Andrew McQuillin

List of Publications by Year
in descending order

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173 papers	46,581 citations	19657 61 h-index	5255 165 g-index
206 all docs	206 docs citations	206 times ranked	41803 citing authors

#	Article	IF	Citations
1	Genetic variation inTERTmodifies the risk of hepatocellular carcinoma in alcohol-related cirrhosis: results from a genome-wide case-control study. Gut, 2023, 72, 381-391.	12.1	19
2	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. Biological Psychiatry, 2022, 91, 102-117.	1.3	61
3	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. Biological Psychiatry, 2022, 91, 313-327.	1.3	114
4	A genetic risk score and diabetes predict development of alcohol-related cirrhosis in drinkers. Journal of Hepatology, 2022, 76, 275-282.	3.7	33
5	Adolescent Verbal Memory as a Psychosis Endophenotype: A Genome-Wide Association Study in an Ancestrally Diverse Sample. Genes, 2022, 13, 106.	2.4	2
6	Interaction Testing and Polygenic Risk Scoring to Estimate the Association of Common Genetic Variants With Treatment Resistance in Schizophrenia. JAMA Psychiatry, 2022, 79, 260.	11.0	44
7	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	27.8	929
8	Rare coding variants in ten genes confer substantial risk for schizophrenia. Nature, 2022, 604, 509-516.	27.8	326
9	Exome sequencing in bipolar disorder identifies AKAP11 as a risk gene shared with schizophrenia. Nature Genetics, 2022, 54, 541-547.	21.4	65
10	Genetic copy number variants, cognition and psychosis: a meta-analysis and a family study. Molecular Psychiatry, 2021, 26, 5307-5319.	7.9	18
11	Identifying risk factors involved in the common versus specific liabilities to substance use: A genetically informed approach. Addiction Biology, 2021, 26, e12944.	2.6	7
12	DNA methylation meta-analysis reveals cellular alterations in psychosis and markers of treatment-resistant schizophrenia. ELife, 2021, 10, .	6.0	72
13	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. Nature Genetics, 2021, 53, 817-829.	21.4	629
14	Characterisation of age and polarity at onset in bipolar disorder. British Journal of Psychiatry, 2021, 219, 659-669.	2.8	20
15	A machine learning caseâ€“control classifier for schizophrenia based on DNA methylation in blood. Translational Psychiatry, 2021, 11, 412.	4.8	16
16	Elevated common variant genetic risk for tourette syndrome in a densely-affected pedigree. Molecular Psychiatry, 2021, 26, 7522-7529.	7.9	8
17	Investigating rare pathogenic/likely pathogenic exonic variation in bipolar disorder. Molecular Psychiatry, 2021, 26, 5239-5250.	7.9	15
18	The influence of regression models on genome-wide association studies of alcohol dependence: a comparison of binary and quantitative analyses. Psychiatric Genetics, 2021, 31, 13-20.	1.1	3

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19	The Influence of CYP2D6 and CYP2C19 Genetic Variation on Diabetes Mellitus Risk in People Taking Antidepressants and Antipsychotics. Genes, 2021, 12, 1758.	2.4	8
20	Genetic Variation in HSD17B13 Reduces the Risk of Developing Cirrhosis and Hepatocellular Carcinoma in Alcohol Misusers. Hepatology, 2020, 72, 88-102.	7.3	76
21	The Genetics of the Mood Disorder Spectrum: Genome-wide Association Analyses of More Than 185,000 Cases and 439,000 Controls. Biological Psychiatry, 2020, 88, 169-184.	1.3	137
22	Predictive power of the ADHD GWAS 2019 polygenic risk scores in independent samples of bipolar patients with childhood ADHD. Journal of Affective Disorders, 2020, 265, 651-659.	4.1	15
