

# Andrew McQuillin

## List of Publications by Year in Descending Order

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**Version:** 2024-04-09

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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.

167 papers	33,302 citations	57 h-index	182 g-index
206 ext. papers	41,183 ext. citations	10.9 avg, IF	8.04 L-index

#	Paper	IF	Citations
167	Adolescent Verbal Memory as a Psychosis Endophenotype: A Genome-Wide Association Study in an Ancestrally Diverse Sample.. <i>Genes</i> , <b>2022</b> , 13,	4.2	1
166	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. <i>Biological Psychiatry</i> , <b>2022</b> , 91, 102-117	7.9	11
165	Mapping genomic loci implicates genes and synaptic biology in schizophrenia.. <i>Nature</i> , <b>2022</b> ,	50.4	35
164	Rare coding variants in ten genes confer substantial risk for schizophrenia.. <i>Nature</i> , <b>2022</b> ,	50.4	16
163	The Influence of CYP2D6 and CYP2C19 Genetic Variation on Diabetes Mellitus Risk in People Taking Antidepressants and Antipsychotics. <i>Genes</i> , <b>2021</b> , 12,	4.2	3
162	A genetic risk score and diabetes predict development of alcohol-related cirrhosis in drinkers. <i>Journal of Hepatology</i> , <b>2021</b> ,	13.4	4
161	The influence of regression models on genome-wide association studies of alcohol dependence: a comparison of binary and quantitative analyses. <i>Psychiatric Genetics</i> , <b>2021</b> , 31, 13-20	2.9	0
160	Genetic copy number variants, cognition and psychosis: a meta-analysis and a family study. <i>Molecular Psychiatry</i> , <b>2021</b> , 26, 5307-5319	15.1	5
159	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. <i>Nature Genetics</i> , <b>2021</b> , 53, 817-829	36.3	83
158	Identifying risk factors involved in the common versus specific liabilities to substance use: A genetically informed approach. <i>Addiction Biology</i> , <b>2021</b> , 26, e12944	4.6	4
157	DNA methylation meta-analysis reveals cellular alterations in psychosis and markers of treatment-resistant schizophrenia. <i>ELife</i> , <b>2021</b> , 10,	8.9	15
156	Characterisation of age and polarity at onset in bipolar disorder.. <i>British Journal of Psychiatry</i> , <b>2021</b> , 219, 659-669	5.4	2
155	A machine learning case-control classifier for schizophrenia based on DNA methylation in blood. <i>Translational Psychiatry</i> , <b>2021</b> , 11, 412	8.6	3
154	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. <i>Biological Psychiatry</i> , <b>2021</b> ,	7.9	11
153	Elevated common variant genetic risk for tourette syndrome in a densely-affected pedigree. <i>Molecular Psychiatry</i> , <b>2021</b> ,	15.1	1
152	Investigating rare pathogenic/likely pathogenic exonic variation in bipolar disorder. <i>Molecular Psychiatry</i> , <b>2021</b> , 26, 5239-5250	15.1	3
151	Genome-Wide Association Study for Alcohol-Related Cirrhosis Identifies Risk Loci in MARC1 and HNRNPUL1. <i>Gastroenterology</i> , <b>2020</b> , 159, 1276-1289.e7	13.3	19

