Andrew McQuillin

List of Publications by Citations

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	Biological insights from 108 schizophrenia-associated genetic loci. *Nature*, **2014**, 511, 421-7	50.4	5249
166	Common polygenic variation contributes to risk of schizophrenia and bipolar disorder. *Nature*, **2009**, 460, 748-52	50.4	3568
165	Genome-wide association study identifies variants at CLU and PICALM associated with Alzheimer's disease. *Nature Genetics*, **2009**, 41, 1088-93	36.3	2018
164	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. *Nature Genetics*, **2013**, 45, 984-94	36.3	1628
163	Genome-wide association study identifies five new schizophrenia loci. *Nature Genetics*, **2011**, 43, 969-76	36.3	1508
162	Common variants at ABCA7, MS4A6A/MS4A4E, EPHA1, CD33 and CD2AP are associated with Alzheimer's disease. *Nature Genetics*, **2011**, 43, 429-35	36.3	1421
161	Rare chromosomal deletions and duplications increase risk of schizophrenia. *Nature*, **2008**, 455, 237-41	50.4	1251
160	Large-scale genome-wide association analysis of bipolar disorder identifies a new susceptibility locus near ODZ4. *Nature Genetics*, **2011**, 43, 977-83	36.3	1094
159	Genome scan meta-analysis of schizophrenia and bipolar disorder, part II: Schizophrenia. *American Journal of Human Genetics*, **2003**, 73, 34-48	11	985
158	Collaborative genome-wide association analysis supports a role for ANK3 and CACNA1C in bipolar disorder. *Nature Genetics*, **2008**, 40, 1056-8	36.3	949
157	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates A□ tau, immunity and lipid processing. *Nature Genetics*, **2019**, 51, 414-430	36.3	917
156	The UK10K project identifies rare variants in health and disease. *Nature*, **2015**, 526, 82-90	50.4	776
155	Analysis of shared heritability in common disorders of the brain. *Science*, **2018**, 360,	33.3	666
154	Genome-wide association study identifies 30 loci associated with bipolar disorder. *Nature Genetics*, **2019**, 51, 793-803	36.3	662
153	Modeling Linkage Disequilibrium Increases Accuracy of Polygenic Risk Scores. *American Journal of Human Genetics*, **2015**, 97, 576-92	11	649
152	Psychiatric genome-wide association study analyses implicate neuronal, immune and histone pathways. *Nature Neuroscience*, **2015**, 18, 199-209	25.5	572
151	Whole-genome association study of bipolar disorder. *Molecular Psychiatry*, **2008**, 13, 558-69	15.1	571

150	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. *Nature Genetics*, **2017**, 49, 27-35	36.3	530
149	Partitioning heritability of regulatory and cell-type-specific variants across 11 common diseases. *American Journal of Human Genetics*, **2014**, 95, 535-52	11	411
148	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. *Cell*, **2019**, 179, 1469-1482.e11	56.2	402
147	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. *Cell*, **2018**, 173, 1705-1715.e16	56.2	360
146	Genome scan meta-analysis of schizophrenia and bipolar disorder, part III: Bipolar disorder. *American Journal of Human Genetics*, **2003**, 73, 49-62	11	353
145	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. *American Journal of Human Genetics*, **2001**, 68, 661-73	11	337
144	A genome-wide association study confirms PNPLA3 and identifies TM6SF2 and MBOAT7 as risk loci for alcohol-related cirrhosis. *Nature Genetics*, **2015**, 47, 1443-8	36.3	303
143	NK1 (TACR1) receptor gene "knockout" mouse phenotype predicts genetic association with ADHD. *Journal of Psychopharmacology*, **2010**, 24, 27-38	4.6	294
142	Rare loss-of-function variants in SETD1A are associated with schizophrenia and developmental disorders. *Nature Neuroscience*, **2016**, 19, 571-7	25.5	284
141	Genetic evidence implicates the immune system and cholesterol metabolism in the aetiology of Alzheimer's disease. *PLoS ONE*, **2010**, 5, e13950	3.7	276
140	Polygenic dissection of diagnosis and clinical dimensions of bipolar disorder and schizophrenia. *Molecular Psychiatry*, **2014**, 19, 1017-1024	15.1	258
139	Meta-analysis of 32 genome-wide linkage studies of schizophrenia. *Molecular Psychiatry*, **2009**, 14, 774-85	15.1	202
138	An integrated genetic-epigenetic analysis of schizophrenia: evidence for co-localization of genetic associations and differential DNA methylation. *Genome Biology*, **2016**, 17, 176	18.3	189
137	A novel Alzheimer disease locus located near the gene encoding tau protein. *Molecular Psychiatry*, **2016**, 21, 108-17	15.1	175
136	Genetic influences on schizophrenia and subcortical brain volumes: large-scale proof of concept. *Nature Neuroscience*, **2016**, 19, 420-431	25.5	163
135	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, 283-94	11	161
134	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	2.9	130
133	DISC1 association, heterogeneity and interplay in schizophrenia and bipolar disorder. *Molecular Psychiatry*, **2009**, 14, 865-73	15.1	127

