Sven Cichon

List of Publications by Year
in descending order

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397 papers	56,804 citations	2091 103 h-index	1719 219 g-index
440 all docs	440 docs citations	440 times ranked	56297 citing authors

#	Article	IF	Citations
1	Effects of copy number variations on brain structure and risk for psychiatric illness: Largeâ€scale studies from the<scp>ENIGMA</scp>working groups on<scp>CNVs</scp>. Human Brain Mapping, 2022, 43, 300-328.	1.9	30
2	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. Biological Psychiatry, 2022, 91, 102-117.	0.7	61
3	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. Biological Psychiatry, 2022, 91, 313-327.	0.7	114
4	Efficacy and safety of exogenous beta-hydroxybutyrate for preventive treatment in episodic migraine: A single-centred, randomised, placebo-controlled, double-blind crossover trial. Cephalalgia, 2022, 42, 302-311.	1.8	6
5	Genome-wide Association Study of Postoperative Cognitive Dysfunction in Older Surgical Patients. Journal of Neurosurgical Anesthesiology, 2022, 34, 248-250.	0.6	3
6	Using polygenic scores and clinical data for bipolar disorder patient stratification and lithium response prediction: machine learning approach. British Journal of Psychiatry, 2022, 220, 219-228.	1.7	11
7	Genetic variants associated with longitudinal changes in brain structure across the lifespan. Nature Neuroscience, 2022, 25, 421-432.	7.1	75
8	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	13.7	929
9	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
10	Shared genetic risk between eating disorderâ€•and substanceâ€useâ€related phenotypes: Evidence from genomeâ€wide association studies. Addiction Biology, 2021, 26, e12880.	1.4	28
11	Bipolar multiplex families have an increased burden of common risk variants for psychiatric disorders. Molecular Psychiatry, 2021, 26, 1286-1298.	4.1	33
12	Genetic factors influencing a neurobiological substrate for psychiatric disorders. Translational Psychiatry, 2021, 11, 192.	2.4	4
13	1q21.1 distal copy number variants are associated with cerebral and cognitive alterations in humans. Translational Psychiatry, 2021, 11, 182.	2.4	24
14	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
15	Analysis of genetic impact on smell impairment in patients with hereditary angioedema typeÂ1 and 2. JDDG - Journal of the German Society of Dermatology, 2021, 19, 1060-1062.	0.4	0
16	Identification of pleiotropy at the gene level between psychiatric disorders and related traits. Translational Psychiatry, 2021, 11, 410.	2.4	7
17	Untersuchung genetischer EinflÃ¼sse auf RiechstÃ¶rungen bei Patienten mit hereditÃrem AngioÃ¶dem Typ 1 und 2. JDDG - Journal of the German Society of Dermatology, 2021, 19, 1060-1063.	0.4	0
18	Characterisation of age and polarity at onset in bipolar disorder. British Journal of Psychiatry, 2021, 219, 659-669.	1.7	20

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19	Identification of Phonology-Related Genes and Functional Characterization of Brocaâ€™s and Wernickeâ€™s Regions in Language and Learning Disorders. Frontiers in Neuroscience, 2021, 15, 680762.	1.4	7
20	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. JAMA Psychiatry, 2021, 78, 1258.	6.0	88
21	HLA-DRB1 and HLA-DQB1 genetic diversity modulates response to lithium in bipolar affective disorders. Scientific Reports, 2021, 11, 17823.	1.6	10
22	Association of Attention-Deficit/Hyperactivity Disorder and Depression Polygenic Scores with Lithium Response: A Consortium for Lithium Genetics Study. Complex Psychiatry, 2021, 7, 80-89.	1.3	6
23	Combining schizophrenia and depression polygenic risk scores improves the genetic prediction of lithium response in bipolar disorder patients. Translational Psychiatry, 2021, 11, 606.	2.4	25
