Aiden P Corvin

List of Publications by Year
in descending order

180 papers	29,816 citations	20817 60 h-index	6131 159 g-index
202 all docs	202 docs citations	202 times ranked	35419 citing authors

#	Article	IF	Citations
1	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. Biological Psychiatry, 2022, 91, 102-117.	1.3	61
2	Minding metabolism: targeted interventions to improve cardio-metabolic monitoring across early and chronic psychosis. Irish Journal of Medical Science, 2022, 191, 337-346.	1.5	3
3	A <scp>metaâ€analysis</scp> of deep brain structural shape and asymmetry abnormalities in 2,833 individuals with schizophrenia compared with 3,929 healthy volunteers via the <scp>ENIGMA Consortium</scp>. Human Brain Mapping, 2022, 43, 352-372.	3.6	39
4	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. Biological Psychiatry, 2022, 91, 313-327.	1.3	114
5	Interaction Testing and Polygenic Risk Scoring to Estimate the Association of Common Genetic Variants With Treatment Resistance in Schizophrenia. JAMA Psychiatry, 2022, 79, 260.	11.0	44
6	Schizophrenia genomics. , 2022, , 17-41.		0
7	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	27.8	929
8	Exome sequencing in bipolar disorder identifies AKAP11 as a risk gene shared with schizophrenia. Nature Genetics, 2022, 54, 541-547.	21.4	65
9	Graph Convolutional Networks Reveal Network-Level Functional Dysconnectivity in Schizophrenia. Schizophrenia Bulletin, 2022, 48, 881-892.	4.3	18
10	Childhood trauma, parental bonding, and social cognition in patients with schizophrenia and healthy adults. Journal of Clinical Psychology, 2021, 77, 241-253.	1.9	22
11	DNA methylation meta-analysis reveals cellular alterations in psychosis and markers of treatment-resistant schizophrenia. ELife, 2021, 10, .	6.0	72
12	Changes in Default-Mode Network Associated With Childhood Trauma in Schizophrenia. Schizophrenia Bulletin, 2021, 47, 1482-1494.	4.3	18
13	Converting single nucleotide variants between genome builds: from cautionary tale to solution. Briefings in Bioinformatics, 2021, 22, .	6.5	15
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.	21.4	629
15	Identifying nootropic drug targets via large-scale cognitive GWAS and transcriptomics. Neuropsychopharmacology, 2021, 46, 1788-1801.	5.4	12
16	Early life Adversity, functional connectivity and cognitive performance in Schizophrenia: The mediating role of IL-6. Brain, Behavior, and Immunity, 2021, 98, 388-396.	4.1	21
17	Characterisation of age and polarity at onset in bipolar disorder. British Journal of Psychiatry, 2021, 219, 659-669.	2.8	20
18	Microglial-expressed genetic risk variants, cognitive function and brain volume in patients with schizophrenia and healthy controls. Translational Psychiatry, 2021, 11, 490.	4.8	10

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19	Investigating rare pathogenic/likely pathogenic exonic variation in bipolar disorder. Molecular Psychiatry, 2021, 26, 5239-5250.	7.9	15
20	Interleukin 6 predicts increased neural response during face processing in a sample of individuals with schizophrenia and healthy participants: A functional magnetic resonance imaging study. NeuroImage: Clinical, 2021, 32, 102851.	2.7	3
21	The Relationship Between Polygenic Risk Scores and Cognition in Schizophrenia. Schizophrenia Bulletin, 2020, 46, 336-344.	4.3	60
22	Dose response of the 16p11.2 distal copy number variant on intracranial volume and basal ganglia. Molecular Psychiatry, 2020, 25, 584-602.	7.9	49
23	Detecting schizophrenia at the level of the individual: relative diagnostic value of whole-brain images, connectome-wide functional connectivity and graph-based metrics. Psychological Medicine, 2020, 50, 1852-1861.	4.5	57
24	Association of Copy Number Variation of the 15q11.2 BP1-BP2 Region With Cortical and Subcortical Morphology and Cognition. JAMA Psychiatry, 2020, 77, 420.	11.0	54
25	Integrating machining learning and multimodal neuroimaging to detect schizophrenia at the level of the individual. Human Brain Mapping, 2020, 41, 1119-1135.	3.6	56