150	The Communication of Metacognition for Social Strategy in Psychosis: An Exploratory Study. <i>Schizophrenia Bulletin Open</i> , <b>2020</b> , 1,	2.2	2
149	Genome-wide meta-analysis of problematic alcohol use in 435,563 individuals yields insights into biology and relationships with other traits. <i>Nature Neuroscience</i> , <b>2020</b> , 23, 809-818	25.5	69
148	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> , <b>2020</b> , 88, 169-184	7.9	57
147	Predictive power of the ADHD GWAS 2019 polygenic risk scores in independent samples of bipolar patients with childhood ADHD. <i>Journal of Affective Disorders</i> , <b>2020</b> , 265, 651-659	6.6	7
146	Genetic Variation in HSD17B13 Reduces the Risk of Developing Cirrhosis and Hepatocellular Carcinoma in Alcohol Misusers. <i>Hepatology</i> , <b>2020</b> , 72, 88-102	11.2	46
145	GWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. <i>American Journal of Psychiatry</i> , <b>2019</b> , 176, 651-660	11.9	103
144	PS-177-HSD17B13 rs72613567 TA is associated with a reduced risk for developing hepatocellular carcinoma in patients with alcohol-related cirrhosis. <i>Journal of Hepatology</i> , <b>2019</b> , 70, e109-e110	13.4	4
143	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , <b>2019</b> , 51, 793-803	36.3	662
142	Genetic association and functional characterization of MCPH1 gene variation in bipolar disorder and schizophrenia. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2019</b> , 180, 258-265	3.5	1
141	Placental imprinted gene expression mediates the effects of maternal psychosocial stress during pregnancy on fetal growth. <i>Journal of Developmental Origins of Health and Disease</i> , <b>2019</b> , 10, 196-205	2.4	4
140	Heterozygous carriage of the alpha1-antitrypsin Pi*Z variant increases the risk to develop liver cirrhosis. <i>Gut</i> , <b>2019</b> , 68, 1099-1107	19.2	62
139	Ultra-Rare Genetic Variation in the Epilepsies: A Whole-Exome Sequencing Study of 17,606 Individuals. <i>American Journal of Human Genetics</i> , <b>2019</b> , 105, 267-282	11	104
138	Physical Health and Clinical Phenotypes <b>2019</b> , 71-86		1
137	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates A $\beta$ tau, immunity and lipid processing. <i>Nature Genetics</i> , <b>2019</b> , 51, 414-430	36.3	917
136	P: 81 Previously Identified Candidate Gene Associations in Hepatic Encephalopathy Do Not Replicate in the STOPAH Cohort. <i>American Journal of Gastroenterology</i> , <b>2019</b> , 114, S39-S39	0.7	
135	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , <b>2019</b> , 179, 1469-1482.e11	56.2	402
134	Delineating the psychiatric and behavioral phenotype of recurrent 2q13 deletions and duplications. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2018</b> , 177, 397-405	3.5	10
133	People who survive an episode of severe alcoholic hepatitis should be advised to maintain total abstinence from alcohol. <i>Hepatology</i> , <b>2018</b> , 67, 2479-2480	11.2	2

