Family-based association of FKBP5 in bipolar disorder. <i>Molecular Psychiatry</i> , 2009 , 14, 261-8	15.1	126
205	Diagnostic reliability of bipolar II disorder. <i>Archives of General Psychiatry</i> , 2002 , 59, 736-40		126
204	Genome-wide association study of 40,000 individuals identifies two novel loci associated with bipolar disorder. <i>Human Molecular Genetics</i> , 2016 , 25, 3383-3394	5.6	125
203	Parental diagnoses in youth with narrow phenotype bipolar disorder or severe mood dysregulation. <i>American Journal of Psychiatry</i> , 2007 , 164, 1238-41	11.9	124
202	Singleton deletions throughout the genome increase risk of bipolar disorder. <i>Molecular Psychiatry</i> , 2009 , 14, 376-80	15.1	121
201	A genome-wide association study of attempted suicide. <i>Molecular Psychiatry</i> , 2012 , 17, 433-44	15.1	117
200	Defining the phenotype in human genetic studies: forward genetics and reverse phenotyping. <i>Human Heredity</i> , 2004 , 58, 131-8	1.1	117
199	Genotype-phenotype studies in bipolar disorder showing association between the DAOA/G30 locus and persecutory delusions: a first step toward a molecular genetic classification of psychiatric phenotypes. <i>American Journal of Psychiatry</i> , 2005 , 162, 2101-8	11.9	116
198	Findings in an independent sample support an association between bipolar affective disorder and the G72/G30 locus on chromosome 13q33. <i>Molecular Psychiatry</i> , 2004 , 9, 87-92; image 5	15.1	114
197	Genome-wide association study of suicide attempts in mood disorder patients. <i>American Journal of Psychiatry</i> , 2010 , 167, 1499-507	11.9	113

196	Assessment of Response to Lithium Maintenance Treatment in Bipolar Disorder: A Consortium on Lithium Genetics (ConLiGen) Report. <i>PLoS ONE</i> , 2013 , 8, e65636	3.7	113
195	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, 3576-81	11.5	112
194	The International Consortium on Lithium Genetics (ConLiGen): an initiative by the NIMH and IGSLI to study the genetic basis of response to lithium treatment. <i>Neuropsychobiology</i> , 2010 , 62, 72-8	4	109
193	Initial genome scan of the NIMH genetics initiative bipolar pedigrees: chromosomes 1, 6, 8, 10, and 12. <i>American Journal of Medical Genetics Part A</i> , 1997 , 74, 247-53		109
192	Two gene co-expression modules differentiate psychotics and controls. <i>Molecular Psychiatry</i> , 2013 , 18, 1308-14	15.1	107
191	Genomic survey of bipolar illness in the NIMH genetics initiative pedigrees: a preliminary report. <i>American Journal of Medical Genetics Part A</i> , 1997 , 74, 227-37		103
190	Linkage of bipolar affective disorder to chromosome 18 markers in a new pedigree series. <i>American Journal of Human Genetics</i> , 1997 , 61, 1397-404	11	101
189	Increased levels of a mitochondrial DNA deletion in the brain of patients with bipolar disorder. <i>Biological Psychiatry</i> , 1997 , 42, 871-5	7.9	97
188	Meta-analysis of two genome-wide association studies of bipolar disorder reveals important points of agreement. <i>Molecular Psychiatry</i> , 2008 , 13, 466-7	15.1	91
187	Genetic association mapping at the crossroads: which test and why? Overview and practical guidelines. <i>American Journal of Medical Genetics Part A</i> , 2002 , 114, 1-11		90
186	Initial genome screen for bipolar disorder in the NIMH genetics initiative pedigrees: Chromosomes 2, 11, 13, 14, and X 1997 , 74, 263-269		89
185	What is familial about familial bipolar disorder? Resemblance among relatives across a broad spectrum of phenotypic characteristics. <i>Archives of General Psychiatry</i> , 2006 , 63, 1368-76		88