26	Effects of complement geneâ€set polygenic risk score on brain volume and cortical measures in patients with psychotic disorders and healthy controls. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2020, 183, 445-453.	1.7	6
27	Childhood trauma, brain structure and emotion recognition in patients with schizophrenia and healthy participants. Social Cognitive and Affective Neuroscience, 2020, 15, 1325-1339.	3.0	26
28	Prevalence of N-Methyl-d-Aspartate Receptor antibody (NMDAR-Ab) encephalitis in patients with first episode psychosis and treatment resistant schizophrenia on clozapine, a population based study. Schizophrenia Research, 2020, 222, 455-461.	2.0	17
29	Deficit not bias: A quantifiable neuropsychological model of delusions. Schizophrenia Research, 2020, 222, 496-498.	2.0	2
30	The genetic architecture of the human cerebral cortex. Science, 2020, 367, .	12.6	450
31	Neuroharmony: A new tool for harmonizing volumetric MRI data from unseen scanners. NeuroImage, 2020, 220, 117127.	4.2	48
32	Identifying schizophrenia patients who carry pathogenic genetic copy number variants using standard clinical assessment: retrospective cohort study. British Journal of Psychiatry, 2020, 216, 275-279.	2.8	12
33	Methyl-CpG-binding protein 2 mediates overlapping mechanisms across brain disorders. Scientific Reports, 2020, 10, 22255.	3.3	10
34	Beyond C4: Analysis of the complement gene pathway shows enrichment for IQ in patients with psychotic disorders and healthy controls. Genes, Brain and Behavior, 2019, 18, e12602.	2.2	13
35	Pleiotropic Meta-Analysis of Cognition, Education, and Schizophrenia Differentiates Roles of Early Neurodevelopmental and Adult Synaptic Pathways. American Journal of Human Genetics, 2019, 105, 334-350.	6.2	86
36	Autoantibodies and Psychosis. Current Topics in Behavioral Neurosciences, 2019, 44, 85-123.	1.7	6

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37	GWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. American Journal of Psychiatry, 2019, 176, 651-660.	7.2	186
38	Cortical patterning of abnormal morphometric similarity in psychosis is associated with brain expression of schizophrenia-related genes. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 9604-9609.	7.1	200
39	Genome-wide association study identifies 30 loci associated with bipolar disorder. Nature Genetics, 2019, 51, 793-803.	21.4	1,191
40	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.	1.7	2
41	Targeted Sequencing of 10,198 Samples Confirms Abnormalities in Neuronal Activity and Implicates Voltage-Gated Sodium Channels in Schizophrenia Pathogenesis. Biological Psychiatry, 2019, 85, 554-562.	1.3	40
42	A randomized controlled trial of cognitive remediation for a national cohort of forensic patients with schizophrenia or schizoaffective disorder. BMC Psychiatry, 2019, 19, 27.	2.6	30
43	Effects of MiRâ€137 genetic risk score on brain volume and cortical measures in patients with schizophrenia and controls. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2018, 177, 369-376.	1.7	10
44	Moral cognition and homicide amongst forensic patients with schizophrenia and schizoaffective disorder: A cross-sectional cohort study. Schizophrenia Research, 2018, 193, 468-469.	2.0	8
45	Study of 300,486 individuals identifies 148 independent genetic loci influencing general cognitive function. Nature Communications, 2018, 9, 2098.	12.8	484
46	Computerised working memoryâ€based cognitive remediation therapy does not affect Reading the Mind in The Eyes test performance or neural activity during a Facial Emotion Recognition test in psychosis. European Journal of Neuroscience, 2018, 48, 1691-1705.	2.6	4
47	Estimation of Genetic Correlation via Linkage Disequilibrium Score Regression and Genomic Restricted Maximum Likelihood. American Journal of Human Genetics, 2018, 102, 1185-1194.	6.2	119
48	Cortical Brain Abnormalities in 4474 Individuals With Schizophrenia and 5098 Control Subjects via the Enhancing Neuro Imaging Genetics Through Meta Analysis (ENIGMA) Consortium. Biological Psychiatry, 2018, 84, 644-654.	1.3	627