24	Common and rare variant association analyses in amyotrophic lateral sclerosis identify 15 risk loci with distinct genetic architectures and neuron-specific biology. Nature Genetics, 2021, 53, 1636-1648.	9.4	223
25	Dose response of the 16p11.2 distal copy number variant on intracranial volume and basal ganglia. Molecular Psychiatry, 2020, 25, 584-602.	4.1	49
26	Identification of ADHD risk genes in extended pedigrees by combining linkage analysis and whole-exome sequencing. Molecular Psychiatry, 2020, 25, 2047-2057.	4.1	17
27	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. Biological Psychiatry, 2020, 87, 419-430.	0.7	27
28	Association of Copy Number Variation of the 15q11.2 BP1-BP2 Region With Cortical and Subcortical Morphology and Cognition. JAMA Psychiatry, 2020, 77, 420.	6.0	54
29	International Consensus on the Use of Genetics in the Management of Hereditary Angioedema. Journal of Allergy and Clinical Immunology: in Practice, 2020, 8, 901-911.	2.0	43
30	A large-scale genome-wide association study meta-analysis of cannabis use disorder. Lancet Psychiatry,the, 2020, 7, 1032-1045.	3.7	200
31	Gene expression is stable in a complete CIB1 knockout keratinocyte model. Scientific Reports, 2020, 10, 14952.	1.6	2
32	The genetic architecture of the human cerebral cortex. Science, 2020, 367, .	6.0	450
33	Quantitative genome-wide association study of six phenotypic subdomains identifies novel genome-wide significant variants in autism spectrum disorder. Translational Psychiatry, 2020, 10, 215.	2.4	13
34	Whole-exome sequencing of 81 individuals from 27 multiply affected bipolar disorder families. Translational Psychiatry, 2020, 10, 57.	2.4	23
35	Insights into the genomics of affective disorders. Medizinische Genetik, 2020, 32, 9-18.	0.1	2
36	Brain imaging genomics: influences of genomic variability on the structure and function of the human brain. Medizinische Genetik, 2020, 32, 47-56.	0.1	3

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37	Investigating polygenic burden in age at disease onset in bipolar disorder: Findings from an international multicentric study. Bipolar Disorders, 2019, 21, 68-75.	1.1	20
38	Pathway-Specific Genetic Risk for Alzheimerâ€™s Disease Differentiates Regional Patterns of Cortical Atrophy in Older Adults. Cerebral Cortex, 2019, 30, 801-811.	1.6	11
39	Genome-wide association study identifies eight risk loci and implicates metabo-psychiatric origins for anorexia nervosa. Nature Genetics, 2019, 51, 1207-1214.	9.4	641
40	SU30ANALYSIS OF WGS DATA FROM 108 INDIVIDUALS OF 8 SPANISH FAMILIES AFFECTED WITH BIPOLAR DISORDER. European Neuropsychopharmacology, 2019, 29, S1283-S1284.	0.3	0
41	Efficacy and safety of exogenous ketone bodies for preventive treatment of migraine: A study protocol for a single-centred, randomised, placebo-controlled, double-blind crossover trial. Trials, 2019, 20, 61.	0.7	17
42	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
43	Associations Between Attention-Deficit/Hyperactivity Disorder and Various Eating Disorders: A Swedish Nationwide Population Study Using Multiple Genetically Informative Approaches. Biological Psychiatry, 2019, 86, 577-586.	0.7	43
44	EXOME SEQUENCING OF MULTIPLY AFFECTED BIPOLAR DISORDER FAMILIES AND FOLLOW-UP RESEQUENCING IMPLICATE RARE VARIANTS IN NEURONAL GENES CONTRIBUTING TO DISEASE ETIOLOGY. European Neuropsychopharmacology, 2019, 29, S836-S837.	0.3	0
45	Genome-wide association study identifies 30 loci associated with bipolar disorder. Nature Genetics, 2019, 51, 793-803.	9.4	1,191
46	Combining lifestyle risks to disentangle brain structure and functional connectivity differences in older adults. Nature Communications, 2019, 10, 621.	5.8	42