23	Genome-Wide Association Study for Alcohol-Related Cirrhosis Identifies Risk Loci in MARC1 and HNRNPUL1. Gastroenterology, 2020, 159, 1276-1289.e7.	1.3	53
24	Genome-wide meta-analysis of problematic alcohol use in 435,563 individuals yields insights into biology and relationships with other traits. Nature Neuroscience, 2020, 23, 809-818.	14.8	242
25	The Communication of Metacognition for Social Strategy in Psychosis: An Exploratory Study. Schizophrenia Bulletin Open, 2020, 1, .	1.7	4
26	Heterozygous carriage of the alpha1-antitrypsin Pi*Z variant increases the risk to develop liver cirrhosis. Gut, 2019, 68, 1099-1107.	12.1	100
27	Ultra-Rare Genetic Variation in the Epilepsies: A Whole-Exome Sequencing Study of 17,606 Individuals. American Journal of Human Genetics, 2019, 105, 267-282.	6.2	237
28	SA140GENETIC RISK FACTORS OF ALCOHOL DEPENDENCE AND ANTISOCIAL PERSONALITY DISORDER. European Neuropsychopharmacology, 2019, 29, S1265-S1266.	0.7	0
29	SA33ADHD POLYGENIC RISK SCORES IN ROMANIAN AND UK BIPOLAR PATIENTS WITH CHILDHOOD ADHD. European Neuropsychopharmacology, 2019, 29, S1206.	0.7	0
30	META-ANALYSIS OF ALCOHOL DEPENDENCE GWAS DATA FROM EUROPEAN SAMPLES ASCERTAINED FROM CLINIC AND POPULATION BASED APPROACHES. European Neuropsychopharmacology, 2019, 29, S1036.	0.7	2
31	F130EXOME SEQUENCE ANALYSIS IDENTIFY RARE GENETIC VARIANT IMPLICATED IN SUSCEPTIBILITY TO SCHIZOPHRENIA. European Neuropsychopharmacology, 2019, 29, S1181.	0.7	0
32	GENETIC ASSOCIATION AND FUNCTIONAL CHARACTERIZATION OF A VARIANT IN THE MCPH1 GENE IN BIPOLAR DISORDER AND SCHIZOPHRENIA. European Neuropsychopharmacology, 2019, 29, S966-S967.	0.7	0
33	GWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. American Journal of Psychiatry, 2019, 176, 651-660.	7.2	186
34	GENOME-WIDE ASSOCIATION STUDY OF SUICIDE ATTEMPT IN MAJOR PSYCHIATRIC DISORDERS. European Neuropsychopharmacology, 2019, 29, S820-S821.	0.7	0
35	PS-177-HSD17B13 rs72613567 TA is associated with a reduced risk for developing hepatocellular carcinoma in patients with alcohol-related cirrhosis. Journal of Hepatology, 2019, 70, e109-e110.	3.7	5
36	Genome-wide association study identifies 30 loci associated with bipolar disorder. Nature Genetics, 2019, 51, 793-803.	21.4	1,191

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37	Genetic association and functional characterization of MCPH1 gene variation in bipolar disorder and schizophrenia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, 258-265.	1.7	2
38	Placental imprinted gene expression mediates the effects of maternal psychosocial stress during pregnancy on fetal growth. Journal of Developmental Origins of Health and Disease, 2019, 10, 196-205.	1.4	9
39	Genetic meta-analysis of diagnosed Alzheimerâ€™s disease identifies new risk loci and implicates AÎ², tau, immunity and lipid processing. Nature Genetics, 2019, 51, 414-430.	21.4	1,962
40	P: 81â€ƒPreviously Identified Candidate Gene Associations in Hepatic Encephalopathy Do Not Replicate in the STOPAH Cohort. American Journal of Gastroenterology, 2019, 114, S39-S39.	0.4	0
41	IDENTIFYING SUSCEPTIBILITY LOCI FOR TOURETTE'S SYNDROME IN A DENSELY AFFECTED PEDIGREE. European Neuropsychopharmacology, 2019, 29, S819.	0.7	0
42	Psychiatric Genetics, where we have been and where we are going. Psychiatric Genetics, 2019, 29, 131.	1.1	0
43	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. Cell, 2019, 179, 1469-1482.e11.	28.9	935
44	Physical Health and Clinical Phenotypes. , 2019, , 71-86.		2
45	Delineating the psychiatric and behavioral phenotype of recurrent 2q13 deletions and duplications. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2018, 177, 397-405.	1.7	16