49	Genome-wide association meta-analysis in 269,867 individuals identifies new genetic and functional links to intelligence. Nature Genetics, 2018, 50, 912-919.	21.4	893
50	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	12.6	1,085
51	Multi-Trait Analysis of GWAS and Biological Insights Into Cognition: A Response to Hill (2018). Twin Research and Human Genetics, 2018, 21, 394-397.	0.6	3
52	The Genetics of Endophenotypes of Neurofunction to Understand Schizophrenia (GENUS) consortium: A collaborative cognitive and neuroimaging genetics project. Schizophrenia Research, 2018, 195, 306-317.	2.0	17
53	Novel genetic loci associated with hippocampal volume. Nature Communications, 2017, 8, 13624.	12.8	250
54	Occurrence and co-occurrence of hallucinations by modality in schizophrenia-spectrum disorders. Psychiatry Research, 2017, 252, 154-160.	3.3	96

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55	Further evidence of alerted default network connectivity and association with theory of mind ability in schizophrenia. Schizophrenia Research, 2017, 184, 52-58.	2.0	26
56	Cognitive Characterization of Schizophrenia Risk Variants Involved in Synaptic Transmission: Evidence of CACNA1C's Role in Working Memory. Neuropsychopharmacology, 2017, 42, 2612-2622.	5.4	28
57	Large-Scale Cognitive GWAS Meta-Analysis Reveals Tissue-Specific Neural Expression and Potential Nootropic Drug Targets. Cell Reports, 2017, 21, 2597-2613.	6.4	103
58	Human subcortical brain asymmetries in 15,847 people worldwide reveal effects of age and sex. Brain Imaging and Behavior, 2017, 11, 1497-1514.	2.1	144
59	Fluorescent nanodiamond tracking reveals intraneuronal transport abnormalities induced by brain-disease-related genetic risk factors. Nature Nanotechnology, 2017, 12, 322-328.	31.5	111
60	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. Nature Genetics, 2017, 49, 27-35.	21.4	838
61	Genetics of Schizophrenia: Ready to Translate?. Current Psychiatry Reports, 2017, 19, 61.	4.5	52
62	Expression of nuclear Methyl-CpG binding protein 2 (Mecp2) is dependent on neuronal stimulation and application of Insulin-like growth factor 1. Neuroscience Letters, 2016, 621, 111-116.	2.1	13
63	Data science for mental health: a UK perspective on a global challenge. Lancet Psychiatry,the, 2016, 3, 993-998.	7.4	47
64	Novel genetic loci underlying human intracranial volume identified through genome-wide association. Nature Neuroscience, 2016, 19, 1569-1582.	14.8	213
65	Cognitive analysis of schizophrenia risk genes that function as epigenetic regulators of gene expression. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2016, 171, 1170-1179.	1.7	43
66	Polymorphism in a lincRNA Associates with a Doubled Risk of Pneumococcal Bacteremia in Kenyan Children. American Journal of Human Genetics, 2016, 98, 1092-1100.	6.2	39
67	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.	1.7	28
68	Psychiatric genetics: what's new in 2015?. Lancet Psychiatry,the, 2016, 3, 10-12.	7.4	2
69	What Next in Schizophrenia Genetics for the Psychiatric Genomics Consortium?. Schizophrenia Bulletin, 2016, 42, 538-541.	4.3	23
70	Common polygenic variation in coeliac disease and confirmation of ZNF335 and NIFA as disease susceptibility loci. European Journal of Human Genetics, 2016, 24, 291-297.	2.8	25
71	Independent evidence for an association between general cognitive ability and a genetic locus for educational attainment. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 363-373.	1.7	25
72	Biomarkers for Psychosis: the Molecular Genetics of Psychosis. Current Behavioral Neuroscience Reports, 2015, 2, 112-118.	1.3	1

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73	Genome-wide Comparative Analysis of Atopic Dermatitis and Psoriasis Gives Insight into Opposing Genetic Mechanisms. American Journal of Human Genetics, 2015, 96, 104-120.	6.2	163
74	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.	6.2	225
75	Common genetic variants influence human subcortical brain structures. Nature, 2015, 520, 224-229.	27.8	772