47	Genetic architecture of subcortical brain structures in 38,851 individuals. Nature Genetics, 2019, 51, 1624-1636.	9.4	192
48	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. Cell, 2019, 179, 1469-1482.e11.	13.5	935
49	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
50	Leptin gene polymorphisms are associated with weight gain during lithium augmentation in patients with major depression. European Neuropsychopharmacology, 2019, 29, 211-221.	0.3	13
51	Genotypeâ€phenotype correlations in Brazilian patients with hereditary angioedema due to C1 inhibitor deficiency. Allergy: European Journal of Allergy and Clinical Immunology, 2019, 74, 1013-1016.	2.7	13
52	Efficient region-based test strategy uncovers genetic risk factors for functional outcome in bipolar disorder. European Neuropsychopharmacology, 2019, 29, 156-170.	0.3	7
53	Association of Polygenic Score for Schizophrenia and HLA Antigen and Inflammation Genes With Response to Lithium in Bipolar Affective Disorder. JAMA Psychiatry, 2018, 75, 65-74.	6.0	102
54	Integration of transcriptomic and cytoarchitectonic data implicates a role for MAOA and TAC1 in the limbic-cortical network. Brain Structure and Function, 2018, 223, 2335-2342.	1.2	19

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55	Hereditary Angioedema with Normal C1 Inhibitor and F12 Mutations in 42 Brazilian Families. Journal of Allergy and Clinical Immunology: in Practice, 2018, 6, 1209-1216.e8.	2.0	43
56	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
57	The 5-HTTLPR Polymorphism Affects Network-Based Functional Connectivity in the Visual-Limbic System in Healthy Adults. Neuropsychopharmacology, 2018, 43, 406-414.	2.8	22
58	El estudio Andalusian Bipolar Family (ABiF): protocolo y descripciÃ³n de la muestra. Revista De PsiquiatrÃa Y Salud Mental, 2018, 11, 199-207.	1.0	5
59	Psychiatric Genomics: An Update and an Agenda. American Journal of Psychiatry, 2018, 175, 15-27.	4.0	518
60	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
61	Common functional variants of the glutamatergic system in Autism spectrum disorder with high and low intellectual abilities. Journal of Neural Transmission, 2018, 125, 259-271.	1.4	6
62	Gene set enrichment analysis and expression pattern exploration implicate an involvement of neurodevelopmental processes in bipolar disorder. Journal of Affective Disorders, 2018, 228, 20-25.	2.0	14
63	Detecting significant genotypeâ€“phenotype association rules in bipolar disorder: market research meets complex genetics. International Journal of Bipolar Disorders, 2018, 6, 24.	0.8	8
64	The Andalusian Bipolar Family (ABiF) Study: Protocol and sample description. Revista De PsiquiatrÃa Y Salud Mental (English Edition), 2018, 11, 199-207.	0.2	0
65	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. Nature Neuroscience, 2018, 21, 1656-1669.	7.1	490
66	Elevated expression of a minor isoform of ANK3 is a risk factor for bipolar disorder. Translational Psychiatry, 2018, 8, 210.	2.4	24
67	Exome sequencing in large, multiplex bipolar disorder families from Cuba. PLoS ONE, 2018, 13, e0205895.	1.1	13
68	Analysis of the Influence of microRNAs in Lithium Response in Bipolar Disorder. Frontiers in Psychiatry, 2018, 9, 207.	1.3	28
69	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	6.0	1,085
70	Shared genetic etiology between alcohol dependence and major depressive disorder. Psychiatric Genetics, 2018, 28, 66-70.	0.6	19
71	Novel genetic loci associated with hippocampal volume. Nature Communications, 2017, 8, 13624.	5.8	250
72	Genetics of schizophrenia: A consensus paper of the WFSBP Task Force on Genetics. World Journal of Biological Psychiatry, 2017, 18, 492-505.	1.3	48

