Srdjan Djurovic

List of Publications by Year
in descending order

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280 papers	35,621 citations	11608 70 h-index	4978 167 g-index
332 all docs	332 docs citations	332 times ranked	35919 citing authors

#	Article	IF	Citations
1	Variant ofTREM2Associated with the Risk of Alzheimer's Disease. New England Journal of Medicine, 2013, 368, 107-116.	13.9	2,085
2	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. Nature Genetics, 2013, 45, 984-994.	9.4	2,067
3	Genome-wide meta-analysis identifies new loci and functional pathways influencing Alzheimerâ€™s disease risk. Nature Genetics, 2019, 51, 404-413.	9.4	1,625
4	Large recurrent microdeletions associated with schizophrenia. Nature, 2008, 455, 232-236.	13.7	1,619
5	Common variants conferring risk of schizophrenia. Nature, 2009, 460, 744-747.	13.7	1,572
6	Identification of common genetic risk variants for autism spectrum disorder. Nature Genetics, 2019, 51, 431-444.	9.4	1,538
7	Common schizophrenia alleles are enriched in mutation-intolerant genes and in regions under strong background selection. Nature Genetics, 2018, 50, 381-389.	9.4	1,332
8	Genome-wide association study identifies 30 loci associated with bipolar disorder. Nature Genetics, 2019, 51, 793-803.	9.4	1,191
9	Modeling Linkage Disequilibrium Increases Accuracy of Polygenic Risk Scores. American Journal of Human Genetics, 2015, 97, 576-592.	2.6	1,098
10	Genome-wide association meta-analysis in 269,867 individuals identifies new genetic and functional links to intelligence. Nature Genetics, 2018, 50, 912-919.	9.4	893
11	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
12	Common genetic variants influence human subcortical brain structures. Nature, 2015, 520, 224-229.	13.7	772
13	New insights into the genetic etiology of Alzheimerâ€™s disease and related dementias. Nature Genetics, 2022, 54, 412-436.	9.4	700
14	The ENIGMA Consortium: large-scale collaborative analyses of neuroimaging and genetic data. Brain Imaging and Behavior, 2014, 8, 153-182.	1.1	696
15	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
16	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. Cell, 2018, 173, 1705-1715.e16.	13.5	623
17	Identification of common variants associated with human hippocampal and intracranial volumes. Nature Genetics, 2012, 44, 552-561.	9.4	594
18	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

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19	Study of 300,486 individuals identifies 148 independent genetic loci influencing general cognitive function. Nature Communications, 2018, 9, 2098.	5.8	484
20	The genetic architecture of the human cerebral cortex. Science, 2020, 367, .	6.0	450
21	A genome-wide association study with 1,126,563 individuals identifies new risk loci for Alzheimerâ€™s disease. Nature Genetics, 2021, 53, 1276-1282.	9.4	430
22	Disruption of the neurexin 1 gene is associated with schizophrenia. Human Molecular Genetics, 2009, 18, 988-996.	1.4	424
23	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
24	Common brain disorders are associated with heritable patterns of apparent aging of the brain. Nature Neuroscience, 2019, 22, 1617-1623.	7.1	358
25	Polygenic dissection of diagnosis and clinical dimensions of bipolar disorder and schizophrenia. Molecular Psychiatry, 2014, 19, 1017-1024.	4.1	333
26	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
27	Loss-of-function variants in ABCA7 confer risk of Alzheimer's disease. Nature Genetics, 2015, 47, 445-447.	9.4	283
28	Genome-wide Association Study Identifies Genetic Variation in Neurocan as a Susceptibility Factor for Bipolar Disorder. American Journal of Human Genetics, 2011, 88, 372-381.	2.6	257
29	Novel genetic loci associated with hippocampal volume. Nature Communications, 2017, 8, 13624.	5.8	250
30	Genome-wide Pleiotropy Between Parkinson Disease and Autoimmune Diseases. JAMA Neurology, 2017, 74, 780.	4.5	245
31	Brain Expressed microRNAs Implicated in Schizophrenia Etiology. PLoS ONE, 2007, 2, e873.	1.1	235
32	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
33	Novel genetic loci underlying human intracranial volume identified through genome-wide association. Nature Neuroscience, 2016, 19, 1569-1582.	7.1	213
34	Brain Heterogeneity in Schizophrenia and Its Association With Polygenic Risk. JAMA Psychiatry, 2019, 76, 739.	6.0	195
35	Common variants at VRK2 and TCF4 conferring risk of schizophrenia. Human Molecular Genetics, 2011, 20, 4076-4081.	1.4	193
36	Genetic architecture of subcortical brain structures in 38,851 individuals. Nature Genetics, 2019, 51, 1624-1636.	9.4	192