76	MIR137HG risk variant rs1625579 genotype is related to corpus callosum volume in schizophrenia. Neuroscience Letters, 2015, 602, 44-49.	2.1	18
77	Greater number of older siblings is associated with decreased theory of mind ability in psychosis. Schizophrenia Research, 2015, 165, 247-248.	2.0	0
78	Modeling Linkage Disequilibrium Increases Accuracy of Polygenic Risk Scores. American Journal of Human Genetics, 2015, 97, 576-592.	6.2	1,098
79	New data and an old puzzle: the negative association between schizophrenia and rheumatoid arthritis. International Journal of Epidemiology, 2015, 44, 1706-1721.	1.9	53
80	Repeated Insulin-Like Growth Factor 1 Treatment in a Patient with Rett Syndrome: A Single Case Study. Frontiers in Pediatrics, 2014, 2, 52.	1.9	28
81	The miRâ€137 schizophrenia susceptibility variant rs1625579 does not predict variability in brain volume in a sample of schizophrenic patients and healthy individuals. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 467-471.	1.7	17
82	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.	2.9	37
83	CNV analysis in a large schizophrenia sample implicates deletions at 16p12.1 and SLC1A1 and duplications at 1p36.33 and CGNL1. Human Molecular Genetics, 2014, 23, 1669-1676.	2.9	82
84	Effects of MIR137 on fronto-amygdala functional connectivity. NeuroImage, 2014, 90, 189-195.	4.2	42
85	Genetic modifiers and subtypes in schizophrenia: Investigations of age at onset, severity, sex and family history. Schizophrenia Research, 2014, 154, 48-53.	2.0	68
86	No evidence that runs of homozygosity are associated with schizophrenia in an Irish genome-wide association dataset. Schizophrenia Research, 2014, 154, 79-82.	2.0	18
87	Variability in Working Memory Performance Explained by Epistasis vs Polygenic Scores in theZNF804APathway. JAMA Psychiatry, 2014, 71, 778.	11.0	28
88	Unlocking the Treasure Trove: From Genes to Schizophrenia Biology. Schizophrenia Bulletin, 2014, 40, 492-496.	4.3	19
89	Partitioning Heritability of Regulatory and Cell-Type-Specific Variants across 11 Common Diseases. American Journal of Human Genetics, 2014, 95, 535-552.	6.2	569
90	Polygenic dissection of diagnosis and clinical dimensions of bipolar disorder and schizophrenia. Molecular Psychiatry, 2014, 19, 1017-1024.	7.9	333

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91	Effects of a novel schizophrenia risk variant rs7914558 atCNNM2on brain structure and attributional style. British Journal of Psychiatry, 2014, 204, 115-121.	2.8	30
92	The correlation between reading and mathematics ability at age twelve has a substantial genetic component. Nature Communications, 2014, 5, 4204.	12.8	72
93	The phenotypic manifestations of rare CNVs in schizophrenia. Schizophrenia Research, 2014, 158, 255-260.	2.0	21
94	Genome-wide schizophrenia variant at MIR137 does not impact white matter microstructure in healthy participants. Neuroscience Letters, 2014, 574, 6-10.	2.1	15
95	Analysis of the hexanucleotide repeat expansion and founder haplotype at C9ORF72 in an Irish psychosis case-control sample. Neurobiology of Aging, 2014, 35, 1510.e1-1510.e5.	3.1	20
96	A Genome-wide Association Analysis of a Broad Psychosis Phenotype Identifies Three Loci for Further Investigation. Biological Psychiatry, 2014, 75, 386-397.	1.3	44
97	Altered medial prefrontal activity during dynamic face processing in schizophrenia spectrum patients. Schizophrenia Research, 2014, 157, 225-230.	2.0	30
98	Genome-wide Association Studies: Findings at the Major Histocompatibility Complex Locus in Psychosis. Biological Psychiatry, 2014, 75, 276-283.	1.3	115
99	Genome-wide association analysis identifies 13 new risk loci for schizophrenia. Nature Genetics, 2013, 45, 1150-1159.	21.4	1,395
100	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. Nature Genetics, 2013, 45, 984-994.	21.4	2,067
101	Implication of a Rare Deletion at Distal 16p11.2 in Schizophrenia. JAMA Psychiatry, 2013, 70, 253.	11.0	69