132	A genomewide association study of response to lithium for prevention of recurrence in bipolar disorder. *American Journal of Psychiatry*, **2009**, 166, 718-25	11.9	121
131	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, 605-17	1.9	117
130	Genome-wide association study of suicide attempts in mood disorder patients. *American Journal of Psychiatry*, **2010**, 167, 1499-507	11.9	113
129	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	11	104
128	GWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. *American Journal of Psychiatry*, **2019**, 176, 651-660	11.9	103
127	Genome-wide association study of borderline personality disorder reveals genetic overlap with bipolar disorder, major depression and schizophrenia. *Translational Psychiatry*, **2017**, 7, e1155	8.6	100
126	Genetics of attention-deficit hyperactivity disorder (ADHD). *Neuropharmacology*, **2009**, 57, 590-600	5.5	100
125	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. *Molecular Psychiatry*, **2009**, 14, 614-20	15.1	92
124	Genome-wide association study of Alzheimer's disease with psychotic symptoms. *Molecular Psychiatry*, **2012**, 17, 1316-27	15.1	90
123	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	85
122	Maternally derived microduplications at 15q11-q13: implication of imprinted genes in psychotic illness. *American Journal of Psychiatry*, **2011**, 168, 408-17	11.9	84
121	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	36.3	83
120	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, 844-54		77
119	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, 744-53	11	72
118	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, 134-42	15.1	70
117	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	25.5	69
116	Genetic Overlap Between Attention-Deficit/Hyperactivity Disorder and Bipolar Disorder: Evidence From Genome-wide Association Study Meta-analysis. *Biological Psychiatry*, **2017**, 82, 634-641	7.9	66
115	Heterozygous carriage of the alpha1-antitrypsin Pi*Z variant increases the risk to develop liver cirrhosis. *Gut*, **2019**, 68, 1099-1107	19.2	62

114	Case-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	2.9	58
113	GWA study data mining and independent replication identify cardiomyopathy-associated 5 (CMYA5) as a risk gene for schizophrenia. *Molecular Psychiatry*, **2011**, 16, 1117-29	15.1	58
112	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, 902-7	11	58
111	Polygenic dissection of the bipolar phenotype. *British Journal of Psychiatry*, **2011**, 198, 284-8	5.4	57
110	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	7.9	57
109	Estimation of Genetic Correlation via Linkage Disequilibrium Score Regression and Genomic Restricted Maximum Likelihood. *American Journal of Human Genetics*, **2018**, 102, 1185-1194	11	55
108	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	0.7	50
107	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	2.9	48
106	The role of variation at APP, PSEN1, PSEN2, and MAPT in late onset Alzheimer's disease. *Journal of Alzheimer's Disease*, **2012**, 28, 377-87	4.3	47
105	Genetic Variation in HSD17B13 Reduces the Risk of Developing Cirrhosis and Hepatocellular Carcinoma in Alcohol Misusers. *Hepatology*, **2020**, 72, 88-102	11.2	46
104	A threonine to isoleucine missense mutation in the pericentriolar material 1 gene is strongly associated with schizophrenia. *Molecular Psychiatry*, **2010**, 15, 615-28	15.1	45
103	New data and an old puzzle: the negative association between schizophrenia and rheumatoid arthritis. *International Journal of Epidemiology*, **2015**, 44, 1706-21	7.8	43
102	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	3.5	42
101	Evidence for Genetic Overlap Between Schizophrenia and Age at First Birth in Women. *JAMA Psychiatry*, **2016**, 73, 497-505	14.5	40
100	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	3.5	39
99	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, 83-7	2.9	39
98	Mutations in the Gabrb1 gene promote alcohol consumption through increased tonic inhibition. *Nature Communications*, **2013**, 4, 2816	17.4	37
97	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	13.4	36