46	People who survive an episode of severe alcoholic hepatitis should be advised to maintain total abstinence from alcohol. Hepatology, 2018, 67, 2479-2480.	7.3	3
47	Timing of prenatal exposure to trauma and altered placental expressions of hypothalamicâ€pituitaryâ€adrenal axis genes and genes driving neurodevelopment. Journal of Neuroendocrinology, 2018, 30, e12581.	2.6	24
48	Neurodevelopmental risk copy number variants in adults with intellectual disabilities and comorbid psychiatric disorders. British Journal of Psychiatry, 2018, 212, 287-294.	2.8	30
49	Rare variant analysis in multiply affected families, association studies and functional analysis suggest a role for the ITGÎ²4 gene in schizophrenia and bipolar disorder. Schizophrenia Research, 2018, 199, 181-188.	2.0	11
50	Genetic variants in PNPLA3 and TM6SF2 predispose to the development of hepatocellular carcinoma in individuals with alcohol-related cirrhosis. American Journal of Gastroenterology, 2018, 113, 1475-1483.	0.4	82
51	Genetic testing in intellectual disability psychiatry: Opinions and practices of <scp>UK</scp> child and intellectual disability psychiatrists. Journal of Applied Research in Intellectual Disabilities, 2018, 31, 273-284.	2.0	16
52	A polygenic risk score analysis of psychosis endophenotypes across brain functional, structural, and cognitive domains. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2018, 177, 21-34.	1.7	57
53	Reply to: â€œThe PNPLA3 SNP rs738409:G allele is associated with increased liver disease-associated mortality but reduced overall mortality in a population-based cohortâ€• Journal of Hepatology, 2018, 68, 860-862.	3.7	2
54	Exome sequence analysis and follow up genotyping implicates rare ULK1 variants to be involved in susceptibility to schizophrenia. Annals of Human Genetics, 2018, 82, 88-92.	0.8	16

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55	PWE-082â€...Genetic variants in CYP2D6 and the propensity to chronic liver disease in men chewing khat. , 2018, , .		0
56	Estimation of Genetic Correlation via Linkage Disequilibrium Score Regression and Genomic Restricted Maximum Likelihood. American Journal of Human Genetics, 2018, 102, 1185-1194.	6.2	119
57	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	12.6	1,085
58	Use of schizophrenia and bipolar disorder polygenic risk scores to identify psychotic disorders. British Journal of Psychiatry, 2018, 213, 535-541.	2.8	37
59	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. Cell, 2018, 173, 1705-1715.e16.	28.9	623
60	Homozygosity for rs738409:G in PNPLA3 is associated with increased mortality following an episode of severe alcoholic hepatitis. Journal of Hepatology, 2017, 67, 120-127.	3.7	52
61	Whole-exome sequencing of individuals from an isolated population implicates rare risk variants in bipolar disorder. Translational Psychiatry, 2017, 7, e1034-e1034.	4.8	24
62	Identification of rare nonsynonymous variants in SYNE1/CPG2 in bipolar affective disorder. Psychiatric Genetics, 2017, 27, 81-88.	1.1	6
63	Genome-wide association study of borderline personality disorder reveals genetic overlap with bipolar disorder, major depression and schizophrenia. Translational Psychiatry, 2017, 7, e1155-e1155.	4.8	150
64	Phenotypic Traits of Bipolar Disorder Predicted By Schizophrenia Associated Snps In Romanian Bipolar I Patients. Preliminary results. European Neuropsychopharmacology, 2017, 27, S386-S387.	0.7	0
65	Mutation intolerant genes and targets of FMRP are enriched for nonsynonymous alleles in schizophrenia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2017, 174, 724-731.	1.7	19