102	Mosaic copy number variation in schizophrenia. European Journal of Human Genetics, 2013, 21, 1007-1011.	2.8	15
103	Common variants in the HLA-DRB1â€“HLA-DQA1 HLA class II region are associated with susceptibility to visceral leishmaniasis. Nature Genetics, 2013, 45, 208-213.	21.4	86
104	Social dysfunction in schizophrenia: An investigation of the GAF scale's sensitivity to deficits in social cognition. Schizophrenia Research, 2013, 146, 363-365.	2.0	25
105	A Comprehensive Family-Based Replication Study of Schizophrenia Genes. JAMA Psychiatry, 2013, 70, 573.	11.0	138
106	The Role of the Major Histocompatibility Complex Region in Cognition and Brain Structure: A Schizophrenia GWAS Follow-Up. American Journal of Psychiatry, 2013, 170, 877-885.	7.2	60
107	Genome-wide association study of intraocular pressure identifies the GLCCI1/ICA1 region as a glaucoma susceptibility locus. Human Molecular Genetics, 2013, 22, 4653-4660.	2.9	29
108	Schizophrenia at a Genetics Crossroads: Where to Now?. Schizophrenia Bulletin, 2013, 39, 490-495.	4.3	12

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109	Neural effects of the <scp>CSMD</scp>1 genomeâ€wide associated schizophrenia risk variant rs10503253. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2013, 162, 530-537.	1.7	30
110	Development of Strategies for SNP Detection in RNA-Seq Data: Application to Lymphoblastoid Cell Lines and Evaluation Using 1000 Genomes Data. PLoS ONE, 2013, 8, e58815.	2.5	108
111	The Effect of the Neurogranin Schizophrenia Risk Variant rs12807809 on Brain Structure and Function. Twin Research and Human Genetics, 2012, 15, 296-303.	0.6	26
112	Functional investigation of a schizophrenia GWAS signal at the CDC42 gene. World Journal of Biological Psychiatry, 2012, 13, 550-554.	2.6	18
113	Common variants at the MHC locus and at chromosome 16q24.1 predispose to Barrett's esophagus. Nature Genetics, 2012, 44, 1131-1136.	21.4	162
114	Genome-wide association study identifies a variant in HDAC9 associated with large vessel ischemic stroke. Nature Genetics, 2012, 44, 328-333.	21.4	375
115	Identification of common variants associated with human hippocampal and intracranial volumes. Nature Genetics, 2012, 44, 552-561.	21.4	594
116	Insulin-like growth factor 1 (IGF1) and its active peptide (1â€“3)IGF1 enhance the expression of synaptic markers in neuronal circuits through different cellular mechanisms. Neuroscience Letters, 2012, 520, 51-56.	2.1	49
117	Social cognition in bipolar disorder versus schizophrenia: comparability in mental state decoding deficits. Bipolar Disorders, 2012, 14, 743-748.	1.9	44
118	The NOS1 variant rs6490121 is associated with variation in prefrontal function and grey matter density in healthy individuals. NeuroImage, 2012, 60, 614-622.	4.2	26
119	A NOS1 variant implicated in cognitive performance influences evoked neural responses during a high density EEG study of early visual perception. Human Brain Mapping, 2012, 33, 1202-1211.	3.6	19
120	Psychiatric genetics. , 2012, , 35-53.		1
121	Genetic risk and a primary role for cell-mediated immune mechanisms in multiple sclerosis. Nature, 2011, 476, 214-219.	27.8	2,400
122	A neuropsychological investigation of the genome wide associated schizophrenia risk variant NRGN rs12807809. Schizophrenia Research, 2011, 125, 304-306.	2.0	23
123	ZNF804A risk allele is associated with relatively intact gray matter volume in patients with schizophrenia. NeuroImage, 2011, 54, 2132-2137.	4.2	78
124	High Frequencies of De Novo CNVs in Bipolar Disorder and Schizophrenia. Neuron, 2011, 72, 951-963.	8.1	290
125	Duplications of the neuropeptide receptor gene VIPR2 confer significant risk for schizophrenia. Nature, 2011, 471, 499-503.	27.8	296
126	Genetic Classification of Populations Using Supervised Learning. PLoS ONE, 2011, 6, e14802.	2.5	16

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127	Common variants near ATM are associated with glycemic response to metformin in type 2 diabetes. Nature Genetics, 2011, 43, 117-120.	21.4	390
