Andrew McQuillin

List of Publications by Year
in descending order

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173 papers	46,581 citations	22548 61 h-index	6177 164 g-index
206 all docs	206 docs citations	206 times ranked	46190 citing authors

#	Article	IF	Citations
1	Biological insights from 108 schizophrenia-associated genetic loci. Nature, 2014, 511, 421-427.	13.7	6,934
2	Common polygenic variation contributes to risk of schizophrenia and bipolar disorder. Nature, 2009, 460, 748-752.	13.7	4,345
3	Genome-wide association study identifies variants at CLU and PICALM associated with Alzheimer's disease. Nature Genetics, 2009, 41, 1088-1093.	9.4	2,697
4	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. Nature Genetics, 2013, 45, 984-994.	9.4	2,067
5	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.	9.4	1,962
6	Genome-wide association study identifies five new schizophrenia loci. Nature Genetics, 2011, 43, 969-976.	9.4	1,758
7	Common variants at ABCA7, MS4A6A/MS4A4E, EPHA1, CD33 and CD2AP are associated with Alzheimer's disease. Nature Genetics, 2011, 43, 429-435.	9.4	1,708
8	Rare chromosomal deletions and duplications increase risk of schizophrenia. Nature, 2008, 455, 237-241.	13.7	1,387
9	Large-scale genome-wide association analysis of bipolar disorder identifies a new susceptibility locus near ODZ4. Nature Genetics, 2011, 43, 977-983.	9.4	1,283
10	Genome-wide association study identifies 30 loci associated with bipolar disorder. Nature Genetics, 2019, 51, 793-803.	9.4	1,191
11	Collaborative genome-wide association analysis supports a role for ANK3 and CACNA1C in bipolar disorder. Nature Genetics, 2008, 40, 1056-1058.	9.4	1,102
12	Modeling Linkage Disequilibrium Increases Accuracy of Polygenic Risk Scores. American Journal of Human Genetics, 2015, 97, 576-592.	2.6	1,098
13	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	6.0	1,085
14	Genome Scan Meta-Analysis of Schizophrenia and Bipolar Disorder, Part II: Schizophrenia. American Journal of Human Genetics, 2003, 73, 34-48.	2.6	1,072
15	The UK10K project identifies rare variants in health and disease. Nature, 2015, 526, 82-90.	13.7	1,014
16	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. Cell, 2019, 179, 1469-1482.e11.	13.5	935
17	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	13.7	929
18	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. Nature Genetics, 2017, 49, 27-35.	9.4	838

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19	Psychiatric genome-wide association study analyses implicate neuronal, immune and histone pathways. Nature Neuroscience, 2015, 18, 199-209.	7.1	701
20	Whole-genome association study of bipolar disorder. Molecular Psychiatry, 2008, 13, 558-569.	4.1	642
21	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.	9.4	629
22	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. Cell, 2018, 173, 1705-1715.e16.	13.5	623
23	Partitioning Heritability of Regulatory and Cell-Type-Specific Variants across 11 Common Diseases. American Journal of Human Genetics, 2014, 95, 535-552.	2.6	569
24	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.	9.4	435
25	Genome Scan Meta-Analysis of Schizophrenia and Bipolar Disorder, Part III: Bipolar Disorder. American Journal of Human Genetics, 2003, 73, 49-62.	2.6	400
26	Rare loss-of-function variants in SETD1A are associated with schizophrenia and developmental disorders. Nature Neuroscience, 2016, 19, 571-577.	7.1	388
27	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, 661-673.	2.6	362
28	Genetic Evidence Implicates the Immune System and Cholesterol Metabolism in the Aetiology of Alzheimer's Disease. PLoS ONE, 2010, 5, e13950.	1.1	347
29	Polygenic dissection of diagnosis and clinical dimensions of bipolar disorder and schizophrenia. Molecular Psychiatry, 2014, 19, 1017-1024.	4.1	333
30	NK₁ (TACR₁) receptor gene â€˜knockoutâ€™ mouse phenotype predicts genetic association with ADHD. Journal of Psychopharmacology, 2010, 24, 27-38.	2.0	329
31	Rare coding variants in ten genes confer substantial risk for schizophrenia. Nature, 2022, 604, 509-516.	13.7	326
32	An integrated genetic-epigenetic analysis of schizophrenia: evidence for co-localization of genetic associations and differential DNA methylation. Genome Biology, 2016, 17, 176.	3.8	287
33	A novel Alzheimer disease locus located near the gene encoding tau protein. Molecular Psychiatry, 2016, 21, 108-117.	4.1	260
34	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.	7.1	242
35	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.	2.6	237
36	Meta-analysis of 32 genome-wide linkage studies of schizophrenia. Molecular Psychiatry, 2009, 14, 774-785.	4.1	235