66	Genetic variation in GABRÎ²1 and the risk for developing alcohol dependence. Psychiatric Genetics, 2017, 27, 110-115.	1.1	6
67	Genetic Overlap Between Attention-Deficit/Hyperactivity Disorder and Bipolar Disorder: Evidence From Genome-wide Association Study Meta-analysis. Biological Psychiatry, 2017, 82, 634-641.	1.3	99
68	Chromosomal microarray testing in adults with intellectual disability presenting with comorbid psychiatric disorders. European Journal of Human Genetics, 2017, 25, 66-72.	2.8	30
69	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. Nature Genetics, 2017, 49, 27-35.	21.4	838
70	Genetic variants in ALDH1B1 and alcohol dependence risk in a British and Irish population: A bioinformatic and genetic study. PLoS ONE, 2017, 12, e0177009.	2.5	6
71	Genetic variant analysis of the putative regulatory regions of the LRRC7 gene in bipolar disorder. Psychiatric Genetics, 2016, 26, 99-100.	1.1	2
72	Association study of rare nonsynonymous variants of FTO in bipolar disorder. Psychiatric Genetics, 2016, 26, 140-141.	1.1	0

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73	Hypomethylation of FAM63B in bipolar disorder patients. Clinical Epigenetics, 2016, 8, 52.	4.1	24
74	An integrated genetic-epigenetic analysis of schizophrenia: evidence for co-localization of genetic associations and differential DNA methylation. Genome Biology, 2016, 17, 176.	8.8	287
75	Genetic variation in the miRâ€708 gene and its binding targets in bipolar disorder. Bipolar Disorders, 2016, 18, 650-656.	1.9	14
76	Genomeâ€wide association study reveals greater polygenic loading for schizophrenia in cases with a family history of illness. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2016, 171, 276-289.	1.7	28
77	Evidence for Genetic Overlap Between Schizophrenia and Age at First Birth in Women. JAMA Psychiatry, 2016, 73, 497.	11.0	51
78	Genetic influences on schizophrenia and subcortical brain volumes: large-scale proof of concept. Nature Neuroscience, 2016, 19, 420-431.	14.8	204
79	Rare loss-of-function variants in SETD1A are associated with schizophrenia and developmental disorders. Nature Neuroscience, 2016, 19, 571-577.	14.8	388
80	A novel Alzheimer disease locus located near the gene encoding tau protein. Molecular Psychiatry, 2016, 21, 108-117.	7.9	260
81	Phenotypic heterogeneity in study populations may significantly confound the results of genetic association studies on alcohol dependence. Psychiatric Genetics, 2015, 25, 234-240.	1.1	6
82	Joint Analysis of Psychiatric Disorders Increases Accuracy of Risk Prediction for Schizophrenia, Bipolar Disorder, and Major Depressive Disorder. American Journal of Human Genetics, 2015, 96, 283-294.	6.2	225
83	Psychiatric genome-wide association study analyses implicate neuronal, immune and histone pathways. Nature Neuroscience, 2015, 18, 199-209.	14.8	701
84	Modeling Linkage Disequilibrium Increases Accuracy of Polygenic Risk Scores. American Journal of Human Genetics, 2015, 97, 576-592.	6.2	1,098
85	Association of rare variation in the glutamate receptor gene SLC1A2 with susceptibility to bipolar disorder and schizophrenia. European Journal of Human Genetics, 2015, 23, 1200-1206.	2.8	45
86	Genetic variants in or nearADH1BandADH1Caffect susceptibility to alcohol dependence in a British and Irish population. Addiction Biology, 2015, 20, 594-604.	2.6	33
87	Analysis of t(9;17)(q33.2;q25.3) chromosomal breakpoint regions and genetic association reveals novel candidate genes for bipolar disorder. Bipolar Disorders, 2015, 17, 205-211.	1.9	19