132	Timing of prenatal exposure to trauma and altered placental expressions of hypothalamic-pituitary-adrenal axis genes and genes driving neurodevelopment. <i>Journal of Neuroendocrinology</i> , <b>2018</b> , 30, e12581	3.8	17
131	Neurodevelopmental risk copy number variants in adults with intellectual disabilities and comorbid psychiatric disorders. <i>British Journal of Psychiatry</i> , <b>2018</b> , 212, 287-294	5.4	21
130	Rare variant analysis in multiply affected families, association studies and functional analysis suggest a role for the ITGA gene in schizophrenia and bipolar disorder. <i>Schizophrenia Research</i> , <b>2018</b> , 199, 181-188	3.6	8
129	Genetic variants in PNPLA3 and TM6SF2 predispose to the development of hepatocellular carcinoma in individuals with alcohol-related cirrhosis. <i>American Journal of Gastroenterology</i> , <b>2018</b> , 113, 1475-1483	0.7	50
128	Genetic testing in intellectual disability psychiatry: Opinions and practices of UK child and intellectual disability psychiatrists. <i>Journal of Applied Research in Intellectual Disabilities</i> , <b>2018</b> , 31, 273-284	2.2	11
127	A polygenic risk score analysis of psychosis endophenotypes across brain functional, structural, and cognitive domains. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2018</b> , 177, 21-34	3.5	39
126	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , <b>2018</b> , 360,	33.3	666
125	Use of schizophrenia and bipolar disorder polygenic risk scores to identify psychotic disorders. <i>British Journal of Psychiatry</i> , <b>2018</b> , 213, 535-541	5.4	21
124	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. <i>Cell</i> , <b>2018</b> , 173, 1705-1715.e16	56.2	360
123	Reply to: "The PNPLA3 SNP rs738409:G allele is associated with increased liver disease-associated mortality but reduced overall mortality in a population-based cohort". <i>Journal of Hepatology</i> , <b>2018</b> , 68, 860-862	13.4	2
122	Exome sequence analysis and follow up genotyping implicates rare ULK1 variants to be involved in susceptibility to schizophrenia. <i>Annals of Human Genetics</i> , <b>2018</b> , 82, 88-92	2.2	13
121	Estimation of Genetic Correlation via Linkage Disequilibrium Score Regression and Genomic Restricted Maximum Likelihood. <i>American Journal of Human Genetics</i> , <b>2018</b> , 102, 1185-1194	11	55
120	Homozygosity for rs738409:G in PNPLA3 is associated with increased mortality following an episode of severe alcoholic hepatitis. <i>Journal of Hepatology</i> , <b>2017</b> , 67, 120-127	13.4	36
119	Whole-exome sequencing of individuals from an isolated population implicates rare risk variants in bipolar disorder. <i>Translational Psychiatry</i> , <b>2017</b> , 7, e1034	8.6	18
118	Identification of rare nonsynonymous variants in SYNE1/CPG2 in bipolar affective disorder. <i>Psychiatric Genetics</i> , <b>2017</b> , 27, 81-88	2.9	4
117	Genome-wide association study of borderline personality disorder reveals genetic overlap with bipolar disorder, major depression and schizophrenia. <i>Translational Psychiatry</i> , <b>2017</b> , 7, e1155	8.6	100
116	Mutation intolerant genes and targets of FMRP are enriched for nonsynonymous alleles in schizophrenia. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2017</b> , 174, 724-731	3.5	16
115	Genetic variation in GABRG1 and the risk for developing alcohol dependence. <i>Psychiatric Genetics</i> , <b>2017</b> , 27, 110-115	2.9	5

114	Genetic Overlap Between Attention-Deficit/Hyperactivity Disorder and Bipolar Disorder: Evidence From Genome-wide Association Study Meta-analysis. <i>Biological Psychiatry</i> , <b>2017</b> , 82, 634-641	7.9	66
113	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. <i>Nature Genetics</i> , <b>2017</b> , 49, 27-35	36.3	530
112	Genetic variants in ALDH1B1 and alcohol dependence risk in a British and Irish population: A bioinformatic and genetic study. <i>PLoS ONE</i> , <b>2017</b> , 12, e0177009	3.7	5
111	A novel Alzheimer disease locus located near the gene encoding tau protein. <i>Molecular Psychiatry</i> , <b>2016</b> , 21, 108-17	15.1	175
110	An integrated genetic-epigenetic analysis of schizophrenia: evidence for co-localization of genetic associations and differential DNA methylation. <i>Genome Biology</i> , <b>2016</b> , 17, 176	18.3	189
109	Genetic variation in the miR-708 gene and its binding targets in bipolar disorder. <i>Bipolar Disorders</i> , <b>2016</b> , 18, 650-656	3.8	10
108	Genome-wide association study reveals greater polygenic loading for schizophrenia in cases with a family history of illness. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2016</b> , 171B, 276-89	3.5	23
107	Evidence for Genetic Overlap Between Schizophrenia and Age at First Birth in Women. <i>JAMA Psychiatry</i> , <b>2016</b> , 73, 497-505	14.5	40
106	Genetic influences on schizophrenia and subcortical brain volumes: large-scale proof of concept. <i>Nature Neuroscience</i> , <b>2016</b> , 19, 420-431	25.5	163
105	Rare loss-of-function variants in SETD1A are associated with schizophrenia and developmental disorders. <i>Nature Neuroscience</i> , <b>2016</b> , 19, 571-7	25.5	284
104	Chromosomal microarray testing in adults with intellectual disability presenting with comorbid psychiatric disorders. <i>European Journal of Human Genetics</i> , <b>2016</b> , 25, 66-72	5.3	24
103	Genetic variant analysis of the putative regulatory regions of the LRR7 gene in bipolar disorder. <i>Psychiatric Genetics</i> , <b>2016</b> , 26, 99-100	2.9	1
102	Association study of rare nonsynonymous variants of FTO in bipolar disorder. <i>Psychiatric Genetics</i> , <b>2016</b> , 26, 140-1	2.9	
101	Hypomethylation of FAM63B in bipolar disorder patients. <i>Clinical Epigenetics</i> , <b>2016</b> , 8, 52	7.7	14
100	Psychiatric genome-wide association study analyses implicate neuronal, immune and histone pathways. <i>Nature Neuroscience</i> , <b>2015</b> , 18, 199-209	25.5	572
99	Modeling Linkage Disequilibrium Increases Accuracy of Polygenic Risk Scores. <i>American Journal of Human Genetics</i> , <b>2015</b> , 97, 576-92	11	649
98	Association of rare variation in the glutamate receptor gene SLC1A2 with susceptibility to bipolar disorder and schizophrenia. <i>European Journal of Human Genetics</i> , <b>2015</b> , 23, 1200-6	5.3	33
97	Genetic variants in or near ADH1B and ADH1C affect susceptibility to alcohol dependence in a British and Irish population. <i>Addiction Biology</i> , <b>2015</b> , 20, 594-604	4.6	22