128	Interaction between ERAP1 and HLA-B27 in ankylosing spondylitis implicates peptide handling in the mechanism for HLA-B27 in disease susceptibility. Nature Genetics, 2011, 43, 761-767.	21.4	778
129	Two patients walk into a clinic...a genomics perspective on the future of schizophrenia. BMC Biology, 2011, 9, 77.	3.8	11
130	Allelic expression imbalance of the schizophrenia susceptibility gene CHI3L1. Psychiatric Genetics, 2011, 21, 281-286.	1.1	2
131	Multiplex Target Enrichment Using DNA Indexing for Ultra-High Throughput SNP Detection. DNA Research, 2011, 18, 31-38.	3.4	41
132	Dissection of the genetics of Parkinson's disease identifies an additional association 5' of SNCA and multiple associated haplotypes at 17q21. Human Molecular Genetics, 2011, 20, 345-353.	2.9	202
133	Mutation of Semaphorin-6A Disrupts Limbic and Cortical Connectivity and Models Neurodevelopmental Psychopathology. PLoS ONE, 2011, 6, e26488.	2.5	40
134	The Shock of the New: Progress in Schizophrenia Genomics. Current Genomics, 2011, 12, 516-524.	1.6	16
135	Population structure and genome-wide patterns of variation in Ireland and Britain. European Journal of Human Genetics, 2010, 18, 1248-1254.	2.8	46
136	Genes predict village of origin in rural Europe. European Journal of Human Genetics, 2010, 18, 1269-1270.	2.8	22
137	A genome-wide association study identifies new psoriasis susceptibility loci and an interaction between HLA-C and ERAP1. Nature Genetics, 2010, 42, 985-990.	21.4	918
138	Evidence for cisâ€acting regulation of ANK3 and CACNA1C gene expression. Bipolar Disorders, 2010, 12, 440-445.	1.9	31
139	Psychosis Susceptibility Gene ZNF804A and Cognitive Performance in Schizophrenia. Archives of General Psychiatry, 2010, 67, 692.	12.3	129
140	Reduced Occipital and Prefrontal Brain Volumes in Dysbindin-Associated Schizophrenia. Neuropsychopharmacology, 2010, 35, 368-373.	5.4	29
141	Genetic Differences between Five European Populations. Human Heredity, 2010, 70, 141-149.	0.8	29
142	Neuronal cell adhesion genes. Cell Adhesion and Migration, 2010, 4, 511-514.	2.7	29
143	The Psychosis Susceptibility Gene ZNF804A: Associations, Functions, and Phenotypes. Schizophrenia Bulletin, 2010, 36, 904-909.	4.3	51
144	Replicated genetic evidence supports a role for HOMER2 in schizophrenia. Neuroscience Letters, 2010, 468, 229-233.	2.1	18

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145	Is â€œclinicalâ€•insight the same as â€œcognitiveâ€•insight in schizophrenia?. Journal of the International Neuropsychological Society, 2009, 15, 471-475.	1.8	32
146	Influence of NOS1 on Verbal Intelligence and Working Memory in Both Patients With Schizophrenia and Healthy Control Subjects. Archives of General Psychiatry, 2009, 66, 1045.	12.3	45
147	The SNP ratio test: pathway analysis of genome-wide association datasets. Bioinformatics, 2009, 25, 2762-2763.	4.1	125
148	Copy-number variants in neurodevelopmental disorders: promises and challenges. Trends in Genetics, 2009, 25, 536-544.	6.7	120
149	Genome-wide association study of ulcerative colitis identifies three new susceptibility loci, including the HNF4A region. Nature Genetics, 2009, 41, 1330-1334.	21.4	483
150	Moodâ€•incongruent psychosis in bipolar disorder: conditional linkage analysis shows genomeâ€•wide suggestive linkage at 1q32.3, 7p13 and 20q13.31. Bipolar Disorders, 2009, 11, 610-620.	1.9	23
151	Does the ability to sustain attention underlie symptom severity in schizophrenia?. Schizophrenia Research, 2009, 107, 319-323.	2.0	26
152	Identification of loci associated with schizophrenia by genome-wide association and follow-up. Nature Genetics, 2008, 40, 1053-1055.	21.4	977
153	Collaborative genome-wide association analysis supports a role for ANK3 and CACNA1C in bipolar disorder. Nature Genetics, 2008, 40, 1056-1058.	21.4	1,102
154	An assessment of the Irish population for large-scale genetic mapping studies involving epilepsy and other complex diseases. European Journal of Human Genetics, 2008, 16, 176-183.	2.8	5
155	Familial patterns and the origins of individual differences in synaesthesia. Cognition, 2008, 106, 871-893.	2.2	144