184	Brain-derived neurotrophic factor (BDNF) gene: no major impact on antidepressant treatment response. <i>International Journal of Neuropsychopharmacology</i> , 2010 , 13, 93-101	5.8	87
183	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	36.3	83
182	Molecular genetic overlap in bipolar disorder, schizophrenia, and major depressive disorder. <i>World Journal of Biological Psychiatry</i> , 2014 , 15, 200-8	3.8	82
181	RNA-sequencing of the brain transcriptome implicates dysregulation of neuroplasticity, circadian rhythms and GTPase binding in bipolar disorder. <i>Molecular Psychiatry</i> , 2014 , 19, 1179-85	15.1	81
180	Panic disorder with familial bipolar disorder. <i>Biological Psychiatry</i> , 1997 , 42, 90-5	7.9	81
179	Genome-wide association study of suicidal ideation emerging during citalopram treatment of depressed outpatients. <i>Pharmacogenetics and Genomics</i> , 2009 , 19, 666-74	1.9	80

178	A novel, heritable, expanding CTG repeat in an intron of the SEF2-1 gene on chromosome 18q21.1. <i>Human Molecular Genetics</i> , 1997 , 6, 1855-63	5.6	80
177	Genetic and clinical predictors of sexual dysfunction in citalopram-treated depressed patients. <i>Neuropsychopharmacology</i> , 2009 , 34, 1819-28	8.7	78
176	Genome-wide scan and conditional analysis in bipolar disorder: evidence for genomic interaction in the National Institute of Mental Health genetics initiative bipolar pedigrees. <i>Biological Psychiatry</i> , 2003 , 54, 1265-73	7.9	78
175	Association of Polygenic Score for Schizophrenia and HLA Antigen and Inflammation Genes With Response to Lithium in Bipolar Affective Disorder: A Genome-Wide Association Study. <i>JAMA Psychiatry</i> , 2018 , 75, 65-74	14.5	75
174	Nested association between genetic variation in tryptophan hydroxylase II, bipolar affective disorder, and suicide attempts. <i>Biological Psychiatry</i> , 2007 , 61, 181-6	7.9	75
173	Mood-incongruent psychotic features in bipolar disorder: familial aggregation and suggestive linkage to 2p11-q14 and 13q21-33. <i>American Journal of Psychiatry</i> , 2007 , 164, 236-47	11.9	75
172	Attempted suicide and alcoholism in bipolar disorder: clinical and familial relationships. <i>American Journal of Psychiatry</i> , 2000 , 157, 2048-50	11.9	75
171	Full-genome scan for linkage in 50 families segregating the bipolar affective disease phenotype. <i>American Journal of Human Genetics</i> , 2000 , 66, 205-15	11	72
170	The DISC locus and schizophrenia: evidence from an association study in a central European sample and from a meta-analysis across different European populations. <i>Human Molecular Genetics</i> , 2009 , 18, 2719-27	5.6	71
169	Genome-wide linkage and follow-up association study of postpartum mood symptoms. <i>American Journal of Psychiatry</i> , 2009 , 166, 1229-37	11.9	68
168	The bipolar disorder phenome database: a resource for genetic studies. <i>American Journal of Psychiatry</i> , 2007 , 164, 1229-37	11.9	66
167	Mitochondrial DNA sequence diversity in bipolar affective disorder. <i>American Journal of Psychiatry</i> , 2000 , 157, 1058-64	11.9	66
166	The genetics of bipolar disorder. <i>Molecular Psychiatry</i> , 2020 , 25, 544-559	15.1	66
165	Replication and meta-analysis of TMEM132D gene variants in panic disorder. <i>Translational Psychiatry</i> , 2012 , 2, e156	8.6	62
164	Loci on chromosomes 6q and 6p interact to increase susceptibility to bipolar affective disorder in the national institute of mental health genetics initiative pedigrees. <i>Biological Psychiatry</i> , 2004 , 56, 18-23	7.9	59