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37	Bivariate causal mixture model quantifies polygenic overlap between complex traits beyond genetic correlation. Nature Communications, 2019, 10, 2417.	5.8	190
38	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
39	Gene variants associated with schizophrenia in a Norwegian genome-wide study are replicated in a large European cohort. Journal of Psychiatric Research, 2010, 44, 748-753.	1.5	183
40	Cell type specificity of neurovascular coupling in cerebral cortex. ELife, 2016, 5, .	2.8	176
41	Genome-wide analysis reveals extensive genetic overlap between schizophrenia, bipolar disorder, and intelligence. Molecular Psychiatry, 2020, 25, 844-853.	4.1	156
42	The Complement Control-Related Genes CSMD1 and CSMD2 Associate to Schizophrenia. Biological Psychiatry, 2011, 70, 35-42.	0.7	149
43	Polygenic Overlap Between C-Reactive Protein, Plasma Lipids, and Alzheimer Disease. Circulation, 2015, 131, 2061-2069.	1.6	145
44	Deep 2-photon imaging and artifact-free optogenetics through transparent graphene microelectrode arrays. Nature Communications, 2018, 9, 2035.	5.8	143
45	Common variants in Alzheimerâ€™s disease and risk stratification by polygenic risk scores. Nature Communications, 2021, 12, 3417.	5.8	140
46	A Comprehensive Family-Based Replication Study of Schizophrenia Genes. JAMA Psychiatry, 2013, 70, 573.	6.0	138
47	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
48	Identification of Genetic Loci Jointly Influencing Schizophrenia Risk and the Cognitive Traits of Verbal-Numerical Reasoning, Reaction Time, and General Cognitive Function. JAMA Psychiatry, 2017, 74, 1065.	6.0	123
49	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
50	Sex-dependent association of common variants of microcephaly genes with brain structure. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 384-388.	3.3	118
51	At-Risk Variant in TCF7L2 for Type II Diabetes Increases Risk of Schizophrenia. Biological Psychiatry, 2011, 70, 59-63.	0.7	114
52	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. Biological Psychiatry, 2022, 91, 313-327.	0.7	114
53	Discovery of shared genomic loci using the conditional false discovery rate approach. Human Genetics, 2020, 139, 85-94.	1.8	109
54	A common MECP2 haplotype associates with reduced cortical surface area in humans in two independent populations. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 15483-15488.	3.3	108