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73	Influence of age and cognitive performance on resting-state brain networks of older adults in a population-based cohort. Cortex, 2017, 89, 28-44.	1.1	53
74	Significant Locus and Metabolic Genetic Correlations Revealed in Genome-Wide Association Study of Anorexia Nervosa. American Journal of Psychiatry, 2017, 174, 850-858.	4.0	410
75	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. Biological Psychiatry, 2017, 82, 322-329.	0.7	84
76	Polygenic Risk For BIP, MDD, And SCZ In Andalusian Multiplex Families. European Neuropsychopharmacology, 2017, 27, S385-S386.	0.3	0
77	T8. GLUTAMATERGIC SIGNALLING AND AUTISM: A FAMILY BASED ASSOCIATION STUDY ON THE GLUTAMATERGIC NEUROTRANSMITTER SYSTEM. European Neuropsychopharmacology, 2017, 27, S436-S437.	0.3	0
78	Expert and self-assessment of lifetime symptoms and diagnosis of major depressive disorder in large-scale genetic studies in the general population. Psychiatric Genetics, 2017, 27, 187-196.	0.6	10
79	Clinical Utility Gene Card for hereditary angioedema with normal C1 inhibitor (HAEnC1). European Journal of Human Genetics, 2017, 25, e1-e4.	1.4	9
80	Common variants at 2q11.2, 8q21.3, and 11q13.2 are associated with major mood disorders. Translational Psychiatry, 2017, 7, 1273.	2.4	9
81	Genome-wide mapping of genetic determinants influencing DNA methylation and gene expression in human hippocampus. Nature Communications, 2017, 8, 1511.	5.8	60
82	Association between neuropeptide Y receptor Y2 promoter variant rs6857715 and major depressive disorder. Psychiatric Genetics, 2017, 27, 34-37.	0.6	13
83	Functional Subclone Profiling for Prediction of Treatment-Induced Intratumor Population Shifts and Discovery of Rational Drug Combinations in Human Glioblastoma. Clinical Cancer Research, 2017, 23, 562-574.	3.2	60
84	Identification of a Bipolar Disorder Vulnerable Gene CHDH at 3p21.1. Molecular Neurobiology, 2017, 54, 5166-5176.	1.9	9
85	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
86	Consensus paper of the WFSBP Task Force on Genetics: Genetics, epigenetics and gene expression markers of major depressive disorder and antidepressant response. World Journal of Biological Psychiatry, 2017, 18, 5-28.	1.3	75
87	Analysis of the joint effect of SNPs to identify independent loci and allelic heterogeneity in schizophrenia GWAS data. Translational Psychiatry, 2017, 7, 1289.	2.4	4
88	Genetic Contribution to Alcohol Dependence: Investigation of a Heterogeneous German Sample of Individuals with Alcohol Dependence, Chronic Alcoholic Pancreatitis, and Alcohol-Related Cirrhosis. Genes, 2017, 8, 183.	1.0	11
89	Agonist Binding to Chemosensory Receptors: A Systematic Bioinformatics Analysis. Frontiers in Molecular Biosciences, 2017, 4, 63.	1.6	36
90	Identification of shared risk loci and pathways for bipolar disorder and schizophrenia. PLoS ONE, 2017, 12, e0171595.	1.1	77

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91	Leveraging Genomic Annotations and Pleiotropic Enrichment for Improved Replication Rates in Schizophrenia GWAS. PLoS Genetics, 2016, 12, e1005803.	1.5	34
92	Altered Functional Subnetwork During Emotional Face Processing. JAMA Psychiatry, 2016, 73, 598.	6.0	59
93	Novel genetic loci underlying human intracranial volume identified through genome-wide association. Nature Neuroscience, 2016, 19, 1569-1582.	7.1	213
94	Genome-wide association analyses identify new risk variants and the genetic architecture of amyotrophic lateral sclerosis. Nature Genetics, 2016, 48, 1043-1048.	9.4	494
95	Identification of rare variants in KCTD13 at the schizophrenia risk locus 16p11.2. Psychiatric Genetics, 2016, 26, 293-296.	0.6	5
96	Genome-wide association study of 40,000 individuals identifies two novel loci associated with bipolar disorder. Human Molecular Genetics, 2016, 25, 3383-3394.	1.4	182