96	Analysis of t(9;17)(q33.2;q25.3) chromosomal breakpoint regions and genetic association reveals novel candidate genes for bipolar disorder. <i>Bipolar Disorders</i> , <b>2015</b> , 17, 205-11	3.8	13
95	A genome-wide association study confirms PNPLA3 and identifies TM6SF2 and MBOAT7 as risk loci for alcohol-related cirrhosis. <i>Nature Genetics</i> , <b>2015</b> , 47, 1443-8	36.3	303
94	New data and an old puzzle: the negative association between schizophrenia and rheumatoid arthritis. <i>International Journal of Epidemiology</i> , <b>2015</b> , 44, 1706-21	7.8	43
93	The UK10K project identifies rare variants in health and disease. <i>Nature</i> , <b>2015</b> , 526, 82-90	50.4	776
92	Phenotypic heterogeneity in study populations may significantly confound the results of genetic association studies on alcohol dependence. <i>Psychiatric Genetics</i> , <b>2015</b> , 25, 234-40	2.9	5
91	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> , <b>2015</b> , 96, 283-94	11	161
90	Genome-wide association study of bipolar disorder in Canadian and UK populations corroborates disease loci including SYNE1 and CSMD1. <i>BMC Medical Genetics</i> , <b>2014</b> , 15, 2	2.1	85
89	Genetic association of the tachykinin receptor 1 TACR1 gene in bipolar disorder, attention deficit hyperactivity disorder, and the alcohol dependence syndrome. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2014</b> , 165B, 373-80	3.5	32
88	Partitioning heritability of regulatory and cell-type-specific variants across 11 common diseases. <i>American Journal of Human Genetics</i> , <b>2014</b> , 95, 535-52	11	411
87	Polygenic dissection of diagnosis and clinical dimensions of bipolar disorder and schizophrenia. <i>Molecular Psychiatry</i> , <b>2014</b> , 19, 1017-1024	15.1	258
86	Biological insights from 108 schizophrenia-associated genetic loci. <i>Nature</i> , <b>2014</b> , 511, 421-7	50.4	5249
85	Memory decline in Down syndrome and its relationship to iPF2alpha, a urinary marker of oxidative stress. <i>PLoS ONE</i> , <b>2014</b> , 9, e97709	3.7	17
84	Does rare matter? Copy number variants at 16p11.2 and the risk of psychosis: a systematic review of literature and meta-analysis. <i>Schizophrenia Research</i> , <b>2014</b> , 159, 340-6	3.6	22
83	A rare functional noncoding variant at the GWAS-implicated MIR137/MIR2682 locus might confer risk to schizophrenia and bipolar disorder. <i>American Journal of Human Genetics</i> , <b>2014</b> , 95, 744-53	11	72
82	Analysis of ANK3 and CACNA1C variants identified in bipolar disorder whole genome sequence data. <i>Bipolar Disorders</i> , <b>2014</b> , 16, 583-91	3.8	36
81	Allelic association, DNA resequencing and copy number variation at the metabotropic glutamate receptor GRM7 gene locus in bipolar disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2014</b> , 165B, 365-72	3.5	24
80	An inherited duplication at the gene p21 Protein-Activated Kinase 7 (PAK7) is a risk factor for psychosis. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 3316-26	5.6	32
79	The functional GRM3 Kozak sequence variant rs148754219 affects the risk of schizophrenia and alcohol dependence as well as bipolar disorder. <i>Psychiatric Genetics</i> , <b>2014</b> , 24, 277-8	2.9	26