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37	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.	2.6	225
38	Genetic influences on schizophrenia and subcortical brain volumes: large-scale proof of concept. Nature Neuroscience, 2016, 19, 420-431.	7.1	204
39	GWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. American Journal of Psychiatry, 2019, 176, 651-660.	4.0	186
40	Genome-wide association study of borderline personality disorder reveals genetic overlap with bipolar disorder, major depression and schizophrenia. Translational Psychiatry, 2017, 7, e1155-e1155.	2.4	150
41	A Genomewide Association Study of Response to Lithium for Prevention of Recurrence in Bipolar Disorder. American Journal of Psychiatry, 2009, 166, 718-725.	4.0	145
42	DISC1 association, heterogeneity and interplay in schizophrenia and bipolar disorder. Molecular Psychiatry, 2009, 14, 865-873.	4.1	140
43	Genome-Wide Association Study of Suicide Attempts in Mood Disorder Patients. American Journal of Psychiatry, 2010, 167, 1499-1507.	4.0	140
44	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.	0.7	137
45	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.	0.6	136
46	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.	0.7	133
47	Estimation of Genetic Correlation via Linkage Disequilibrium Score Regression and Genomic Restricted Maximum Likelihood. American Journal of Human Genetics, 2018, 102, 1185-1194.	2.6	119
48	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. Biological Psychiatry, 2022, 91, 313-327.	0.7	114
49	Genetics of attention-deficit hyperactivity disorder (ADHD). Neuropharmacology, 2009, 57, 590-600.	2.0	113
50	Genome-wide association study of Alzheimer's disease with psychotic symptoms. Molecular Psychiatry, 2012, 17, 1316-1327.	4.1	110
51	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
52	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.	4.1	101
53	Heterozygous carriage of the alpha1-antitrypsin Pi*Z variant increases the risk to develop liver cirrhosis. Gut, 2019, 68, 1099-1107.	6.1	100
54	Genetic Overlap Between Attention-Deficit/Hyperactivity Disorder and Bipolar Disorder: Evidence From Genome-wide Association Study Meta-analysis. Biological Psychiatry, 2017, 82, 634-641.	0.7	99

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55	Maternally Derived Microduplications at 15q11-q13: Implication of Imprinted Genes in Psychotic Illness. American Journal of Psychiatry, 2011, 168, 408-417.	4.0	95
56	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.	2.6	91
57	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.	13.8	82
58	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.	0.2	82
59	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.	4.1	81
60	Genetic Variation in HSD17B13 Reduces the Risk of Developing Cirrhosis and Hepatocellular Carcinoma in Alcohol Misusers. Hepatology, 2020, 72, 88-102.	3.6	76
61	DNA methylation meta-analysis reveals cellular alterations in psychosis and markers of treatment-resistant schizophrenia. ELife, 2021, 10, .	2.8	72
62	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.	0.6	70
63	GWA study data mining and independent replication identify cardiomyopathy-associated 5 (CMYA5) as a risk gene for schizophrenia. Molecular Psychiatry, 2011, 16, 1117-1129.	4.1	67
64	Polygenic dissection of the bipolar phenotype. British Journal of Psychiatry, 2011, 198, 284-288.	1.7	67
65	Exome sequencing in bipolar disorder identifies AKAP11 as a risk gene shared with schizophrenia. Nature Genetics, 2022, 54, 541-547.	9.4	65
66	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.	2.6	62
67	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. Biological Psychiatry, 2022, 91, 102-117.	0.7	61
68	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.	0.6	59
69	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.	1.1	57
70	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.	1.2	53
71	New data and an old puzzle: the negative association between schizophrenia and rheumatoid arthritis. International Journal of Epidemiology, 2015, 44, 1706-1721.	0.9	53
72	Genome-Wide Association Study for Alcohol-Related Cirrhosis Identifies Risk Loci in MARC1 and HNRNPUL1. Gastroenterology, 2020, 159, 1276-1289.e7.	0.6	53