97	Genetics of structural connectivity and information processing in the brain. Brain Structure and Function, 2016, 221, 4643-4661.	1.2	17
98	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
99	Genetic variants associated with response to lithium treatment in bipolar disorder: a genome-wide association study. Lancet, The, 2016, 387, 1085-1093.	6.3	306
100	Convergent Lines of Evidence Support LRP8 as a Susceptibility Gene for Psychosis. Molecular Neurobiology, 2016, 53, 6608-6619.	1.9	20
101	Impact of a cis-associated gene expression SNP on chromosome 20q11.22 on bipolar disorder susceptibility, hippocampal structure and cognitive performance. British Journal of Psychiatry, 2016, 208, 128-137.	1.7	11
102	Association of the OPRM1 Variant rs1799971 (A118G) with Non-Specific Liability to Substance Dependence in a Collaborative de novo Meta-Analysis of European-Ancestry Cohorts. Behavior Genetics, 2016, 46, 151-169.	1.4	98
103	No Reliable Association between Runs of Homozygosity and Schizophrenia in a Well-Powered Replication Study. PLoS Genetics, 2016, 12, e1006343.	1.5	24
104	A common risk variant in CACNA1C supports a sex-dependent effect on longitudinal functioning and functional recovery from episodes of schizophrenia-spectrum but not bipolar disorder. European Neuropsychopharmacology, 2015, 25, 2262-2270.	0.3	13
105	Investigation of the role of TCF4 rare sequence variants in schizophrenia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 354-362.	1.1	12
106	Meta-analysis identifies seven susceptibility loci involved in the atopic march. Nature Communications, 2015, 6, 8804.	5.8	148
107	Genetic Differences in the Immediate Transcriptome Response to Stress Predict Risk-Related Brain Function and Psychiatric Disorders. Neuron, 2015, 86, 1189-1202.	3.8	102
108	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

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109	Systems genetics identifies Sestrin 3 as a regulator of a proconvulsant gene network in human epileptic hippocampus. Nature Communications, 2015, 6, 6031.	5.8	158
110	Common genetic variants influence human subcortical brain structures. Nature, 2015, 520, 224-229.	13.7	772
111	Distinct Conditions Support a Novel Classification for Bradykinin-Mediated Angio-Oedema. Dermatology, 2015, 230, 324-331.	0.9	13
112	MicroRNA hsaâ€miRâ€4717â€5p regulates RGS2 and may be a risk factor for anxietyâ€related traits. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 296-306.	1.1	23
113	XRCC5 as a Risk Gene for Alcohol Dependence: Evidence from a Genome-Wide Gene-Set-Based Analysis and Follow-up Studies in Drosophila and Humans. Neuropsychopharmacology, 2015, 40, 361-371.	2.8	12
114	Systematic Integration of Brain eQTL and GWAS IdentifiesZNF323as a Novel Schizophrenia Risk Gene and Suggests Recent Positive Selection Based on Compensatory Advantage on Pulmonary Function. Schizophrenia Bulletin, 2015, 41, 1294-1308.	2.3	48
115	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
116	Multi-ancestry genome-wide association study of 21,000 cases and 95,000 controls identifies new risk loci for atopic dermatitis. Nature Genetics, 2015, 47, 1449-1456.	9.4	529
117	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
118	Genome-wide Studies of Verbal Declarative Memory in Nondemented Older People: The Cohorts for Heart and Aging Research in Genomic Epidemiology Consortium. Biological Psychiatry, 2015, 77, 749-763.	0.7	67
119	Studying variability in human brain aging in a population-based German cohortâ€”rationale and design of 1000BRAINS. Frontiers in Aging Neuroscience, 2014, 6, 149.	1.7	97