78	Evidence for genetic susceptibility to the alcohol dependence syndrome from the thiamine transporter 2 gene solute carrier SLC19A3. <i>Psychiatric Genetics</i> , <b>2014</b> , 24, 122-3	2.9	1
77	Mutations in the Gabrb1 gene promote alcohol consumption through increased tonic inhibition. <i>Nature Communications</i> , <b>2013</b> , 4, 2816	17.4	37
76	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. <i>Nature Genetics</i> , <b>2013</b> , 45, 984-94	36.3	1628
75	The effect of clozapine on mRNA expression for genes encoding G protein-coupled receptors and the protein components of clathrin-mediated endocytosis. <i>Psychiatric Genetics</i> , <b>2013</b> , 23, 153-62	2.9	7
74	Schizophrenia genetic variants are not associated with intelligence. <i>Psychological Medicine</i> , <b>2013</b> , 43, 2563-70	6.9	34
73	Genetic association, mutation screening, and functional analysis of a Kozak sequence variant in the metabotropic glutamate receptor 3 gene in bipolar disorder. <i>JAMA Psychiatry</i> , <b>2013</b> , 70, 591-8	14.5	28
72	Tests of linkage and allelic association between markers in the 1p36 PRKCZ (protein kinase C zeta) gene region and bipolar affective disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2012</b> , 159B, 201-9	3.5	8
71	Genome-wide association study of Alzheimer's disease with psychotic symptoms. <i>Molecular Psychiatry</i> , <b>2012</b> , 17, 1316-27	15.1	90
70	Sequencing of the ANKYRIN 3 gene (ANK3) encoding ankyrin G in bipolar disorder reveals a non-conservative amino acid change in a short isoform of ankyrin G. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2012</b> , 159B, 328-35	3.5	14
69	A nonconservative amino acid change in the UPF3B gene in a patient with schizophrenia. <i>Psychiatric Genetics</i> , <b>2012</b> , 22, 150-1	2.9	13
68	A gene expression and systems pathway analysis of the effects of clozapine compared to haloperidol in the mouse brain implicates susceptibility genes for schizophrenia. <i>Journal of Psychopharmacology</i> , <b>2012</b> , 26, 1218-30	4.6	28
67	The role of variation at APP, PSEN1, PSEN2, and MAPT in late onset Alzheimer's disease. <i>Journal of Alzheimer's Disease</i> , <b>2012</b> , 28, 377-87	4.3	47
66	Genome-wide association study identifies five new schizophrenia loci. <i>Nature Genetics</i> , <b>2011</b> , 43, 969-76	36.3	1508
65	Genetic association study of GABRA2 single nucleotide polymorphisms and electroencephalography in alcohol dependence. <i>Neuroscience Letters</i> , <b>2011</b> , 500, 162-6	3.3	27
64	Case-case genome-wide association analysis shows markers differentially associated with schizophrenia and bipolar disorder and implicates calcium channel genes. <i>Psychiatric Genetics</i> , <b>2011</b> , 21, 1-4	2.9	58
63	Common variants at ABCA7, MS4A6A/MS4A4E, EPHA1, CD33 and CD2AP are associated with Alzheimer's disease. <i>Nature Genetics</i> , <b>2011</b> , 43, 429-35	36.3	1421
62	Analysis of genetic deletions and duplications in the University College London bipolar disorder case control sample. <i>European Journal of Human Genetics</i> , <b>2011</b> , 19, 588-92	5.3	35
61	GWA study data mining and independent replication identify cardiomyopathy-associated 5 (CMYA5) as a risk gene for schizophrenia. <i>Molecular Psychiatry</i> , <b>2011</b> , 16, 1117-29	15.1	58