88	A genome-wide association study confirms PNPLA3 and identifies TM6SF2 and MBOAT7 as risk loci for alcohol-related cirrhosis. Nature Genetics, 2015, 47, 1443-1448.	21.4	435
89	New data and an old puzzle: the negative association between schizophrenia and rheumatoid arthritis. International Journal of Epidemiology, 2015, 44, 1706-1721.	1.9	53
90	The UK10K project identifies rare variants in health and disease. Nature, 2015, 526, 82-90.	27.8	1,014

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91	Memory Decline in Down Syndrome and Its Relationship to iPF2alpha, a Urinary Marker of Oxidative Stress. PLoS ONE, 2014, 9, e97709.	2.5	17
92	Does rare matter? Copy number variants at 16p11.2 and the risk of psychosis: A systematic review of literature and meta-analysis. Schizophrenia Research, 2014, 159, 340-346.	2.0	27
93	A Rare Functional Noncoding Variant at the GWAS-Implicated MIR137/MIR2682 Locus Might Confer Risk to Schizophrenia and Bipolar Disorder. American Journal of Human Genetics, 2014, 95, 744-753.	6.2	91
94	Analysis of ANK3 and CACNA1C variants identified in bipolar disorder whole genome sequence data. Bipolar Disorders, 2014, 16, 583-591.	1.9	44
95	Allelic association, DNA resequencing and copy number variation at the metabotropic glutamate receptor GRM7 gene locus in bipolar disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 365-372.	1.7	31
96	An inherited duplication at the gene p21 Protein-Activated Kinase 7 (PAK7) is a risk factor for psychosis. Human Molecular Genetics, 2014, 23, 3316-3326.	2.9	37
97	The functional GRM3 Kozak sequence variant rs148754219 affects the risk of schizophrenia and alcohol dependence as well as bipolar disorder. Psychiatric Genetics, 2014, 24, 277-278.	1.1	33
98	Evidence for genetic susceptibility to the alcohol dependence syndrome from the thiamine transporter 2 gene solute carrier SLC19A3. Psychiatric Genetics, 2014, 24, 122-123.	1.1	2
99	Genome-wide association study of bipolar disorder in Canadian and UK populations corroborates disease loci including SYNE1 and CSMD1. BMC Medical Genetics, 2014, 15, 2.	2.1	106
100	Genetic association of the tachykinin receptor 1 TACR1 gene in bipolar disorder, attention deficit hyperactivity disorder, and the alcohol dependence syndrome. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 373-380.	1.7	39
101	Partitioning Heritability of Regulatory and Cell-Type-Specific Variants across 11 Common Diseases. American Journal of Human Genetics, 2014, 95, 535-552.	6.2	569
102	Polygenic dissection of diagnosis and clinical dimensions of bipolar disorder and schizophrenia. Molecular Psychiatry, 2014, 19, 1017-1024.	7.9	333
103	Biological insights from 108 schizophrenia-associated genetic loci. Nature, 2014, 511, 421-427.	27.8	6,934
104	Mutations in the Gabrb1 gene promote alcohol consumption through increased tonic inhibition. Nature Communications, 2013, 4, 2816.	12.8	44
105	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. Nature Genetics, 2013, 45, 984-994.	21.4	2,067
106	The effect of clozapine on mRNA expression for genes encoding G protein-coupled receptors and the protein components of clathrin-mediated endocytosis. Psychiatric Genetics, 2013, 23, 153-162.	1.1	10
107	Schizophrenia genetic variants are not associated with intelligence. Psychological Medicine, 2013, 43, 2563-2570.	4.5	40
108	Genetic Association, Mutation Screening, and Functional Analysis of a Kozak Sequence Variant in the Metabotropic Glutamate Receptor 3 Gene in Bipolar Disorder. JAMA Psychiatry, 2013, 70, 591.	11.0	43