96	Analysis of ANK3 and CACNA1C variants identified in bipolar disorder whole genome sequence data. *Bipolar Disorders*, **2014**, 16, 583-91	3.8	36
95	Analysis of genetic deletions and duplications in the University College London bipolar disorder case control sample. *European Journal of Human Genetics*, **2011**, 19, 588-92	5.3	35
94	Mapping genomic loci implicates genes and synaptic biology in schizophrenia.. *Nature*, **2022**,	50.4	35
93	Schizophrenia genetic variants are not associated with intelligence. *Psychological Medicine*, **2013**, 43, 2563-70	6.9	34
92	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	4.1	34
91	No evidence for excess runs of homozygosity in bipolar disorder. *Psychiatric Genetics*, **2009**, 19, 165-70	2.9	34
90	Association of rare variation in the glutamate receptor gene SLC1A2 with susceptibility to bipolar disorder and schizophrenia. *European Journal of Human Genetics*, **2015**, 23, 1200-6	5.3	33
89	Linkage disequilibrium and demographic history of the isolated population of the Faroe Islands. *European Journal of Human Genetics*, **2002**, 10, 381-7	5.3	33
88	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**, 165B, 373-80	3.5	32
87	An inherited duplication at the gene p21 Protein-Activated Kinase 7 (PAK7) is a risk factor for psychosis. *Human Molecular Genetics*, **2014**, 23, 3316-26	5.6	32
86	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	2.6	32
85	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, 873-9	7.9	30
84	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	3.5	30
83	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, 591-8	14.5	28
82	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, 1218-30	4.6	28
81	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). *European Journal of Human Genetics*, **2001**, 9, 469-72	5.3	28
80	Genome-wide association study identifies 30 Loci Associated with Bipolar Disorder		28
79	Genetic association study of GABRA2 single nucleotide polymorphisms and electroencephalography in alcohol dependence. *Neuroscience Letters*, **2011**, 500, 162-6	3.3	27

78	Failure to confirm allelic association between markers at the CAPON gene locus and schizophrenia in a British sample. *Biological Psychiatry*, **2006**, 59, 195-7	7.9	27
77	The functional GRM3 Kozak sequence variant rs148754219 affects the risk of schizophrenia and alcohol dependence as well as bipolar disorder. *Psychiatric Genetics*, **2014**, 24, 277-8	2.9	26
76	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, 664-72	11	26
75	Optimization of liposome mediated transfection of a neuronal cell line. *NeuroReport*, **1997**, 8, 1481-4	1.7	25
74	Chromosomal microarray testing in adults with intellectual disability presenting with comorbid psychiatric disorders. *European Journal of Human Genetics*, **2016**, 25, 66-72	5.3	24
73	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**, 165B, 365-72	3.5	24
72	Genome-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** , 171B, 276-89	3.5	23
71	Genetic variants in or near ADH1B and ADH1C affect susceptibility to alcohol dependence in a British and Irish population. *Addiction Biology*, **2015**, 20, 594-604	4.6	22
70	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, 340-6	3.6	22
69	Genetic association and sequencing of the insulin-like growth factor 1 gene in bipolar affective disorder. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2011**, 156, 177-87	3.5	22
68	Neurodevelopmental risk copy number variants in adults with intellectual disabilities and comorbid psychiatric disorders. *British Journal of Psychiatry*, **2018**, 212, 287-294	5.4	21
67	Use of schizophrenia and bipolar disorder polygenic risk scores to identify psychotic disorders. *British Journal of Psychiatry*, **2018**, 213, 535-541	5.4	21
66	Genome-Wide Association Study for Alcohol-Related Cirrhosis Identifies Risk Loci in MARC1 and HNRNPUL1. *Gastroenterology*, **2020**, 159, 1276-1289.e7	13.3	19
65	Whole-exome sequencing of individuals from an isolated population implicates rare risk variants in bipolar disorder. *Translational Psychiatry*, **2017**, 7, e1034	8.6	18
64	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	4.1	18
63	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, 17-9	3.5	18
62	Timing of prenatal exposure to trauma and altered placental expressions of hypothalamic-pituitary-adrenal axis genes and genes driving neurodevelopment. *Journal of Neuroendocrinology*, **2018**, 30, e12581	3.8	17
61	Memory decline in Down syndrome and its relationship to iPF2alpha, a urinary marker of oxidative stress. *PLoS ONE*, **2014**, 9, e97709	3.7	17