163	Pharmacogenetics Studies in STAR*D: Strengths, Limitations, and Results. <i>Psychiatric Services</i> , 2009 , 60, 1446-1457	3.3	58
162	Association study of Wnt signaling pathway genes in bipolar disorder. <i>Archives of General Psychiatry</i> , 2008 , 65, 785-93		58
161	Suggestive evidence of a locus on chromosome 10p using the NIMH genetics initiative bipolar affective disorder pedigrees 2000 , 96, 18-23		58

160	A network-based approach to prioritize results from genome-wide association studies. <i>PLoS ONE</i> , 2011 , 6, e24220	3.7	57
159	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, 344-52	7.9	56
158	Pharmacogenomics and personalized medicine in neuropsychiatry. <i>Neuron</i> , 2012 , 74, 773-6	13.9	55
157	Familial variation in episode frequency in bipolar affective disorder. <i>American Journal of Psychiatry</i> , 2005 , 162, 1266-72	11.9	55
156	Lack of support for a genetic association of the XBP1 promoter polymorphism with bipolar disorder in probands of European origin. <i>Nature Genetics</i> , 2004 , 36, 783-4; author reply 784-5	36.3	54
155	Genome-wide association of bipolar disorder suggests an enrichment of replicable associations in regions near genes. <i>PLoS Genetics</i> , 2011 , 7, e1002134	6	53
154	SSRI response in depression may be influenced by SNPs in HTR1B and HTR1A. <i>Psychiatric Genetics</i> , 2009 , 19, 281-91	2.9	53
153	Familial aggregation of psychotic symptoms in a replication set of 69 bipolar disorder pedigrees. <i>American Journal of Medical Genetics Part A</i> , 2003 , 116B, 90-7		51
152	Rapid mood switching and suicidality in familial bipolar disorder. <i>Bipolar Disorders</i> , 2005 , 7, 441-8	3.8	51
151	Bipolar disorder and panic disorder in families: an analysis of chromosome 18 data. <i>American Journal of Psychiatry</i> , 1998 , 155, 829-31	11.9	51
150	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	3.5	48
149	Evidence of association between brain-derived neurotrophic factor gene and bipolar disorder. <i>Psychiatric Genetics</i> , 2008 , 18, 267-74	2.9	47
148	Gene expression and genetic variation data implicate PCLO in bipolar disorder. <i>Biological Psychiatry</i> , 2011 , 69, 353-9	7.9	46
147	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, 407-18	15.1	46
146	NEDD4L on human chromosome 18q21 has multiple forms of transcripts and is a homologue of the mouse Nedd4-2 gene. <i>European Journal of Human Genetics</i> , 2001 , 9, 922-30	5.3	46
145	Familiarity of polarity at illness onset in bipolar affective disorder. <i>American Journal of Psychiatry</i> , 2006 , 163, 1754-9	11.9	43
144	Convergent genome wide association results for bipolar disorder and substance dependence. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2009 , 150B, 182-90	3.5	42
143	Familial aggregation of postpartum mood symptoms in bipolar disorder pedigrees. <i>Bipolar Disorders</i> , 2008 , 10, 38-44	3.8	42

142	Family-based association of YWHAH in psychotic bipolar disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2009 , 150B, 977-83	3.5	41
141	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, 850-77	3.5	41
140	A rare truncating mutation in ADH1C (G78Stop) shows significant association with Parkinson disease in a large international sample. <i>Archives of Neurology</i> , 2005 , 62, 74-8		41
139	Pharmacogenetics studies in STAR*D: strengths, limitations, and results. <i>Psychiatric Services</i> , 2009 , 60, 1446-57	3.3	41
138	Neurocognitive functioning in euthymic patients with bipolar disorder and unaffected relatives: A review of the literature. <i>Neuroscience and Biobehavioral Reviews</i> , 2016 , 69, 193-215	9	41