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109	A nonconservative amino acid change in the UPF3B gene in a patient with schizophrenia. Psychiatric Genetics, 2012, 22, 150-151.	1.1	18
110	A gene expression and systems pathway analysis of the effects of clozapine compared to haloperidol in the mouse brain implicates susceptibility genes for schizophrenia. Journal of Psychopharmacology, 2012, 26, 1218-1230.	4.0	32
111	The Role of Variation at AÎ²PP, PSEN1, PSEN2, and MAPT in Late Onset Alzheimer's Disease. Journal of Alzheimer's Disease, 2012, 28, 377-387.	2.6	53
112	Genome-wide association study of Alzheimer's disease with psychotic symptoms. Molecular Psychiatry, 2012, 17, 1316-1327.	7.9	110
113	Sequencing of the ANKYRIN 3 gene (ANK3) encoding ankyrin G in bipolar disorder reveals a nonâ€conservative amino acid change in a short isoform of ankyrin G. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2012, 159B, 328-335.	1.7	14
114	Tests of linkage and allelic association between markers in the 1p36 PRKCZ (Protein Kinase C Zeta) gene region and bipolar affective disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2012, 159B, 201-209.	1.7	10
115	Genome-wide association study identifies five new schizophrenia loci. Nature Genetics, 2011, 43, 969-976.	21.4	1,758
116	Genetic association study of GABRA2 single nucleotide polymorphisms and electroencephalography in alcohol dependence. Neuroscience Letters, 2011, 500, 162-166.	2.1	38
117	Caseâ€"case genome-wide association analysis shows markers differentially associated with schizophrenia and bipolar disorder and implicates calcium channel genes. Psychiatric Genetics, 2011, 21, 1-4.	1.1	70
118	Common variants at ABCA7, MS4A6A/MS4A4E, EPHA1, CD33 and CD2AP are associated with Alzheimer's disease. Nature Genetics, 2011, 43, 429-435.	21.4	1,708
119	Analysis of genetic deletions and duplications in the University College London bipolar disorder case control sample. European Journal of Human Genetics, 2011, 19, 588-592.	2.8	38
120	GWA study data mining and independent replication identify cardiomyopathy-associated 5 (CMYA5) as a risk gene for schizophrenia. Molecular Psychiatry, 2011, 16, 1117-1129.	7.9	67
121	Genetic association and sequencing of the insulin-like growth factor 1 gene in bipolar affective disorder. , 2011, 156, 177-187.		26
122	No evidence that extended tracts of homozygosity are associated with Alzheimer's disease. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 764-771.	1.7	17
123	Polygenic dissection of the bipolar phenotype. British Journal of Psychiatry, 2011, 198, 284-288.	2.8	67
124	Maternally Derived Microduplications at 15q11-q13: Implication of Imprinted Genes in Psychotic Illness. American Journal of Psychiatry, 2011, 168, 408-417.	7.2	95
125	Large-scale genome-wide association analysis of bipolar disorder identifies a new susceptibility locus near ODZ4. Nature Genetics, 2011, 43, 977-983.	21.4	1,283
126	Lack of allelic association between markers at the DRD2 and ANKK1 gene loci with the alcohol-dependence syndrome and criminal activity. Psychiatric Genetics, 2011, 21, 323-324.	1.1	6

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127	Confirmation of prior evidence of genetic susceptibility to alcoholism in a genome-wide association study of comorbid alcoholism and bipolar disorder. Psychiatric Genetics, 2011, 21, 294-306.	1.1	59
128	Support of association between BRD1 and both schizophrenia and bipolar affective disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 582-591.	1.7	47
