

Genome-wide association and meta-analysis of bipolar ancestry

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Ankyrin protein networks in membrane formation and stabilization. <i>Journal of Cellular and Molecular Medicine</i> , 2009, 13, 4364-4376.	1.6	121
2	Simultaneous Genotype Calling and Haplotype Phasing Improves Genotype Accuracy and Reduces False-Positive Associations for Genome-wide Association Studies. <i>American Journal of Human Genetics</i> , 2009, 85, 847-861.	2.6	186
3	Lithium: a key to the genetics of bipolar disorder. <i>Genome Medicine</i> , 2009, 1, 79.	3.6	35
4	Implication of synapse-related genes in bipolar disorder by linkage and gene expression analyses. <i>International Journal of Neuropsychopharmacology</i> , 2010, 13, 1397-1410.	1.0	47
5	Genetic Studies of Schizophrenia and Bipolar Disorder. <i>Focus (American Psychiatric Publishing)</i> , 2010, 8, 323-338.	0.4	2
8	Identifying disease polymorphisms from case-control genetic association data. <i>Genetica</i> , 2010, 138, 1147-1159.	0.5	4
9	A genome-wide association study of bipolar disorder in Norwegian individuals, followed by replication in Icelandic sample. <i>Journal of Affective Disorders</i> , 2010, 126, 312-316.	2.0	100
10	miRNA genes and the brain: implications for psychiatric disorders. <i>Human Mutation</i> , 2010, 31, 1195-1204.	1.1	64
11	Functional variants of <i>TSPAN8</i> are associated with bipolar disorder and schizophrenia. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2010, 153B, 967-972.	1.1	18
12	European collaborative study of early-onset bipolar disorder: Evidence for genetic heterogeneity on 2q14 according to age at onset. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2010, 153B, 1425-1433.	1.1	16
13	High-throughput behavioral phenotyping in the expanded panel of BXD recombinant inbred strains. <i>Genes, Brain and Behavior</i> , 2010, 9, 129-159.	1.1	199
14	Replication of GWAS of bipolar disorder: association of SNPs near <i>CDH7</i> with bipolar disorder and visual processing. <i>Molecular Psychiatry</i> , 2010, 15, 4-6.	4.1	27
15	Meta-analysis of genome-wide association data identifies a risk locus for major mood disorders on 3p21.1. <i>Nature Genetics</i> , 2010, 42, 128-131.	9.4	152
16	Meta-analysis and imputation refines the association of 15q25 with smoking quantity. <i>Nature Genetics</i> , 2010, 42, 436-440.	9.4	581
17	Increased peripheral blood expression of electron transport chain genes in bipolar depression. <i>Bipolar Disorders</i> , 2010, 12, 813-824.	1.1	50
18	Family-based association analysis to finemap bipolar linkage peak on chromosome 8q24 using 2,500 genotyped SNPs and 15,000 imputed SNPs. <i>Bipolar Disorders</i> , 2010, 12, 786-792.	1.1	34
19	Uncovering Molecular Biomarkers That Correlate Cognitive Decline with the Changes of Hippocampus' Gene Expression Profiles in Alzheimer's Disease. <i>PLoS ONE</i> , 2010, 5, e10153.	1.1	121
20	A Transcription Factor Map as Revealed by a Genome-Wide Gene Expression Analysis of Whole-Blood mRNA Transcriptome in Multiple Sclerosis. <i>PLoS ONE</i> , 2010, 5, e14176.	1.1	51

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21	Efficient population assignment and outlier detection in human populations using biallelic markers chosen by principal component-based rankings. <i>BioTechniques</i> , 2010, 48, 449-454.	0.8	5
22	A computationally fast measure of epistasis for 2 SNPs and a categorical phenotype. , 2010, 2010, 6194-7.		3
23	Pharmacogenomics of Mood Stabilizers in the Treatment of Bipolar Disorder. <i>Human Genomics and Proteomics</i> , 2010, 2, 159761.	1.5	19
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25	The Internet-Based MGS2 Control Sample: Self Report of Mental Illness. <i>American Journal of Psychiatry</i> , 2010, 167, 854-865.	4.0	48
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28	Predictive and Diagnostic Genetic Testing in Psychiatry. <i>Clinics in Laboratory Medicine</i> , 2010, 30, 829-846.	0.7	21
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30	A genome-wide association study of Cloninger's temperament scales: Implications for the evolutionary genetics of personality. <i>Biological Psychology</i> , 2010, 85, 306-317.	1.1	150
31	Convergent integration of animal model and human studies of bipolar disorder (manic-depressive) Tj ETQq0 0 0 rgBT./Overlock 10 Tf 50	1.7	12
32	The involvement of GSK3 β in bipolar disorder: Integrating evidence from multiple types of genetic studies. <i>European Neuropsychopharmacology</i> , 2010, 20, 357-368.	0.3	28
33	Genetic risk prediction in complex disease. <i>Human Molecular Genetics</i> , 2011, 20, R182-R188.	1.4	154
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39	Genome-wide Association: From Confounded to Confident. <i>Neuroscientist</i> , 2011, 17, 174-184.	2.6	4
40	Admixture analysis of age at onset in bipolar disorder. <i>Psychiatry Research</i> , 2011, 185, 27-32.	1.7	51
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43	Efficient Replication of Over 180 Genetic Associations with Self-Reported Medical Data. <i>Nature Precedings</i> , 2011, , .	0.1	1
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56	Random-Effects Model Aimed at Discovering Associations in Meta-Analysis of Genome-wide Association Studies. <i>American Journal of Human Genetics</i> , 2011, 88, 586-598.	2.6	515
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67	Disease-associated epigenetic changes in monozygotic twins discordant for schizophrenia and bipolar disorder. Human Molecular Genetics, 2011, 20, 4786-4796.	1.4	407
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156	Replication of genome-wide association study (GWAS) susceptibility loci in a Latino bipolar disorder cohort. Bipolar Disorders, 2016, 18, 520-527.	1.1	25
157	Genetic factors in the etiology of bipolar disorder. , 0, , 144-168.		0
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