120	Genome-wide study of association and interaction with maternal cytomegalovirus infection suggests new schizophrenia loci. Molecular Psychiatry, 2014, 19, 325-333.	4.1	163
121	The KMO allele encoding Arg452 is associated with psychotic features in bipolar disorder type 1, and with increased CSF KYNA level and reduced KMO expression. Molecular Psychiatry, 2014, 19, 334-341.	4.1	91
122	Genome-wide analysis of rare copy number variations reveals PARK2 as a candidate gene for attention-deficit/hyperactivity disorder. Molecular Psychiatry, 2014, 19, 115-121.	4.1	76
123	Investigation of the involvement of MIR185 and its target genes in the development of schizophrenia. Journal of Psychiatry and Neuroscience, 2014, 39, 386-396.	1.4	23
124	Integrated Pathway-Based Approach Identifies Association between Genomic Regions at CTCF and CACNB2 and Schizophrenia. PLoS Genetics, 2014, 10, e1004345.	1.5	44
125	Striatal Response to Reward Anticipation. JAMA Psychiatry, 2014, 71, 531.	6.0	96
126	Identification of gene ontologies linked to prefrontalâ€“hippocampal functional coupling in the human brain. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 9657-9662.	3.3	9

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127	Efficient Strategy for Detecting Gene Ã— Gene Joint Action and Its Application in Schizophrenia. Genetic Epidemiology, 2014, 38, 60-71.	0.6	5
128	Further Evidence for the Impact of a Genome-Wide-Supported Psychosis Risk Variant in ZNF804A on the Theory of Mind Network. Neuropsychopharmacology, 2014, 39, 1196-1205.	2.8	42
129	Analysis of genome-wide significant bipolar disorder genes in borderline personality disorder. Psychiatric Genetics, 2014, 24, 262-265.	0.6	26
130	Smoking behaviour. Psychiatric Genetics, 2014, 24, 279-280.	0.6	1
131	A common microdeletion affecting a hippocampusâ€•and amygdalaâ€šspecific isoform of tryptophan hydroxylase 2 is not associated with affective disorders. Bipolar Disorders, 2014, 16, 764-768.	1.1	2
132	Epigenetic alteration of the dopamine transporter gene in alcoholâ€dependent patients is associated with age. Addiction Biology, 2014, 19, 305-311.	1.4	44
133	Molecular genetic overlap in bipolar disorder, schizophrenia, and major depressive disorder. World Journal of Biological Psychiatry, 2014, 15, 200-208.	1.3	120
134	Common variant at 16p11.2 conferring risk of psychosis. Molecular Psychiatry, 2014, 19, 108-114.	4.1	85
135	GENDER-SPECIFIC ASSOCIATION OF VARIANTS IN THEAKR1C1GENE WITH DIMENSIONAL ANXIETY IN PATIENTS WITH PANIC DISORDER: ADDITIONAL EVIDENCE FOR THE IMPORTANCE OF NEUROSTEROIDS IN ANXIETY?. Depression and Anxiety, 2014, 31, 843-850.	2.0	15
136	Genome-wide association study reveals two new risk loci for bipolar disorder. Nature Communications, 2014, 5, 3339.	5.8	294
137	The ENIGMA Consortium: large-scale collaborative analyses of neuroimaging and genetic data. Brain Imaging and Behavior, 2014, 8, 153-182.	1.1	696
138	FARVAT: a family-based rare variant association test. Bioinformatics, 2014, 30, 3197-3205.	1.8	34
139	ZNF804A and Cortical Structure in Schizophrenia: In Vivo and Postmortem Studies. Schizophrenia Bulletin, 2014, 40, 532-541.	2.3	28
140	Replication of brain function effects of a genome-wide supported psychiatric risk variant in the CACNA1C gene and new multi-locus effects. NeuroImage, 2014, 94, 147-154.	2.1	32
141	Hippocampal and Frontolimbic Function as Intermediate Phenotype for Psychosis: Evidence from Healthy Relatives and a Common Risk Variant in CACNA1C. Biological Psychiatry, 2014, 76, 466-475.	0.7	57
142	Common and Rare Variant Analysis in Early-Onset Bipolar Disorder Vulnerability. PLoS ONE, 2014, 9, e104326.	1.1	34
143	Bipolar disorder risk alleles in children with ADHD. Journal of Neural Transmission, 2013, 120, 1611-1617.	1.4	15