129	A threonine to isoleucine missense mutation in the pericentriolar material 1 gene is strongly associated with schizophrenia. Molecular Psychiatry, 2010, 15, 615-628.	7.9	50
130	Genetic Evidence Implicates the Immune System and Cholesterol Metabolism in the Aetiology of Alzheimer's Disease. PLoS ONE, 2010, 5, e13950.	2.5	347
131	Genetic power of a Brazilian three-generation family with generalized aggressive periodontitis. Brazilian Dental Journal, 2010, 21, 137-141.	1.1	5
132	NK₁ (TACR₁) receptor gene â€˜knockoutâ€™ mouse phenotype predicts genetic association with ADHD. Journal of Psychopharmacology, 2010, 24, 27-38.	4.0	329
133	Genome-Wide Association Study of Suicide Attempts in Mood Disorder Patients. American Journal of Psychiatry, 2010, 167, 1499-1507.	7.2	140
134	Effect of the 2004 tsunami on suicide rates in Sri Lanka. Psychiatric Bulletin, 2009, 33, 179-180.	0.3	14
135	A Genomewide Association Study of Response to Lithium for Prevention of Recurrence in Bipolar Disorder. American Journal of Psychiatry, 2009, 166, 718-725.	7.2	145
136	Meta-analysis of 32 genome-wide linkage studies of schizophrenia. Molecular Psychiatry, 2009, 14, 774-785.	7.9	235
137	DISC1 association, heterogeneity and interplay in schizophrenia and bipolar disorder. Molecular Psychiatry, 2009, 14, 865-873.	7.9	140
138	Caseâ€“control studies show that a non-conservative amino-acid change from a glutamine to arginine in the P2RX7 purinergic receptor protein is associated with both bipolar- and unipolar-affective disorders. Molecular Psychiatry, 2009, 14, 614-620.	7.9	101
139	Common polygenic variation contributes to risk of schizophrenia and bipolar disorder. Nature, 2009, 460, 748-752.	27.8	4,345
140	Genome-wide association study identifies variants at CLU and PICALM associated with Alzheimer's disease. Nature Genetics, 2009, 41, 1088-1093.	21.4	2,697
141	Genetics of attention-deficit hyperactivity disorder (ADHD). Neuropharmacology, 2009, 57, 590-600.	4.1	113
142	Evidence for the association of the DAOA (G72) gene with schizophrenia and bipolar disorder but not for the association of the DAO gene with schizophrenia. Behavioral and Brain Functions, 2009, 5, 28.	3.3	40
143	No evidence for excess runs of homozygosity in bipolar disorder. Psychiatric Genetics, 2009, 19, 165-170.	1.1	35
144	Confirmation of the genetic association between the U2AF homology motif (UHM) kinase 1 (UHMK1) gene and schizophrenia on chromosome 1q23.3. European Journal of Human Genetics, 2008, 16, 1275-1282.	2.8	18

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145	Rare chromosomal deletions and duplications increase risk of schizophrenia. Nature, 2008, 455, 237-241.	27.8	1,387
146	Collaborative genome-wide association analysis supports a role for ANK3 and CACNA1C in bipolar disorder. Nature Genetics, 2008, 40, 1056-1058.	21.4	1,102
147	Whole-genome association study of bipolar disorder. Molecular Psychiatry, 2008, 13, 558-569.	7.9	642
148	A Genetic Association Study of Chromosome 11q22-24 in Two Different Samples Implicates the FXYD6 Gene, Encoding Phosphohippolin, in Susceptibility to Schizophrenia. American Journal of Human Genetics, 2007, 80, 664-672.	6.2	32
149	A microarray gene expression study of the molecular pharmacology of lithium carbonate on mouse brain mRNA to understand the neurobiology of mood stabilization and treatment of bipolar affective disorder. Pharmacogenetics and Genomics, 2007, 17, 605-617.	1.5	133
150	Replication of genetic association studies between markers at the Epsin 4 gene locus and schizophrenia in two Han Chinese samples. Schizophrenia Research, 2007, 89, 357-359.	2.0	9