60	Genetic association and sequencing of the insulin-like growth factor 1 gene in bipolar affective disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2011</b> , 156, 177-87	3.5	22
59	No evidence that extended tracts of homozygosity are associated with Alzheimer's disease. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2011</b> , 156B, 764-71	3.5	15
58	Polygenic dissection of the bipolar phenotype. <i>British Journal of Psychiatry</i> , <b>2011</b> , 198, 284-8	5.4	57
57	Maternally derived microduplications at 15q11-q13: implication of imprinted genes in psychotic illness. <i>American Journal of Psychiatry</i> , <b>2011</b> , 168, 408-17	11.9	84
56	Large-scale genome-wide association analysis of bipolar disorder identifies a new susceptibility locus near ODZ4. <i>Nature Genetics</i> , <b>2011</b> , 43, 977-83	36.3	1094
55	Lack of allelic association between markers at the DRD2 and ANKK1 gene loci with the alcohol-dependence syndrome and criminal activity. <i>Psychiatric Genetics</i> , <b>2011</b> , 21, 323-4	2.9	5
54	Confirmation of prior evidence of genetic susceptibility to alcoholism in a genome-wide association study of comorbid alcoholism and bipolar disorder. <i>Psychiatric Genetics</i> , <b>2011</b> , 21, 294-306	2.9	48
53	A threonine to isoleucine missense mutation in the pericentriolar material 1 gene is strongly associated with schizophrenia. <i>Molecular Psychiatry</i> , <b>2010</b> , 15, 615-28	15.1	45
52	Genetic evidence implicates the immune system and cholesterol metabolism in the aetiology of Alzheimer's disease. <i>PLoS ONE</i> , <b>2010</b> , 5, e13950	3.7	276
51	Genetic power of a Brazilian three-generation family with generalized aggressive periodontitis. <i>Brazilian Dental Journal</i> , <b>2010</b> , 21, 137-41	1.9	5
50	NK1 (TACR1) receptor gene knockout mouse phenotype predicts genetic association with ADHD. <i>Journal of Psychopharmacology</i> , <b>2010</b> , 24, 27-38	4.6	294
49	Genome-wide association study of suicide attempts in mood disorder patients. <i>American Journal of Psychiatry</i> , <b>2010</b> , 167, 1499-507	11.9	113
48	Support of association between BRD1 and both schizophrenia and bipolar affective disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2010</b> , 153B, 582-591	3.5	42
47	Effect of the 2004 tsunami on suicide rates in Sri Lanka. <i>Psychiatric Bulletin</i> , <b>2009</b> , 33, 179-180		12
46	A genomewide association study of response to lithium for prevention of recurrence in bipolar disorder. <i>American Journal of Psychiatry</i> , <b>2009</b> , 166, 718-25	11.9	121
45	Meta-analysis of 32 genome-wide linkage studies of schizophrenia. <i>Molecular Psychiatry</i> , <b>2009</b> , 14, 774-85	5.1	202
44	DISC1 association, heterogeneity and interplay in schizophrenia and bipolar disorder. <i>Molecular Psychiatry</i> , <b>2009</b> , 14, 865-73	15.1	127
43	Case-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. <i>Molecular Psychiatry</i> , <b>2009</b> , 14, 614-20	15.1	92