156	Mental state decoding v. mental state reasoning as a mediator between cognitive and social function in psychosis. British Journal of Psychiatry, 2008, 193, 77-78.	2.8	61
157	Dysbindin (DTNBP1) and the Biogenesis of Lysosome-Related Organelles Complex 1 (BLOC-1): Main and Epistatic Gene Effects Are Potential Contributors to Schizophrenia Susceptibility. Biological Psychiatry, 2008, 63, 24-31.	1.3	54
158	Early Visual Processing Deficits in Dysbindin-Associated Schizophrenia. Biological Psychiatry, 2008, 63, 484-489.	1.3	62
159	Chitinase-3-Like 1 (CHI3L1) Gene and Schizophrenia: Genetic Association and a Potential Functional Mechanism. Biological Psychiatry, 2008, 64, 98-103.	1.3	28
160	A dysbindin risk haplotype associated with less severe manic-type symptoms in psychosis. Neuroscience Letters, 2008, 431, 146-149.	2.1	30
161	Are relational style and neuropsychological performance predictors of social attributions in chronic schizophrenia?. Psychiatry Research, 2008, 161, 19-27.	3.3	33
162	Physical health and attendance at primary care in people with schizophrenia. Irish Journal of Psychological Medicine, 2008, 25, 57-60.	1.0	2

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163	d-Amino acid oxidase (DAO) genotype and mood symptomatology in schizophrenia. Neuroscience Letters, 2007, 426, 97-100.	2.1	26
164	Functional Genomics and Schizophrenia: Endophenotypes and Mutant Models. Psychiatric Clinics of North America, 2007, 30, 365-399.	1.3	40
165	Variance in facial recognition performance associated with BDNF in schizophrenia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2007, 144B, 578-579.	1.7	6
166	Variance in neurocognitive performance is associated with dysbindin-1 in schizophrenia: A preliminary study. Neuropsychologia, 2007, 45, 454-458.	1.6	109
167	Evaluation of a Susceptibility Gene for Schizophrenia: Genotype Based Meta-Analysis of RGS4 Polymorphisms from Thirteen Independent Samples. Biological Psychiatry, 2006, 60, 152-162.	1.3	87
168	Neurocognition and suicidal behaviour in an Irish population with major psychotic disorders. Schizophrenia Research, 2006, 85, 196-200.	2.0	69
169	Are deficits in executive sub-processes simply reflecting more general cognitive decline in schizophrenia?. Schizophrenia Research, 2006, 85, 168-173.	2.0	24
170	Do antisaccade deficits in schizophrenia provide evidence of a specific inhibitory function?. Journal of the International Neuropsychological Society, 2006, 12, 901-6.	1.8	15
171	Evidence that interaction between neuregulin 1 and its receptor erbB4 increases susceptibility to schizophrenia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2006, 141B, 96-101.	1.7	162
172	Evidence that specific executive functions predict symptom variance among schizophrenia patients with a predominantly negative symptom profile. Cognitive Neuropsychiatry, 2006, 11, 13-32.	1.3	47
173	Are the Cognitive Deficits Associated With Impaired Insight in Schizophrenia Specific to Executive Task Performance?. Journal of Nervous and Mental Disease, 2005, 193, 803-808.	1.0	40
174	Genomewide Linkage Scan in Schizoaffective Disorder. Archives of General Psychiatry, 2005, 62, 1081.	12.3	177
175	Investigation of the apolipoprotein-L (APOL) gene family and schizophrenia using a novel DNA pooling strategy for public database SNPs. Schizophrenia Research, 2005, 76, 231-238.	2.0	12
176	ConfirmingRGS4 as a susceptibility gene for schizophrenia. American Journal of Medical Genetics Part A, 2004, 125B, 50-53.	2.4	125
177	No evidence for association of the dysbindin gene [DTNBP1] with schizophrenia in an Irish population-based study. Schizophrenia Research, 2003, 60, 167-172.	2.0	85
178	Psychiatric genetics in the post-genome age. British Journal of Psychiatry, 2003, 182, 95-96.	2.8	0
179	The attitudes of Irish trainees to their training and its supervision: a five-year follow up study. Irish Journal of Psychological Medicine, 2001, 18, 120-125.	1.0	6
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