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55	Large-Scale Cognitive GWAS Meta-Analysis Reveals Tissue-Specific Neural Expression and Potential Nootropic Drug Targets. Cell Reports, 2017, 21, 2597-2613.	2.9	103
56	A Comparison of Ten Polygenic Score Methods for Psychiatric Disorders Applied Across Multiple Cohorts. Biological Psychiatry, 2021, 90, 611-620.	0.7	103
57	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
58	A genome-wide association study of bipolar disorder in Norwegian individuals, followed by replication in Icelandic sample. Journal of Affective Disorders, 2010, 126, 312-316.	2.0	100
59	Interplay between childhood trauma and BDNF val66met variants on blood BDNF mRNA levels and on hippocampus subfields volumes in schizophrenia spectrum and bipolar disorders. Journal of Psychiatric Research, 2014, 59, 14-21.	1.5	97
60	Maternally Derived Microduplications at 15q11-q13: Implication of Imprinted Genes in Psychotic Illness. American Journal of Psychiatry, 2011, 168, 408-417.	4.0	95
61	Exploring the Wnt signaling pathway in schizophrenia and bipolar disorder. Translational Psychiatry, 2018, 8, 55.	2.4	94
62	Genetic Markers of Human Evolution Are Enriched in Schizophrenia. Biological Psychiatry, 2016, 80, 284-292.	0.7	92
63	Distinct multivariate brain morphological patterns and their added predictive value with cognitive and polygenic risk scores in mental disorders. NeuroImage: Clinical, 2017, 15, 719-731.	1.4	89
64	BDNF val66met modulates the association between childhood trauma, cognitive and brain abnormalities in psychoses. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2013, 46, 181-188.	2.5	87
65	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.	2.6	86
66	Inflammatory markers are associated with general cognitive abilities in schizophrenia and bipolar disorder patients and healthy controls. Schizophrenia Research, 2015, 165, 188-194.	1.1	85
67	Genetic Overlap Between Schizophrenia and Volumes of Hippocampus, Putamen, and Intracranial Volume Indicates Shared Molecular Genetic Mechanisms. Schizophrenia Bulletin, 2018, 44, 854-864.	2.3	85
68	Identifying Common Genetic Variants in Blood Pressure Due to Polygenic Pleiotropy With Associated Phenotypes. Hypertension, 2014, 63, 819-826.	1.3	83
69	Shared Genetic Loci Between Body Mass Index and Major Psychiatric Disorders. JAMA Psychiatry, 2020, 77, 503.	6.0	82
70	Chronotype and cellular circadian rhythms predict the clinical response to lithium maintenance treatment in patients with bipolar disorder. Neuropsychopharmacology, 2019, 44, 620-628.	2.8	80
71	Brain scans from 21,297 individuals reveal the genetic architecture of hippocampal subfield volumes. Molecular Psychiatry, 2020, 25, 3053-3065.	4.1	80
72	Identification of genetic loci shared between schizophrenia and the Big Five personality traits. Scientific Reports, 2017, 7, 2222.	1.6	79

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73	CACNA1C Risk Variant and Amygdala Activity in Bipolar Disorder, Schizophrenia and Healthy Controls. PLoS ONE, 2013, 8, e56970.	1.1	76
74	Attention-deficit hyperactivity disorder shares copy number variant risk with schizophrenia and autism spectrum disorder. Translational Psychiatry, 2019, 9, 258.	2.4	75
75	Absence of enhanced systemic inflammatory response at 18 weeks of gestation in women with subsequent pre-eclampsia. BJOG: an International Journal of Obstetrics and Gynaecology, 2002, 109, 759-764.	1.1	73
76	Syndecan-4 Is Essential for Development of Concentric Myocardial Hypertrophy via Stretch-Induced Activation of the Calcineurin-NFAT Pathway. PLoS ONE, 2011, 6, e28302.	1.1	72
77	Plasma concentrations of Lp(a) lipoprotein and TGFâ€î²₁ are altered in preeclampsia. Clinical Genetics, 1997, 52, 371-376.	1.0	71
78	Genome-Wide Analysis of Attention Deficit Hyperactivity Disorder in Norway. PLoS ONE, 2015, 10, e0122501.	1.1	71
79	Comparison of Nonviral Transfection and Adeno-Associated Viral Transduction on Cardiomyocytes. Molecular Biotechnology, 2004, 28, 21-32.	1.3	64
80	The Endogenous Hallucinogen and Trace Amine N,N-Dimethyltryptamine (DMT) Displays Potent Protective Effects against Hypoxia via Sigma-1 Receptor Activation in Human Primary iPSC-Derived Cortical Neurons and Microglia-Like Immune Cells. Frontiers in Neuroscience, 2016, 10, 423.	1.4	64
81	Polygenic Risk for Schizophrenia Associated With Working Memory-related Prefrontal Brain Activation in Patients With Schizophrenia and Healthy Controls. Schizophrenia Bulletin, 2015, 41, 736-743.	2.3	62
82	Dyslipidemia in early second trimester is mainly a feature of women with early onset pre-eclampsia. British Journal of Obstetrics and Gynaecology, 2001, 108, 1081-1087.	0.9	61
83	Review: Epidemiology of Lp(a) lipoprotein: its role in atherosclerotic/thrombotic disease. Clinical Genetics, 1997, 52, 281-292.	1.0	61
84	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. Biological Psychiatry, 2022, 91, 102-117.	0.7	61
85	The Relationship Between Polygenic Risk Scores and Cognition in Schizophrenia. Schizophrenia Bulletin, 2020, 46, 336-344.	2.3	60
86	Serotonin Transporter Gene Polymorphism, Childhood Trauma, and Cognition in Patients With Psychotic Disorders. Schizophrenia Bulletin, 2012, 38, 15-22.	2.3	58
87	Altered circulating levels of adhesion molecules at 18 weeksâ€™ gestation among women with eventual preeclampsia: Indicators of disturbed placentation in absence of evidence of endothelial dysfunction?. American Journal of Obstetrics and Gynecology, 2000, 182, 321-325.	0.7	56
88	Identification of Gene Loci That Overlap Between Schizophrenia and Educational Attainment. Schizophrenia Bulletin, 2017, 43, sbw085.	2.3	56
89	Dyslipidemia in early second trimester is mainly a feature of women with early onset pre-eclampsia. BJOG: an International Journal of Obstetrics and Gynaecology, 2001, 108, 1081-1087.	1.1	55
90	Altered Brain Activation during Emotional Face Processing in Relation to Both Diagnosis and Polygenic Risk of Bipolar Disorder. PLoS ONE, 2015, 10, e0134202.	1.1	54

