Thomas F. Hansen

List of Publications by Year
in descending order

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194 papers	38,603 citations	28736 57 h-index	4217 180 g-index
211 all docs	211 docs citations	211 times ranked	41149 citing authors

#	Article	IF	Citations
1	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. Biological Psychiatry, 2022, 91, 102-117.	0.7	61
2	Identifying the Common Genetic Basis of Antidepressant Response. Biological Psychiatry Global Open Science, 2022, 2, 115-126.	1.0	31
3	Genome-wide association study of febrile seizures implicates fever response and neuronal excitability genes. Brain, 2022, 145, 555-568.	3.7	29
4	Cohort Profile: COVIDMENT: COVID-19 cohorts on mental health across six nations. International Journal of Epidemiology, 2022, 51, e108-e122.	0.9	16
5	Smooth muscle ATP-sensitive potassium channels mediate migraine-relevant hypersensitivity in mouse models. Cephalalgia, 2022, 42, 93-107.	1.8	11
6	Rare SLC13A1 variants associate with intervertebral disc disorder highlighting role of sulfate in disc pathology. Nature Communications, 2022, 13, 634.	5.8	21
7	Genome-wide analysis of 102,084 migraine cases identifies 123 risk loci and subtype-specific risk alleles. Nature Genetics, 2022, 54, 152-160.	9.4	135
8	Elucidating the relationship between migraine risk and brain structure using genetic data. Brain, 2022, 145, 3214-3224.	3.7	7
9	A genome-wide meta-analysis identifies 50 genetic loci associated with carpal tunnel syndrome. Nature Communications, 2022, 13, 1598.	5.8	8
10	Hyperhidrosis and human leucocyte antigens in the Danish Blood Donor Study. Scandinavian Journal of Immunology, 2022, , e13150.	1.3	0
11	Incidence and remission rates of selfâ€reported hidradenitis suppurativa â€•A prospective cohort study conducted in Danish blood donors. Journal of the European Academy of Dermatology and Venereology, 2022, 36, 717-725.	1.3	6
12	Low adherence to the guideline for the acute treatment of migraine. Scientific Reports, 2022, 12, 8487.	1.6	12
13	Population-based prevalence of cranial autonomic symptoms in migraine and proposed diagnostic appendix criteria. Cephalalgia, 2022, 42, 1160-1171.	1.8	7
14	Genome-wide association study of panic disorder reveals genetic overlap with neuroticism and depression. Molecular Psychiatry, 2021, 26, 4179-4190.	4.1	58
15	Association of polygenic score for major depression with response to lithium in patients with bipolar disorder. Molecular Psychiatry, 2021, 26, 2457-2470.	4.1	44
16	Left ventricular systolic ejection time is an independent predictor of allâ€cause mortality in heart failure with reduced ejection fraction. European Journal of Heart Failure, 2021, 23, 240-249.	2.9	17
17	Genome-wide association study identifies 48 common genetic variants associated with handedness. Nature Human Behaviour, 2021, 5, 59-70.	6.2	79
18	Superficial fungal infections and patients with hidradenitis suppurativa: a study under the Danish Blood Donor Study. Clinical and Experimental Dermatology, 2021, 46, 571-573.	0.6	0