151	Failure to confirm allelic and haplotypic association between markers at the chromosome 6p22.3 dystrobrevin-binding protein 1 (DTNBP1) locus and schizophrenia. Behavioral and Brain Functions, 2007, 3, 50.	3.3	20
152	Fine Mapping by Genetic Association Implicates the Chromosome 1q23.3 Gene UHMK1, Encoding a Serine/Threonine Protein Kinase, as a Novel Schizophrenia Susceptibility Gene. Biological Psychiatry, 2007, 61, 873-879.	1.3	35
153	Geneâ€“Brain Structure Relationships: Arbitrary Assumptions of Heterogeneity Generate Unfalsifiable Claimsâ€”Reply. Archives of General Psychiatry, 2007, 64, 1098.	12.3	0
154	Failure to Confirm Allelic Association Between Markers at the CAPON Gene Locus and Schizophrenia in a British Sample. Biological Psychiatry, 2006, 59, 195-197.	1.3	28
155	Genetic Association and Brain Morphology Studies and the Chromosome 8p22 Pericentriolar Material 1 (PCM1) Gene in Susceptibility to Schizophrenia. Archives of General Psychiatry, 2006, 63, 844.	12.3	82
156	Identification of the Slynar Gene (AY070435) and Related Brain Expressed Sequences as a Candidate Gene for Susceptibility to Affective Disorders Through Allelic and Haplotypic Association With Bipolar Disorder on Chromosome 12q24. American Journal of Psychiatry, 2006, 163, 1767-1776.	7.2	13
157	Fine mapping of a susceptibility locus for bipolar and genetically related unipolar affective disorders, to a region containing the C21ORF29 and TRPM2 genes on chromosome 21q22.3. Molecular Psychiatry, 2006, 11, 134-142.	7.9	81
158	Failure to confirm genetic association between schizophrenia and markers on chromosome 1q23.3 in the region of the gene encoding the regulator of G-protein signaling 4 protein (RGS4). American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2006, 141B, 296-300.	1.7	32
159	Direct genomic PCR sequencing of the high affinity thiamine transporter (SLC19A2) gene identifies three genetic variants in Wernicke Korsakoff syndrome (WKS). American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2005, 137B, 17-19.	1.7	27
160	Genetic linkage analysis supports the presence of two susceptibility loci for alcoholism and heavy drinking on chromosome 1p22.1-11.2 and 1q21.3-24.2. BMC Genetics, 2005, 6, 11.	2.7	36
161	The Epsin 4 Gene on Chromosome 5q, Which Encodes the Clathrin-Associated Protein Enthoprotin, Is Involved in the Genetic Susceptibility to Schizophrenia. American Journal of Human Genetics, 2005, 76, 902-907.	6.2	62
162	No association between a neuronal nitric oxide synthase (NOS1) gene polymorphism on chromosome 12q24 and bipolar disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2004, 124B, 73-75.	1.7	19

#	Article	IF	Citations
163	Genome scan of Tourette syndrome in a single large pedigree shows some support for linkage to regions of chromosomes 5, 10 and 13. Psychiatric Genetics, 2004, 14, 83-87.	1.1	41
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166	Title is missing!. Psychiatric Genetics, 2003, 13, 77-84.	1.1	16
167	Genome scan of pedigrees multiply affected with bipolar disorder provides further support for the presence of a susceptibility locus on chromosome 12q23-q24, and suggests the presence of additional loci on 1p and 1q. Psychiatric Genetics, 2003, 13, 77-84.	1.1	136
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169	A novel polymorphism in exon 11 of the WKL1 gene, shows no association with schizophrenia. European Journal of Human Genetics, 2002, 10, 491-494.	2.8	14
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173	The genetics of schizophrenia. , 0, , 230-261.		1