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91	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
92	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
93	GBA and APOE Îµ4 associate with sporadic dementia with Lewy bodies in European genome wide association study. Scientific Reports, 2019, 9, 7013.	1.6	53
94	Genome-wide Association Analysis of Parkinsonâ€™s Disease and Schizophrenia Reveals Shared Genetic Architecture and Identifies Novel Risk Loci. Biological Psychiatry, 2021, 89, 227-235.	0.7	53
95	Identifying Novel Gene Variants in Coronary Artery Disease and Shared Genes With Several Cardiovascular Risk Factors. Circulation Research, 2016, 118, 83-94.	2.0	52
96	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
97	Evidence for Genetic Overlap Between Schizophrenia and Age at First Birth in Women. JAMA Psychiatry, 2016, 73, 497.	6.0	51
98	Association of common genetic variants in GPCPD1 with scaling of visual cortical surface area in humans. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 3985-3990.	3.3	50
99	A genetic association study of CSMD1 and CSMD2 with cognitive function. Brain, Behavior, and Immunity, 2017, 61, 209-216.	2.0	49
100	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
101	Decreased IL-1Î²-induced CCL20 response in human iPSC-astrocytes in schizophrenia: Potential attenuating effects on recruitment of regulatory T cells. Brain, Behavior, and Immunity, 2020, 87, 634-644.	2.0	49
102	Catechol O-methyltransferase variants and cognitive performance in schizophrenia and bipolar disorder versus controls. Schizophrenia Research, 2010, 122, 31-37.	1.1	47
103	A Study of TNF Pathway Activation in Schizophrenia and Bipolar Disorder in Plasma and Brain Tissue. Schizophrenia Bulletin, 2017, 43, sbw183.	2.3	47
104	Meta-analysis of Alzheimerâ€™s disease on 9,751 samples from Norway and IGAP study identifies four risk loci. Scientific Reports, 2018, 8, 18088.	1.6	47
105	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
106	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
107	Shared common variants in prostate cancer and blood lipids. International Journal of Epidemiology, 2014, 43, 1205-1214.	0.9	45
108	Up-Regulation ofNOTCH4Gene Expression in Bipolar Disorder. American Journal of Psychiatry, 2012, 169, 1292-1300.	4.0	44