42	Common polygenic variation contributes to risk of schizophrenia and bipolar disorder. <i>Nature</i> , <b>2009</b> , 460, 748-52	50.4	3568
41	Genome-wide association study identifies variants at CLU and PICALM associated with Alzheimer's disease. <i>Nature Genetics</i> , <b>2009</b> , 41, 1088-93	36.3	2018
40	Genetics of attention-deficit hyperactivity disorder (ADHD). <i>Neuropharmacology</i> , <b>2009</b> , 57, 590-600	5.5	100
39	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. <i>Behavioral and Brain Functions</i> , <b>2009</b> , 5, 28	4.1	34
38	No evidence for excess runs of homozygosity in bipolar disorder. <i>Psychiatric Genetics</i> , <b>2009</b> , 19, 165-70	2.9	34
37	Confirmation of the genetic association between the U2AF homology motif (UHM) kinase 1 (UHMK1) gene and schizophrenia on chromosome 1q23.3. <i>European Journal of Human Genetics</i> , <b>2008</b> , 16, 1275-82	5.3	13
36	Rare chromosomal deletions and duplications increase risk of schizophrenia. <i>Nature</i> , <b>2008</b> , 455, 237-41	50.4	1251
35	Collaborative genome-wide association analysis supports a role for ANK3 and CACNA1C in bipolar disorder. <i>Nature Genetics</i> , <b>2008</b> , 40, 1056-8	36.3	949
34	Whole-genome association study of bipolar disorder. <i>Molecular Psychiatry</i> , <b>2008</b> , 13, 558-69	15.1	571
33	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. <i>Pharmacogenetics and Genomics</i> , <b>2007</b> , 17, 605-17	1.9	117
32	Replication of genetic association studies between markers at the Epsin 4 gene locus and schizophrenia in two Han Chinese samples. <i>Schizophrenia Research</i> , <b>2007</b> , 89, 357-9	3.6	9
31	Failure to confirm allelic and haplotypic association between markers at the chromosome 6p22.3 dystrobrevin-binding protein 1 (DTNBP1) locus and schizophrenia. <i>Behavioral and Brain Functions</i> , <b>2007</b> , 3, 50	4.1	18
30	Fine mapping by genetic association implicates the chromosome 1q23.3 gene UHMK1, encoding a serine/threonine protein kinase, as a novel schizophrenia susceptibility gene. <i>Biological Psychiatry</i> , <b>2007</b> , 61, 873-9	7.9	30
29	GeneBrain Structure Relationships: Arbitrary Assumptions of Heterogeneity Generate Unfalsifiable ClaimsReply. <i>Archives of General Psychiatry</i> , <b>2007</b> , 64, 1098		
28	A genetic association study of chromosome 11q22-24 in two different samples implicates the FXRD6 gene, encoding phosphohippolin, in susceptibility to schizophrenia. <i>American Journal of Human Genetics</i> , <b>2007</b> , 80, 664-72	11	26
27	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). <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2006</b> , 141B, 296-300	3.5	30
26	Failure to confirm allelic association between markers at the CAPON gene locus and schizophrenia in a British sample. <i>Biological Psychiatry</i> , <b>2006</b> , 59, 195-7	7.9	27
25	Genetic association and brain morphology studies and the chromosome 8p22 pericentriolar material 1 (PCM1) gene in susceptibility to schizophrenia. <i>Archives of General Psychiatry</i> , <b>2006</b> , 63, 844-54		77