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19	Chronic migraine: Genetics or environment?. European Journal of Neurology, 2021, 28, 1726-1736.	1.7	10
20	Epidemiology of Hyperhidrosis in Danish Blood Donors. Acta Dermato-Venereologica, 2021, 101, adv00435.	0.6	7
21	A meta-analysis uncovers the first sequence variant conferring risk of Bellâ€™s palsy. Scientific Reports, 2021, 11, 4188.	1.6	8
22	A genome-wide meta-analysis yields 46 new loci associating with biomarkers of iron homeostasis. Communications Biology, 2021, 4, 156.	2.0	72
23	Genetic insight into sick sinus syndrome. European Heart Journal, 2021, 42, 1959-1971.	1.0	27
24	Twenty-five years of triptans â€“ a nationwide population study. Cephalalgia, 2021, 41, 894-904.	1.8	9
25	Combinations of selfâ€reported rhinitis, conjunctivitis, and asthma predicts IgE sensitization in more than 25,000 Danes. Clinical and Translational Allergy, 2021, 11, e12013.	1.4	7
26	Changes in the gene expression profile during spontaneous migraine attacks. Scientific Reports, 2021, 11, 8294.	1.6	10
27	A Comparison of Ten Polygenic Score Methods for Psychiatric Disorders Applied Across Multiple Cohorts. Biological Psychiatry, 2021, 90, 611-620.	0.7	103
28	The genetic structure of Norway. European Journal of Human Genetics, 2021, 29, 1710-1718.	1.4	10
29	Eleven genomic loci affect plasma levels of chronic inflammation marker soluble urokinase-type plasminogen activator receptor. Communications Biology, 2021, 4, 655.	2.0	12
30	Impact of COVID-19 Pandemic on Sleep Quality, Stress Level and Health-Related Quality of Lifeâ€”A Large Prospective Cohort Study on Adult Danes. International Journal of Environmental Research and Public Health, 2021, 18, 7610.	1.2	13
31	Genetic Susceptibility Loci in Genomewide Association Study of Cluster Headache. Annals of Neurology, 2021, 90, 203-216.	2.8	22
32	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. JAMA Psychiatry, 2021, 78, 1258.	6.0	88
33	Pain sensitivity in men who have never experienced a headache: an observer blinded case control study. Journal of Headache and Pain, 2021, 22, 134.	2.5	2
34	Headache provocation by nitric oxide in men who have never experienced a headache. Cephalalgia, 2021, , 033310242110600.	1.8	0
35	Proposed new diagnostic criteria for chronic migraine. Cephalalgia, 2020, 40, 399-406.	1.8	50
36	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. Biological Psychiatry, 2020, 87, 419-430.	0.7	27

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37	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
38	Von Frey testing revisited: Provision of an online algorithm for improved accuracy of 50% thresholds. European Journal of Pain, 2020, 24, 783-790.	1.4	40
39	PCM1 is necessary for focal ciliary integrity and is a candidate for severe schizophrenia. Nature Communications, 2020, 11, 5903.	5.8	13
40	Genetic variability in the absorption of dietary sterols affects the risk of coronary artery disease. European Heart Journal, 2020, 41, 2618-2628.	1.0	61
41	The impact of low-risk genetic variants in self-limited epilepsy with centrotemporal spikes aka Rolandic epilepsy. EBioMedicine, 2020, 58, 102896.	2.7	2
42	Higher burden of rare frameshift indels in genes related to synaptic transmission separate familial hemiplegic migraine from common types of migraine. Journal of Medical Genetics, 2020, 57, 610-616.	1.5	10
43	Functional gene networks reveal distinct mechanisms segregating in migraine families. Brain, 2020, 143, 2945-2956.	3.7	15
44	Habitual sleep disturbances and migraine: a Mendelian randomization study. Annals of Clinical and Translational Neurology, 2020, 7, 2370-2380.	1.7	18
45	The donors perceived positive and negative effects of blood donation. Transfusion, 2020, 60, 553-560.	0.8	6
46	Familial analysis reveals rare risk variants for migraine in regulatory regions. Neurogenetics, 2020, 21, 149-157.	0.7	11
47	A genome-wide cross-phenotype meta-analysis of the association of blood pressure with migraine. Nature Communications, 2020, 11, 3368.	5.8	49
48	Letter to the editor regarding â€œHave you ever experienced a headache of any kind?â€• Cephalalgia, 2020, 40, 1134-1135.	1.8	0
49	Coronary CT Angiography in Patients With Non-ST-Segment Elevation Acute CoronaryÂSyndrome. Journal of the American College of Cardiology, 2020, 75, 453-463.	1.2	123
50	Crossâ€sectional study identifies lower risk of Staphylococcus aureus nasal colonization in Danish blood donors with hidradenitis suppurativa symptoms. British Journal of Dermatology, 2020, 183, 387-389.	1.4	5
51	Genome-wide gene-environment analyses of major depressive disorder and reported lifetime traumatic experiences in UK Biobank. Molecular Psychiatry, 2020, 25, 1430-1446.	4.1	116
52	Genetic identification of cell types underlying brain complex traits yields insights into the etiology of Parkinsonâ€™s disease. Nature Genetics, 2020, 52, 482-493.	9.4	216
53	Prevalence and socio-demographic characteristics of persons who have never had a headache among healthy voluntary blood donors â€“ a population-based study. Cephalalgia, 2020, 40, 1055-1062.	1.8	4
54	Cross-trait analyses with migraine reveal widespread pleiotropy and suggest a vascular component to migraine headache. International Journal of Epidemiology, 2020, 49, 1022-1031.	0.9	34