137	Review and Consensus on Pharmacogenomic Testing in Psychiatry. <i>Pharmacopsychiatry</i> , 2021 , 54, 5-17	2	40
136	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 , 156B, 370-8	3.5	39
135	Common and rare alleles of the serotonin transporter gene, SLC6A4, associated with Tourette's disorder. <i>Movement Disorders</i> , 2013 , 28, 1263-70	7	38
134	The genetics of panic disorder. <i>Journal of Medical Genetics</i> , 2011 , 48, 361-8	5.8	38
133	Genome-wide association studies of antidepressant outcome: a brief review. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2011 , 35, 1553-7	5.5	38
132	Attempted suicide in bipolar disorder pedigrees: evidence for linkage to 2p12. <i>Biological Psychiatry</i> , 2007 , 61, 725-7	7.9	37
131	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, 141-9	6.6	36
130	SERT Ileu425Val in autism, Asperger syndrome and obsessive-compulsive disorder. <i>Psychiatric Genetics</i> , 2008 , 18, 31-9	2.9	36
129	Increased gene expression of diacylglycerol kinase β in bipolar disorder. <i>International Journal of Neuropsychopharmacology</i> , 2010 , 13, 1127-8	5.8	35
128	Genome-wide linkage scan of 98 bipolar pedigrees and analysis of clinical covariates. <i>Molecular Psychiatry</i> , 2007 , 12, 630-9	15.1	33
127	Defining haplotype blocks and tag single-nucleotide polymorphisms in the human genome. <i>Human Molecular Genetics</i> , 2004 , 13, 335-42	5.6	33
126	Genome scan of a second wave of NIMH genetics initiative bipolar pedigrees: chromosomes 2, 11, 13, 14, and X. <i>American Journal of Medical Genetics Part A</i> , 2003 , 119B, 69-76		33
125	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, 360-7	4.6	32

124	Apparent replication of suggestive linkage on chromosome 16 in the NIMH genetics initiative bipolar pedigrees. <i>American Journal of Medical Genetics Part A</i> , 2002 , 114, 407-12		32
123	Genome scan of the fifty-six bipolar pedigrees from the NIMH genetics initiative replication sample: chromosomes 4, 7, 9, 18, 19, 20, and 21. <i>American Journal of Medical Genetics Part A</i> , 2003 , 121B, 21-7		32
122	Common and rare variant analysis in early-onset bipolar disorder vulnerability. <i>PLoS ONE</i> , 2014 , 9, e104336		32
121	Prediction of treatment outcomes in psychiatry--where do we stand ?. <i>Dialogues in Clinical Neuroscience</i> , 2014 , 16, 455-64	5.7	32
120	Variant GADL1 and response to lithium in bipolar I disorder. <i>New England Journal of Medicine</i> , 2014 , 370, 1857-9	59.2	31
119	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, 328-41	3.5	31
118	Genetic association studies in mood disorders: issues and promise. <i>International Review of Psychiatry</i> , 2004 , 16, 301-10	3.6	31
117	Rediscovering the value of families for psychiatric genetics research. <i>Molecular Psychiatry</i> , 2019 , 24, 523-535	5.5	30
116	Evidence of association between bipolar disorder and Citron on chromosome 12q24. <i>Molecular Psychiatry</i> , 2005 , 10, 807-9	15.1	30
115	Family-based association study of Neuregulin 1 with psychotic bipolar disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2009 , 150B, 693-702	3.5	29
114	Genetic variation in HTR2A influences serotonin transporter binding potential as measured using PET and [11C]DASB. <i>International Journal of Neuropsychopharmacology</i> , 2010 , 13, 715-24	5.8	28
113	Additional, physically ordered markers increase linkage signal for bipolar disorder on chromosome 18q22. <i>Biological Psychiatry</i> , 2003 , 53, 239-43	7.9	28
112	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	3.7	28