60	No association between a neuronal nitric oxide synthase (NOS1) gene polymorphism on chromosome 12q24 and bipolar disorder **2004**, 124B, 73-5		17
59	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	3.5	16
58	Rare coding variants in ten genes confer substantial risk for schizophrenia.. *Nature*, **2022**,	50.4	16
57	No evidence that extended tracts of homozygosity are associated with Alzheimer's disease. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2011**, 156B, 764-71	3.5	15
56	. *Psychiatric Genetics*, **2003**, 13, 77-84	2.9	15
55	DNA methylation meta-analysis reveals cellular alterations in psychosis and markers of treatment-resistant schizophrenia. *ELife*, **2021**, 10,	8.9	15
54	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. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2012**, 159B, 328-35	3.5	14
53	Hypomethylation of FAM63B in bipolar disorder patients. *Clinical Epigenetics*, **2016**, 8, 52	7.7	14
52	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, 205-11	3.8	13
51	A nonconservative amino acid change in the UPF3B gene in a patient with schizophrenia. *Psychiatric Genetics*, **2012**, 22, 150-1	2.9	13
50	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, 1275-82	5.3	13
49	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	2.2	13
48	Effect of the 2004 tsunami on suicide rates in Sri Lanka. *Psychiatric Bulletin*, **2009**, 33, 179-180		12
47	A novel polymorphism in exon 11 of the WKL1 gene, shows no association with schizophrenia. *European Journal of Human Genetics*, **2002**, 10, 491-4	5.3	12
46	Genetic testing in intellectual disability psychiatry: Opinions and practices of UK child and intellectual disability psychiatrists. *Journal of Applied Research in Intellectual Disabilities*, **2018**, 31, 273-284	2.2	11
45	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, 1767-76	11.9	11
44	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. *Biological Psychiatry*, **2022**, 91, 102-117	7.9	11
43	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. *Biological Psychiatry*, **2021**,	7.9	11

42	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	3.5	10
41	Genetic variation in the miR-708 gene and its binding targets in bipolar disorder. *Bipolar Disorders*, **2016**, 18, 650-656	3.8	10
40	Replication of genetic association studies between markers at the Epsin 4 gene locus and schizophrenia in two Han Chinese samples. *Schizophrenia Research*, **2007**, 89, 357-9	3.6	9
39	Rare variant analysis in multiply affected families, association studies and functional analysis suggest a role for the ITG4 gene in schizophrenia and bipolar disorder. *Schizophrenia Research*, **2018**, 199, 181-188	3.6	8
38	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, 201-9	3.5	8
37	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, 153-62	2.9	7
36	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	6.6	7
35	Meta-analysis of problematic alcohol use in 435,563 individuals identifies 29 risk variants and yields insights into biology, pleiotropy and causality		6
34	Genetic variation in GABR1 and the risk for developing alcohol dependence. *Psychiatric Genetics*, **2017**, 27, 110-115	2.9	5
33	Phenotypic heterogeneity in study populations may significantly confound the results of genetic association studies on alcohol dependence. *Psychiatric Genetics*, **2015**, 25, 234-40	2.9	5
32	Genetic power of a Brazilian three-generation family with generalized aggressive periodontitis. *Brazilian Dental Journal*, **2010**, 21, 137-41	1.9	5
31	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, 323-4	2.9	5
30	Genetic variants in ALDH1B1 and alcohol dependence risk in a British and Irish population: A bioinformatic and genetic study. *PLoS ONE*, **2017**, 12, e0177009	3.7	5
29	Genetic copy number variants, cognition and psychosis: a meta-analysis and a family study. *Molecular Psychiatry*, **2021**, 26, 5307-5319	15.1	5
28	Identification of rare nonsynonymous variants in SYNE1/CPG2 in bipolar affective disorder. *Psychiatric Genetics*, **2017**, 27, 81-88	2.9	4
27	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	13.4	4
26	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	2.4	4
25	A genetic risk score and diabetes predict development of alcohol-related cirrhosis in drinkers. *Journal of Hepatology*, **2021**,	13.4	4