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73	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.	1.8	52
74	Evidence for Genetic Overlap Between Schizophrenia and Age at First Birth in Women. JAMA Psychiatry, 2016, 73, 497.	6.0	51
75	A threonine to isoleucine missense mutation in the pericentriolar material 1 gene is strongly associated with schizophrenia. Molecular Psychiatry, 2010, 15, 615-628.	4.1	50
76	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.	1.1	47
77	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.	1.4	45
78	Mutations in the Gabrb1 gene promote alcohol consumption through increased tonic inhibition. Nature Communications, 2013, 4, 2816.	5.8	44
79	Analysis of ANK3 and CACNA1C variants identified in bipolar disorder whole genome sequence data. Bipolar Disorders, 2014, 16, 583-591.	1.1	44
80	Interaction Testing and Polygenic Risk Scoring to Estimate the Association of Common Genetic Variants With Treatment Resistance in Schizophrenia. JAMA Psychiatry, 2022, 79, 260.	6.0	44
81	Linkage disequilibrium and demographic history of the isolated population of the Faroe Islands. European Journal of Human Genetics, 2002, 10, 381-387.	1.4	43
82	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.	6.0	43
83	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.	0.6	41
84	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.	1.4	40
85	Schizophrenia genetic variants are not associated with intelligence. Psychological Medicine, 2013, 43, 2563-2570.	2.7	40
86	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.	1.1	39
87	Genetic association study of GABRA2 single nucleotide polymorphisms and electroencephalography in alcohol dependence. Neuroscience Letters, 2011, 500, 162-166.	1.0	38
88	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.	1.4	38
89	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.	1.4	37
90	Use of schizophrenia and bipolar disorder polygenic risk scores to identify psychotic disorders. British Journal of Psychiatry, 2018, 213, 535-541.	1.7	37

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91	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
92	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.	0.7	35
93	No evidence for excess runs of homozygosity in bipolar disorder. Psychiatric Genetics, 2009, 19, 165-170.	0.6	35
94	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.	0.6	33
95	Genetic variants in or nearADH1BandADH1Caffect susceptibility to alcohol dependence in a British and Irish population. Addiction Biology, 2015, 20, 594-604.	1.4	33
96	A genetic risk score and diabetes predict development of alcohol-related cirrhosis in drinkers. Journal of Hepatology, 2022, 76, 275-282.	1.8	33
97	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.	1.1	32
98	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.	2.6	32
99	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.	2.0	32
100	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.	1.1	31
101	Chromosomal microarray testing in adults with intellectual disability presenting with comorbid psychiatric disorders. European Journal of Human Genetics, 2017, 25, 66-72.	1.4	30
102	Neurodevelopmental risk copy number variants in adults with intellectual disabilities and comorbid psychiatric disorders. British Journal of Psychiatry, 2018, 212, 287-294.	1.7	30
103	Optimization of liposome mediated transfection of a neuronal cell line. NeuroReport, 1997, 8, 1481-1484.	0.6	29
104	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, 469-472.	1.4	29
105	Failure to Confirm Allelic Association Between Markers at the CAPON Gene Locus and Schizophrenia in a British Sample. Biological Psychiatry, 2006, 59, 195-197.	0.7	28
106	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.	1.1	28
107	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.	1.1	27
108	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.	1.1	27

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109	Genetic association and sequencing of the insulin-like growth factor 1 gene in bipolar affective disorder. , 2011, 156, 177-187.		26
110	Hypomethylation of FAM63B in bipolar disorder patients. Clinical Epigenetics, 2016, 8, 52.	1.8	24
111	Whole-exome sequencing of individuals from an isolated population implicates rare risk variants in bipolar disorder. Translational Psychiatry, 2017, 7, e1034-e1034.	2.4	24
112	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.	1.2	24
113	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.	1.4	20
114	Characterisation of age and polarity at onset in bipolar disorder. British Journal of Psychiatry, 2021, 219, 659-669.	1.7	20
115	No association between a neuronal nitric oxide synthase (NOS1) gene polymorphism on chromosome 12q24 and bipolar disorder. , 2004, 124B, 73-75.		19
116	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.	1.1	19
117	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.	1.1	19
118	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.	6.1	19
119	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.	1.4	18
120	A nonconservative amino acid change in the UPF3B gene in a patient with schizophrenia. Psychiatric Genetics, 2012, 22, 150-151.	0.6	18
121	Genetic copy number variants, cognition and psychosis: a meta-analysis and a family study. Molecular Psychiatry, 2021, 26, 5307-5319.	4.1	18
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.	1.1	17
123	Memory Decline in Down Syndrome and Its Relationship to iPF2alpha, a Urinary Marker of Oxidative Stress. PLoS ONE, 2014, 9, e97709.	1.1	17
124	Title is missing!. Psychiatric Genetics, 2003, 13, 77-84.	0.6	16
125	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.	1.1	16
126	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.	1.3	16