111	Genome-wide association study identifies 30 Loci Associated with Bipolar Disorder		28
110	Circadian genes and lithium response in bipolar disorders: associations with PPARGC1A (PGC-1 β) and RORA. <i>Genes, Brain and Behavior</i> , 2016 , 15, 660-8	3.6	28
109	Race, genetic ancestry and response to antidepressant treatment for major depression. <i>Neuropsychopharmacology</i> , 2013 , 38, 2598-606	8.7	27
108	An Integrative Genomic Study Implicates the Postsynaptic Density in the Pathogenesis of Bipolar Disorder. <i>Neuropsychopharmacology</i> , 2016 , 41, 886-95	8.7	25
107	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	7.2	25

106	Sequence variation in DOCK9 and heterogeneity in bipolar disorder. <i>Psychiatric Genetics</i> , 2007 , 17, 274-86.9	25
105	Response to Zhang et al. (2005): loss-of-function mutation in tryptophan hydroxylase-2 identified in unipolar major depression. <i>Neuron</i> 45, 11-16. <i>Neuron</i> , 2005 , 48, 702-3; author reply 705-6	13.9 25
104	Evaluation of Recipients of Positive and Negative Secondary Findings Evaluations in a Hybrid CLIA-Research Sequencing Pilot. <i>American Journal of Human Genetics</i> , 2018 , 103, 358-366	11 24
103	Genome-wide linkage analysis of 972 bipolar pedigrees using single-nucleotide polymorphisms. <i>Molecular Psychiatry</i> , 2012 , 17, 818-26	15.1 24
102	Genetic linkage and association studies in bipolar affective disorder: a time for optimism. <i>American Journal of Medical Genetics Part A</i> , 2003 , 123C, 36-47	24
101	Cutaneous cryptococcosis without evidence of systemic involvement. <i>Journal of the American Academy of Dermatology</i> , 1984 , 11, 371-4	4.5 24
100	Bcl-2 polymorphism influences gray matter volume in the ventral striatum in healthy humans. <i>Biological Psychiatry</i> , 2009 , 66, 804-7	7.9 23
99	Neurotransmission and bipolar disorder: a systematic family-based association study. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008 , 147B, 1270-7	3.5 22
98	Association study of phosphodiesterase genes in the Sequenced Treatment Alternatives to Relieve Depression sample. <i>Pharmacogenetics and Genomics</i> , 2009 , 19, 235-8	1.9 22
97	The pharmacogenetics of major depression: past, present, and future. <i>Biological Psychiatry</i> , 2007 , 62, 1205-7	7.9 21
96	Exome sequencing of a large family identifies potential candidate genes contributing risk to bipolar disorder. <i>Gene</i> , 2018 , 645, 119-123	3.8 20
95	Using duplicate genotyped data in genetic analyses: testing association and estimating error rates. <i>Statistical Applications in Genetics and Molecular Biology</i> , 2007 , 6, Article4	1.2 20
94	Can long-range microsatellite data be used to predict short-range linkage disequilibrium?. <i>Human Molecular Genetics</i> , 2002 , 11, 1363-72	5.6 20
93	A functional alternative splicing mutation in human tryptophan hydroxylase-2. <i>Molecular Psychiatry</i> , 2011 , 16, 1169-76	15.1 19
92	Anticipation in schizophrenia: a review and reconsideration. <i>American Journal of Medical Genetics Part A</i> , 1999 , 88, 686-93	19
91	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, 584-5	15.1 18
90	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	8.6 18
89	Association of polygenic score for major depression with response to lithium in patients with bipolar disorder. <i>Molecular Psychiatry</i> , 2021 , 26, 2457-2470	15.1 17

88	Amish revisited: next-generation sequencing studies of psychiatric disorders among the Plain people. <i>Trends in Genetics</i> , 2013 , 29, 412-8	8.5	17
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