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55	Genome-wide association study identifies 16 genomic regions associated with circulating cytokines at birth. PLoS Genetics, 2020, 16, e1009163.	1.5	12
56	Title is missing!. , 2020, 16, e1009163.		0
57	Title is missing!. , 2020, 16, e1009163.		0
58	Title is missing!. , 2020, 16, e1009163.		0
59	Title is missing!. , 2020, 16, e1009163.		0
60	Herpes Simplex Virus Type 1 infection is associated with suicidal behavior and first registered psychiatric diagnosis in a healthy population. Psychoneuroendocrinology, 2019, 108, 150-154.	1.3	10
61	Echocardiography improves prediction of major adverse cardiovascular events in a population with type 1 diabetes and without known heart disease: the Thousand & 1 Study. Diabetologia, 2019, 62, 2354-2364.	2.9	23
62	The first step towards personalized risk prediction for common epilepsies. Brain, 2019, 142, 3316-3318.	3.7	2
63	Characterization of Familial and Sporadic Migraine. Headache, 2019, 59, 1802-1807.	1.8	5
64	Comparing migraine with and without aura to healthy controls using RNA sequencing. Cephalalgia, 2019, 39, 1435-1444.	1.8	12
65	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
66	DBDS Genomic Cohort, a prospective and comprehensive resource for integrative and temporal analysis of genetic, environmental and lifestyle factors affecting health of blood donors. BMJ Open, 2019, 9, e028401.	0.8	68
67	Predicting treatment response using pharmacy register in migraine. Journal of Headache and Pain, 2019, 20, 31.	2.5	8
68	Populationâ€based identityâ€byâ€descent mapping combined with exome sequencing to detect rare risk variants for schizophrenia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, 223-231.	1.1	2
69	The preferences of potential stakeholders in psychiatric genomic research regarding consent procedures and information delivery. European Psychiatry, 2019, 55, 29-35.	0.1	10
70	Self-reported restless legs syndrome and involuntary leg movements during sleep are associated with symptoms of attention deficit hyperactivity disorder. Sleep Medicine, 2019, 57, 115-121.	0.8	10
71	Migraine polygenic risk score associates with efficacy of migraine-specific drugs. Neurology: Genetics, 2019, 5, e364.	0.9	28
72	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. Cell, 2019, 179, 1469-1482.e11.	13.5	935

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73	Association of Whole-Genome and NETRIN1 Signaling Pathwayâ€"Derived Polygenic Risk Scores for Major Depressive Disorder and White Matter Microstructure in the UK Biobank. Biological Psychiatry: Cognitive Neuroscience and Neuroimaging, 2019, 4, 91-100.	1.1	16
74	Nosographic analysis of osmophobia and field testing of diagnostic criteria including osmophobia. Cephalalgia, 2019, 39, 38-43.	1.8	13
75	Multi-view Consensus CNN for 3D Facial Landmark Placement. Lecture Notes in Computer Science, 2019, , 706-719.	1.0	13
76	UGT polymorphisms and lamotrigine clearance during pregnancy. Epilepsy Research, 2018, 140, 199-208.	0.8	29
77	Value of Myocardial Perfusion Assessment With Coronary Computed Tomography Angiography in Patients With RecentÂAcute-Onset Chest Pain. JACC: Cardiovascular Imaging, 2018, 11, 1611-1621.	2.3	34
78	Attitudes of stakeholders in psychiatry towards the inclusion of children in genomic research. Human Genomics, 2018, 12, 12.	1.4	15
79	Transcriptomic profiling of trigeminal nucleus caudalis and spinal cord dorsal horn. Brain Research, 2018, 1692, 23-33.	1.1	5
80	Common Variant Burden Contributes to the Familial Aggregation of Migraine in 1,589 Families. Neuron, 2018, 98, 743-753.e4.	3.8	63
81	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. Nature Genetics, 2018, 50, 668-681.	9.4	2,224
82	Does Childhood Trauma Moderate Polygenic Risk for Depression? A Meta-analysis of 5765 Subjects From the Psychiatric Genomics Consortium. Biological Psychiatry, 2018, 84, 138-147.	0.7	87
83	Comorbidity of migraine with ADHD in adults. BMC Neurology, 2018, 18, 147.	0.8	24
84	Restless legs syndrome is associated with increased risk of migraine. Cephalalgia Reports, 2018, 1, 251581631878074.	0.2	3
85	Ancient genomes from Iceland reveal the making of a human population. Science, 2018, 360, 1028-1032.	6.0	62
86	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
87	Age at first birth in women is genetically associated with increased risk of schizophrenia. Scientific Reports, 2018, 8, 10168.	1.6	17
88	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	6.0	1,085
89	Molecular genetic overlap between migraine and major depressive disorder. European Journal of Human Genetics, 2018, 26, 1202-1216.	1.4	56
90	Polygenic risk score: use in migraine research. Journal of Headache and Pain, 2018, 19, 29.	2.5	24