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127	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.	0.3	16
128	A machine learning caseâ€“control classifier for schizophrenia based on DNA methylation in blood. Translational Psychiatry, 2021, 11, 412.	2.4	16
129	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.	2.0	15
130	Investigating rare pathogenic/likely pathogenic exonic variation in bipolar disorder. Molecular Psychiatry, 2021, 26, 5239-5250.	4.1	15
131	A novel polymorphism in exon 11 of the WKL1 gene, shows no association with schizophrenia. European Journal of Human Genetics, 2002, 10, 491-494.	1.4	14
132	Effect of the 2004 tsunami on suicide rates in Sri Lanka. Psychiatric Bulletin, 2009, 33, 179-180.	0.3	14
133	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.	1.1	14
134	Genetic variation in the miRâ€708 gene and its binding targets in bipolar disorder. Bipolar Disorders, 2016, 18, 650-656.	1.1	14
135	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.	4.0	13
136	Rare variant analysis in multiply affected families, association studies and functional analysis suggest a role for the ITGÎ'4 gene in schizophrenia and bipolar disorder. Schizophrenia Research, 2018, 199, 181-188.	1.1	11
137	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.	1.1	10
138	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.	0.6	10
139	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.	1.1	9
140	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.	0.7	9
141	Elevated common variant genetic risk for tourette syndrome in a densely-affected pedigree. Molecular Psychiatry, 2021, 26, 7522-7529.	4.1	8
142	The Influence of CYP2D6 and CYP2C19 Genetic Variation on Diabetes Mellitus Risk in People Taking Antidepressants and Antipsychotics. Genes, 2021, 12, 1758.	1.0	8
143	Identifying risk factors involved in the common versus specific liabilities to substance use: A genetically informed approach. Addiction Biology, 2021, 26, e12944.	1.4	7
144	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.	0.6	6

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145	Phenotypic heterogeneity in study populations may significantly confound the results of genetic association studies on alcohol dependence. Psychiatric Genetics, 2015, 25, 234-240.	0.6	6
146	Identification of rare nonsynonymous variants in SYNE1/CPG2 in bipolar affective disorder. Psychiatric Genetics, 2017, 27, 81-88.	0.6	6
147	Genetic variation in GABRÎ²1 and the risk for developing alcohol dependence. Psychiatric Genetics, 2017, 27, 110-115.	0.6	6
148	Genetic variants in ALDH1B1 and alcohol dependence risk in a British and Irish population: A bioinformatic and genetic study. PLoS ONE, 2017, 12, e0177009.	1.1	6
149	Genetic power of a Brazilian three-generation family with generalized aggressive periodontitis. Brazilian Dental Journal, 2010, 21, 137-141.	0.5	5
150	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.	1.8	5
151	The Communication of Metacognition for Social Strategy in Psychosis: An Exploratory Study. Schizophrenia Bulletin Open, 2020, 1, .	0.9	4
152	People who survive an episode of severe alcoholic hepatitis should be advised to maintain total abstinence from alcohol. Hepatology, 2018, 67, 2479-2480.	3.6	3
153	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.	0.6	3
154	Evidence for genetic susceptibility to the alcohol dependence syndrome from the thiamine transporter 2 gene solute carrier SLC19A3. Psychiatric Genetics, 2014, 24, 122-123.	0.6	2
155	Genetic variant analysis of the putative regulatory regions of the LRRC7 gene in bipolar disorder. Psychiatric Genetics, 2016, 26, 99-100.	0.6	2
156	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.	1.8	2
157	META-ANALYSIS OF ALCOHOL DEPENDENCE GWAS DATA FROM EUROPEAN SAMPLES ASCERTAINED FROM CLINIC AND POPULATION BASED APPROACHES. European Neuropsychopharmacology, 2019, 29, S1036.	0.3	2
158	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.	1.1	2
159	Physical Health and Clinical Phenotypes. , 2019, , 71-86.		2
160	Adolescent Verbal Memory as a Psychosis Endophenotype: A Genome-Wide Association Study in an Ancestrally Diverse Sample. Genes, 2022, 13, 106.	1.0	2
161	The genetics of schizophrenia. , 0, , 230-261.		1
162	Geneâ€“Brain Structure Relationships: Arbitrary Assumptions of Heterogeneity Generate Unfalsifiable Claimsâ€”Reply. Archives of General Psychiatry, 2007, 64, 1098.	13.8	0

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163	Association study of rare nonsynonymous variants of FTO in bipolar disorder. Psychiatric Genetics, 2016, 26, 140-141.	0.6	0
164	Phenotypic Traits of Bipolar Disorder Predicted By Schizophrenia Associated Snps In Romanian Bipolar I Patients. Preliminary results. European Neuropsychopharmacology, 2017, 27, S386-S387.	0.3	0
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