24	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. <i>American Journal of Psychiatry</i> , <b>2006</b> , 163, 1767-76	11.9	11
23	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. <i>Molecular Psychiatry</i> , <b>2006</b> , 11, 134-42	15.1	70
22	The Epsin 4 gene on chromosome 5q, which encodes the clathrin-associated protein enthoprotin, is involved in the genetic susceptibility to schizophrenia. <i>American Journal of Human Genetics</i> , <b>2005</b> , 76, 902-7	11	58
21	Direct genomic PCR sequencing of the high affinity thiamine transporter (SLC19A2) gene identifies three genetic variants in Wernicke Korsakoff syndrome (WKS). <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2005</b> , 137B, 17-9	3.5	18
20	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. <i>BMC Genetics</i> , <b>2005</b> , 6, 11	2.6	32
19	No association between a neuronal nitric oxide synthase (NOS1) gene polymorphism on chromosome 12q24 and bipolar disorder <b>2004</b> , 124B, 73-5		17
18	Genome scan of Tourette syndrome in a single large pedigree shows some support for linkage to regions of chromosomes 5, 10 and 13. <i>Psychiatric Genetics</i> , <b>2004</b> , 14, 83-7	2.9	39
17	. <i>Psychiatric Genetics</i> , <b>2003</b> , 13, 77-84	2.9	15
16	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. <i>Psychiatric Genetics</i> , <b>2003</b> , 13, 77-84	2.9	130
15	Genome scan meta-analysis of schizophrenia and bipolar disorder, part III: Bipolar disorder. <i>American Journal of Human Genetics</i> , <b>2003</b> , 73, 49-62	11	353
14	Genome scan meta-analysis of schizophrenia and bipolar disorder, part II: Schizophrenia. <i>American Journal of Human Genetics</i> , <b>2003</b> , 73, 34-48	11	985
13	Linkage disequilibrium and demographic history of the isolated population of the Faroe Islands. <i>European Journal of Human Genetics</i> , <b>2002</b> , 10, 381-7	5.3	33
12	A novel polymorphism in exon 11 of the WKL1 gene, shows no association with schizophrenia. <i>European Journal of Human Genetics</i> , <b>2002</b> , 10, 491-4	5.3	12
11	Genetic association studies of schizophrenia using the 8p21-22 genes: prepronociceptin (PNOC), neuronal nicotinic cholinergic receptor alpha polypeptide 2 (CHRNA2) and arylamine N-acetyltransferase 1 (NAT1). <i>European Journal of Human Genetics</i> , <b>2001</b> , 9, 469-72	5.3	28
10	Genomewide genetic linkage analysis confirms the presence of susceptibility loci for schizophrenia, on chromosomes 1q32.2, 5q33.2, and 8p21-22 and provides support for linkage to schizophrenia, on chromosomes 11q23.3-24 and 20q12.1-11.23. <i>American Journal of Human Genetics</i> , <b>2001</b> , 68, 661-73	11	337
9	Optimization of liposome mediated transfection of a neuronal cell line. <i>NeuroReport</i> , <b>1997</b> , 8, 1481-4	1.7	25
8	The genetics of schizophrenia230-261		1
7	Genome-wide association study of suicide attempt in psychiatric disorders identifies association with major depression polygenic risk scores		2

6	Identifying risk factors involved in the common versus specific liabilities to substance abuse: A genetically informed approach	1
5	Genome-wide Association Study of Clinical Features in the Schizophrenia Psychiatric Genomics Consortium: Confirmation of Polygenic Effect on Negative Symptoms	3
4	Genome-wide association study identifies 30 Loci Associated with Bipolar Disorder	28
3	Dissecting the shared genetic architecture of suicide attempt, psychiatric disorders and known risk factors	2
2	Meta-analysis of problematic alcohol use in 435,563 individuals identifies 29 risk variants and yields insights into biology, pleiotropy and causality	6
1	Characterization of Age and Polarity at Onset in Bipolar Disorder	1