24	Identifying risk factors involved in the common versus specific liabilities to substance use: A genetically informed approach. *Addiction Biology*, **2021**, 26, e12944	4.6	4
23	The Influence of CYP2D6 and CYP2C19 Genetic Variation on Diabetes Mellitus Risk in People Taking Antidepressants and Antipsychotics. *Genes*, **2021**, 12,	4.2	3
22	Genome-wide Association Study of Clinical Features in the Schizophrenia Psychiatric Genomics Consortium: Confirmation of Polygenic Effect on Negative Symptoms		3
21	A machine learning case-control classifier for schizophrenia based on DNA methylation in blood. *Translational Psychiatry*, **2021**, 11, 412	8.6	3
20	Investigating rare pathogenic/likely pathogenic exonic variation in bipolar disorder. *Molecular Psychiatry*, **2021**, 26, 5239-5250	15.1	3
19	People who survive an episode of severe alcoholic hepatitis should be advised to maintain total abstinence from alcohol. *Hepatology*, **2018**, 67, 2479-2480	11.2	2
18	The Communication of Metacognition for Social Strategy in Psychosis: An Exploratory Study. *Schizophrenia Bulletin Open*, **2020**, 1,	2.2	2
17	Genome-wide association study of suicide attempt in psychiatric disorders identifies association with major depression polygenic risk scores		2
16	Dissecting the shared genetic architecture of suicide attempt, psychiatric disorders and known risk factors		2
15	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". *Journal of Hepatology*, **2018**, 68, 860-862	13.4	2
14	Characterisation of age and polarity at onset in bipolar disorder.. *British Journal of Psychiatry*, **2021**, 219, 659-669	5.4	2
13	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	3.5	1
12	Evidence for genetic susceptibility to the alcohol dependence syndrome from the thiamine transporter 2 gene solute carrier SLC19A3. *Psychiatric Genetics*, **2014**, 24, 122-3	2.9	1
11	The genetics of schizophrenia230-261		1
10	Adolescent Verbal Memory as a Psychosis Endophenotype: A Genome-Wide Association Study in an Ancestrally Diverse Sample.. *Genes*, **2022**, 13,	4.2	1
9	Physical Health and Clinical Phenotypes **2019**, 71-86		1
8	Identifying risk factors involved in the common versus specific liabilities to substance abuse: A genetically informed approach		1
7	Characterization of Age and Polarity at Onset in Bipolar Disorder		1

6	Genetic variant analysis of the putative regulatory regions of the LRRC7 gene in bipolar disorder. *Psychiatric Genetics*, **2016**, 26, 99-100	2.9	1
5	Elevated common variant genetic risk for tourette syndrome in a densely-affected pedigree. *Molecular Psychiatry*, **2021**,	15.1	1
4	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	2.9	0
3	GeneBrain Structure Relationships: Arbitrary Assumptions of Heterogeneity Generate Unfalsifiable ClaimsReply. *Archives of General Psychiatry*, **2007**, 64, 1098		
2	Association study of rare nonsynonymous variants of FTO in bipolar disorder. *Psychiatric Genetics*, **2016**, 26, 140-1	2.9	
1	P: 81 Previously Identified Candidate Gene Associations in Hepatic Encephalopathy Do Not Replicate in the STOPAH Cohort. *American Journal of Gastroenterology*, **2019**, 114, S39-S39	0.7	