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91	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. Cell, 2018, 173, 1705-1715.e16.	13.5	623
92	Socio-demographic characteristics of adults with self-reported ADHD symptoms in a Danish population of 12,415 blood donors. Journal of Psychiatry and Behavioral Science, 2018, 1, .	0.1	1
93	Identification of Gene Loci That Overlap Between Schizophrenia and Educational Attainment. Schizophrenia Bulletin, 2017, 43, sbw085.	2.3	56
94	Whole transcriptome expression of trigeminal ganglia compared to dorsal root ganglia in Rattus Norvegicus. Neuroscience, 2017, 350, 169-179.	1.1	37
95	Genetic correlation between amyotrophic lateral sclerosis and schizophrenia. Nature Communications, 2017, 8, 14774.	5.8	114
96	CNV-association meta-analysis in 191,161 European adults reveals new loci associated with anthropometric traits. Nature Communications, 2017, 8, 744.	5.8	64
97	Genome-wide meta-analysis associates HLA-DQA1/DRB1 and LPA and lifestyle factors with human longevity. Nature Communications, 2017, 8, 910.	5.8	118
98	Stakeholders in psychiatry and their attitudes toward receiving pertinent and incident findings in genomic research. American Journal of Medical Genetics, Part A, 2017, 173, 2649-2658.	0.7	20
99	Evaluation of shared genetic susceptibility loci between autoimmune diseases and schizophrenia based on genome-wide association studies. Nordic Journal of Psychiatry, 2017, 71, 20-25.	0.7	10
100	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
101	Investigation of SNP rs2060546 Immediately Upstream to NTN4 in a Danish Gilles de la Tourette Syndrome Cohort. Frontiers in Neuroscience, 2016, 10, 531.	1.4	5
102	Leveraging Genomic Annotations and Pleiotropic Enrichment for Improved Replication Rates in Schizophrenia GWAS. PLoS Genetics, 2016, 12, e1005803.	1.5	34
103	Digital questionnaire platform in the Danish Blood Donor Study. Computer Methods and Programs in Biomedicine, 2016, 135, 101-104.	2.6	38
104	Influence of Septal Thickness on the Clinical Outcome After Alcohol Septal Alation in Hypertrophic Cardiomyopathy. Circulation: Cardiovascular Interventions, 2016, 9, .	1.4	13
105	Meta-analysis of 375,000 individuals identifies 38 susceptibility loci for migraine. Nature Genetics, 2016, 48, 856-866.	9.4	520
106	Genome-Wide Association Studies Suggest Limited Immune Gene Enrichment in Schizophrenia Compared to 5 Autoimmune Diseases. Schizophrenia Bulletin, 2016, 42, 1176-1184.	2.3	62
107	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
108	Genetic risk for autism spectrum disorders and neuropsychiatric variation in the general population. Nature Genetics, 2016, 48, 552-555.	9.4	326