144	Levetiracetam resistance: Synaptic signatures & corresponding promoter SNPs in epileptic hippocampi. Neurobiology of Disease, 2013, 60, 115-125.	2.1	19

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145	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. Nature Genetics, 2013, 45, 984-994.	9.4	2,067
146	Contact System Activation in Patients with HAE and Normal C1ÂInhibitor Function. Immunology and Allergy Clinics of North America, 2013, 33, 513-533.	0.7	8
147	Implication of a Rare Deletion at Distal 16p11.2 in Schizophrenia. JAMA Psychiatry, 2013, 70, 253.	6.0	69
148	Genome-wide association data provide further support for an association between 5-HTTLPR and major depressive disorder. Journal of Affective Disorders, 2013, 146, 438-440.	2.0	24
149	A mega-analysis of genome-wide association studies for major depressive disorder. Molecular Psychiatry, 2013, 18, 497-511.	4.1	1,002
150	Rs6295 promoter variants of the serotonin type 1A receptor are differentially activated by c-Jun in vitro and correlate to transcript levels in human epileptic brain tissue. Brain Research, 2013, 1499, 136-144.	1.1	17
151	Genetic Schizophrenia Risk Variants Jointly Modulate Total Brain and White Matter Volume. Biological Psychiatry, 2013, 73, 525-531.	0.7	119
152	TLR4, ATF-3 and IL8 inflammation mediator expression correlates with seizure frequency in human epileptic brain tissue. Seizure: the Journal of the British Epilepsy Association, 2013, 22, 675-678.	0.9	74
153	Neuregulin 3 is associated with attention deficits in schizophrenia and bipolar disorder. International Journal of Neuropsychopharmacology, 2013, 16, 549-556.	1.0	30
154	Genome-wide association study meta-analysis of European and Asian-ancestry samples identifies three novel loci associated with bipolar disorder. Molecular Psychiatry, 2013, 18, 195-205.	4.1	180
155	Î±CaMKII Autophosphorylation Controls the Establishment of Alcohol Drinking Behavior. Neuropsychopharmacology, 2013, 38, 1636-1647.	2.8	63
156	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
157	No evidence for an involvement of copy number variation in ABCA13 in schizophrenia, bipolar disorder, or major depressive disorder. Psychiatric Genetics, 2013, 23, 45-46.	0.6	6
158	Distinct Loci in the CHRNA5/CHRNA3/CHRNB4 Gene Cluster Are Associated With Onset of Regular Smoking. Genetic Epidemiology, 2013, 37, 846-859.	0.6	32
159	Common obesity risk alleles in childhood attentionâ€deficit/hyperactivity disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2013, 162, 295-305.	1.1	77
160	Functional impact of a recently identified quantitative trait locus for hippocampal volume with genome-wide support. Translational Psychiatry, 2013, 3, e287-e287.	2.4	8
161	Identifying bipolar disorder susceptibility loci in a densely affected pedigree. Molecular Psychiatry, 2013, 18, 1245-1246.	4.1	21
162	Genome-wide association study implicates NDST3 in schizophrenia and bipolar disorder. Nature Communications, 2013, 4, 2739.	5.8	101

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163	Duplications in RB1CC1 are associated with schizophrenia; identification in large European sample sets. Translational Psychiatry, 2013, 3, e326-e326.	2.4	9
164	Copy Number Variants in German Patients with Schizophrenia. PLoS ONE, 2013, 8, e64035.	1.1	24
165	Enzymatic Assays for the Diagnosis of Bradykinin-Dependent Angioedema. PLoS ONE, 2013, 8, e70140.	1.1	54
166	A Genetic Deconstruction of Neurocognitive Traits in Schizophrenia and Bipolar Disorder. PLoS ONE, 2013, 8, e81052.	1.1	20
167	A Genome-Wide Association Study Suggests Novel Loci Associated with a Schizophrenia-Related Brain-Based Phenotype. PLoS ONE, 2013, 8, e64872.	1.1	21
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