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109	Evidence for Genetic Overlap Between Schizophrenia and Age at First Birth in Women. JAMA Psychiatry, 2016, 73, 497.	6.0	51
110	Comprehensive analysis of schizophrenia-associated loci highlights ion channel pathways and biologically plausible candidate causal genes. Human Molecular Genetics, 2016, 25, 1247-1254.	1.4	69
111	Genetic influences on schizophrenia and subcortical brain volumes: large-scale proof of concept. Nature Neuroscience, 2016, 19, 420-431.	7.1	204
112	Identification of rare high-risk copy number variants affecting the dopamine transporter gene in mental disorders. Nordic Journal of Psychiatry, 2016, 70, 276-279.	0.7	2
113	Genetic Markers of Human Evolution Are Enriched in Schizophrenia. Biological Psychiatry, 2016, 80, 284-292.	0.7	92
114	The association between candidate migraine susceptibility loci and severe migraine phenotype in a clinical sample. Cephalalgia, 2016, 36, 615-623.	1.8	24
115	Investigating the Causal Relationship of C-Reactive Protein with 32 Complex Somatic and Psychiatric Outcomes: A Large-Scale Cross-Consortium Mendelian Randomization Study. PLoS Medicine, 2016, 13, e1001976.	3.9	150
116	RNA Sequencing of Trigeminal Ganglia in Rattus Norvegicus after Glyceryl Trinitrate Infusion with Relevance to Migraine. PLoS ONE, 2016, 11, e0155039.	1.1	8
117	An Empirical Bayes Mixture Model for Effect Size Distributions in Genome-Wide Association Studies. PLoS Genetics, 2015, 11, e1005717.	1.5	22
118	Combinations of Genetic Data Present in Bipolar Patients, but Absent in Control Persons. PLoS ONE, 2015, 10, e0143432.	1.1	4
119	Linkage and whole genome sequencing identify a locus on 6q25â€“26 for formal thought disorder and implicate MEF2A regulation. Schizophrenia Research, 2015, 169, 441-446.	1.1	12
120	Psychiatric genome-wide association study analyses implicate neuronal, immune and histone pathways. Nature Neuroscience, 2015, 18, 199-209.	7.1	701
121	Modeling Linkage Disequilibrium Increases Accuracy of Polygenic Risk Scores. American Journal of Human Genetics, 2015, 97, 576-592.	2.6	1,098
122	An atlas of genetic correlations across human diseases and traits. Nature Genetics, 2015, 47, 1236-1241.	9.4	3,145
123	Contrasting genetic architectures of schizophrenia and other complex diseases using fast variance-components analysis. Nature Genetics, 2015, 47, 1385-1392.	9.4	431
124	Usefulness of the SNP microarray technology to identify rare mutations in the case of perinatal death. Case Reports in Perinatal Medicine, 2015, 4, .	0.1	1
125	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
126	3D facial landmarks: Inter-operator variability of manual annotation. BMC Medical Imaging, 2014, 14, 35.	1.4	31

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127	Common variant at 16p11.2 conferring risk of psychosis. Molecular Psychiatry, 2014, 19, 108-114.	4.1	85
128	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
129	Biological insights from 108 schizophrenia-associated genetic loci. Nature, 2014, 511, 421-427.	13.7	6,934
130	Sequence analysis of 17 NRXN1 deletions. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 52-61.	1.1	11
131	Genetic analyses of the human eye colours using a novel objective method for eye colour classification. Forensic Science International: Genetics, 2013, 7, 508-515.	1.6	31
132	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. Nature Genetics, 2013, 45, 984-994.	9.4	2,067
133	Improved Detection of Common Variants Associated with Schizophrenia by Leveraging Pleiotropy with Cardiovascular-Disease Risk Factors. American Journal of Human Genetics, 2013, 92, 197-209.	2.6	422
134	Identification of risk loci with shared effects on five major psychiatric disorders: a genome-wide analysis. Lancet, The, 2013, 381, 1371-1379.	6.3	2,643
135	Genome-wide significant associations in schizophrenia to ITIH3/4, CACNA1C and SDCCAG8, and extensive replication of associations reported by the Schizophrenia PGC. Molecular Psychiatry, 2013, 18, 708-712.	4.1	216
136	All SNPs Are Not Created Equal: Genome-Wide Association Studies Reveal a Consistent Pattern of Enrichment among Functionally Annotated SNPs. PLoS Genetics, 2013, 9, e1003449.	1.5	268
137	Improved Detection of Common Variants Associated with Schizophrenia and Bipolar Disorder Using Pleiotropy-Informed Conditional False Discovery Rate. PLoS Genetics, 2013, 9, e1003455.	1.5	298
138	Replication and metaâ€analysis of common variants identifies a genomeâ€wide significant locus in migraine. European Journal of Neurology, 2013, 20, 765-772.	1.7	67
139	3D gender recognition using cognitive modeling. , 2013, , .		0
140	Genome-Wide Association Study of Genetic Variants in LPS-Stimulated IL-6, IL-8, IL-10, IL-1ra and TNF-Î± Cytokine Response in a Danish Cohort. PLoS ONE, 2013, 8, e66262.	1.1	10
141	No association of polymorphisms in human endogenous retrovirus K18 and CD48 with schizophrenia. Psychiatric Genetics, 2012, 22, 146-148.	0.6	10
142	Investigation of the Genetic Association between Quantitative Measures of Psychosis and Schizophrenia: A Polygenic Risk Score Analysis. PLoS ONE, 2012, 7, e37852.	1.1	60
143	The gene encoding the melanin-concentrating hormone receptor 1 is associated with schizophrenia in a Danish caseâ€“control sample. Psychiatric Genetics, 2012, 22, 62-69.	0.6	11
144	Estimating the proportion of variation in susceptibility to schizophrenia captured by common SNPs. Nature Genetics, 2012, 44, 247-250.	9.4	578

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145	Expression QTL analysis of top loci from GWAS meta-analysis highlights additional schizophrenia candidate genes. European Journal of Human Genetics, 2012, 20, 1004-1008.	1.4	60
146	Gene-Based Analysis of Regionally Enriched Cortical Genes in GWAS Data Sets of Cognitive Traits and Psychiatric Disorders. PLoS ONE, 2012, 7, e31687.	1.1	40
147	Connection between Genetic and Clinical Data in Bipolar Disorder. PLoS ONE, 2012, 7, e44623.	1.1	8
148	Association between genetic variation in a region on chromosome 11 and schizophrenia in large samples from Europe. Molecular Psychiatry, 2012, 17, 906-917.	4.1	105
149	Promoter variants in IL18 are associated with onset of depression in patients previously exposed to stressful-life events. Journal of Affective Disorders, 2012, 136, 134-138.	2.0	47
150	Candidate Gene Analysis of the Human Natural Killer-1 Carbohydrate Pathway and Perineuronal Nets in Schizophrenia: B3GAT2 Is Associated with Disease Risk and Cortical Surface Area. Biological Psychiatry, 2011, 69, 90-96.	0.7	42
151	The Complement Control-Related Genes CSMD1 and CSMD2 Associate to Schizophrenia. Biological Psychiatry, 2011, 70, 35-42.	0.7	149
152	At-Risk Variant in TCF7L2 for Type II Diabetes Increases Risk of Schizophrenia. Biological Psychiatry, 2011, 70, 59-63.	0.7	114
153	Genome-wide association study identifies five new schizophrenia loci. Nature Genetics, 2011, 43, 969-976.	9.4	1,758
154	Kynurenine 3-monooxygenase (KMO) polymorphisms in schizophrenia: An association study. Schizophrenia Research, 2011, 127, 270-272.	1.1	19
155	Combinations of SNPs Related to Signal Transduction in Bipolar Disorder. PLoS ONE, 2011, 6, e23812.	1.1	20
156	Dual association of a TRKA polymorphism with schizophrenia. Psychiatric Genetics, 2011, 21, 125-131.	0.6	8
157	Copy number variations of chromosome 16p13.1 region associated with schizophrenia. Molecular Psychiatry, 2011, 16, 17-25.	4.1	227
158	Expanding the range of ZNF804A variants conferring risk of psychosis. Molecular Psychiatry, 2011, 16, 59-66.	4.1	140
159	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
160	Maternally Derived Microduplications at 15q11-q13: Implication of Imprinted Genes in Psychotic Illness. American Journal of Psychiatry, 2011, 168, 408-417.	4.0	95
161	Copy number variations in affective disorders and meta-analysis. Psychiatric Genetics, 2011, 21, 319-322.	0.6	3
162	Using Electronic Patient Records to Discover Disease Correlations and Stratify Patient Cohorts. PLoS Computational Biology, 2011, 7, e1002141.	1.5	236

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163	Genome-Wide Association Study Identifies Four Loci Associated with Eruption of Permanent Teeth. PLoS Genetics, 2011, 7, e1002275.	1.5	42
164	Association study of PDE4B gene variants in scandinavian schizophrenia and bipolar disorder multicenter caseâ€“control samples. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 86-96.	1.1	25
165	The tryptophan hydroxylase 1 (TPH1) gene, schizophrenia susceptibility, and suicidal behavior: A multiâ€centre caseâ€“control study and metaâ€analysis. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 387-396.	1.1	45
166	Association between methylenetetrahydrofolate reductase (MTHFR) C677T polymorphism and age of onset in schizophrenia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 610-618.	1.1	32
167	Association analysis of PALB2 and BRCA2 in bipolar disorder and schizophrenia in a scandinavian caseâ€“control sample. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 1276-1282.	1.1	20
168	Polymorphisms in SREBF1 and SREBF2, two antipsychotic-activated transcription factors controlling cellular lipogenesis, are associated with schizophrenia in German and Scandinavian samples. Molecular Psychiatry, 2010, 15, 463-472.	4.1	66
169	A large replication study and meta-analysis in European samples provides further support for association of AHI1 markers with schizophrenia. Human Molecular Genetics, 2010, 19, 1379-1386.	1.4	51
170	Diastolic dysfunction predicts new-onset atrial fibrillation and cardiovascular events in patients with acute myocardial infarction and depressed left ventricular systolic function: a CARISMA substudy. European Journal of Echocardiography, 2010, 11, 602-607.	2.3	59
171	CYP2D6 Genotype Predicts Antipsychotic Side Effects in Schizophrenia Inpatients: A Retrospective Matched Case-Control Study. Neuropsychobiology, 2009, 59, 222-226.	0.9	51
172	Disruption of the neurexin 1 gene is associated with schizophrenia. Human Molecular Genetics, 2009, 18, 988-996.	1.4	424
173	Evidence for a possible association of neurotrophin receptor (NTRK-3) gene polymorphisms with hippocampal function and schizophrenia. Neurobiology of Disease, 2009, 34, 518-524.	2.1	46
174	Three-cohort targeted gene screening reveals a non-synonymous TRKA polymorphism associated with schizophrenia. Journal of Psychiatric Research, 2009, 43, 1195-1199.	1.5	25
175	Common variants conferring risk of schizophrenia. Nature, 2009, 460, 744-747.	13.7	1,572
176	An intron 1 polymorphism in the cholecystokininâ€A receptor gene associated with schizophrenia in males. Acta Psychiatrica Scandinavica, 2009, 120, 281-287.	2.2	20
177	A possible association between schizophrenia and GRIK3 polymorphisms in a multicenter sample of Scandinavian origin (SCOPE). Schizophrenia Research, 2009, 107, 242-248.	1.1	17
178	Tyrosine hydroxylase Val81Met polymorphism: lack of association with schizophrenia. Psychiatric Genetics, 2009, 19, 273-274.	0.6	3
179	The estrogen hypothesis of Schizophrenia implicates glucose metabolism: Association study in three independent samples. BMC Medical Genetics, 2008, 9, 39.	2.1	31
180	Two methylenetetrahydrofolate reductase gene (MTHFR) polymorphisms, schizophrenia and bipolar disorder: An association study. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2008, 147B, 976-982.	1.1	51

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181	Association analysis of schizophrenia on 18 genes involved in neuronal migration: MDGA1 as a new susceptibility gene. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2008, 147B, 1089-1100.	1.1	101
182	Large recurrent microdeletions associated with schizophrenia. Nature, 2008, 455, 232-236.	13.7	1,619
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