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109	Linkage-Disequilibrium-Based Binning Affects the Interpretation of GWASs. American Journal of Human Genetics, 2012, 90, 727-733.	2.6	44
110	Association between Genetic Variation in the Oxytocin Receptor Gene and Emotional Withdrawal, but not between Oxytocin Pathway Genes and Diagnosis in Psychotic Disorders. Frontiers in Human Neuroscience, 2015, 9, 9.	1.0	43
111	Genetic Association Between Schizophrenia and Cortical Brain Surface Area and Thickness. JAMA Psychiatry, 2021, 78, 1020.	6.0	43
112	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
113	Genetic Overlap Between Alzheimerâ€™s Disease and Bipolar Disorder Implicates the MARK2 and VAC14 Genes. Frontiers in Neuroscience, 2019, 13, 220.	1.4	42
114	Cardiometabolic risk factors associated with brain age and accelerated brain ageing. Human Brain Mapping, 2022, 43, 700-720.	1.9	42
115	No evidence for association between bipolar disorder risk gene variants and brain structural phenotypes. Journal of Affective Disorders, 2013, 151, 291-297.	2.0	41
116	The roadmap for estimation of cell-type-specific neuronal activity from non-invasive measurements. Philosophical Transactions of the Royal Society B: Biological Sciences, 2016, 371, 20150356.	1.8	41
117	Task modulations and clinical manifestations in the brain functional connectome in 1615 fMRI datasets. NeuroImage, 2017, 147, 243-252.	2.1	41
118	In Vivo Two-Photon Voltage Imaging with Sulfonated Rhodamine Dyes. ACS Central Science, 2018, 4, 1371-1378.	5.3	41
119	Telomere length is associated with childhood trauma in patients with severe mental disorders. Translational Psychiatry, 2019, 9, 97.	2.4	41
120	Increased levels of intercellular adhesion molecules and vascular cell adhesion molecules in pre-eclampsia. BJOG: an International Journal of Obstetrics and Gynaecology, 1997, 104, 466-470.	1.1	40
121	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
122	Inflammatory markers are altered in severe mental disorders independent of comorbid cardiometabolic disease risk factors. Psychological Medicine, 2019, 49, 1749-1757.	2.7	40
123	Abundant Genetic Overlap between Blood Lipids and Immune-Mediated Diseases Indicates Shared Molecular Genetic Mechanisms. PLoS ONE, 2015, 10, e0123057.	1.1	40
124	Association between a disrupted-in-schizophrenia 1 (DISC1) single nucleotide polymorphism and schizophrenia in a combined Scandinavian caseâ€“control sample. Schizophrenia Research, 2008, 106, 237-241.	1.1	39
125	Inflammatory evidence for the psychosis continuum model. Psychoneuroendocrinology, 2016, 67, 189-197.	1.3	39
126	Association analysis of ANK3 gene variants in nordic bipolar disorder and schizophrenia caseâ€“control samples. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 969-974.	1.1	37

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127	Attenuated Notch signaling in schizophrenia and bipolar disorder. Scientific Reports, 2018, 8, 5349.	1.6	37
128	Genome-wide study identifies PTPRO and WDR72 and FOXQ1-SUMO1P1 interaction associated with neurocognitive function. Journal of Psychiatric Research, 2012, 46, 271-278.	1.5	36
129	Examining the association between genetic liability for schizophrenia and psychotic symptoms in Alzheimerâ€™s disease. Translational Psychiatry, 2019, 9, 273.	2.4	36
130	Leveraging Genomic Annotations and Pleiotropic Enrichment for Improved Replication Rates in Schizophrenia GWAS. PLoS Genetics, 2016, 12, e1005803.	1.5	34
131	Electroporation by nucleofector is the best nonviral transfection technique in human endothelial and smooth muscle cells. Genetic Vaccines and Therapy, 2005, 3, 2.	1.5	33
132	<i>DTNBP1, NRG1, DAOA</i>, <i>DAO</i> and <i>GRM3</i> Polymorphisms and Schizophrenia: An Association Study. Neuropsychobiology, 2009, 59, 142-150.	0.9	33
133	Identification of shared genetic variants between schizophrenia and lung cancer. Scientific Reports, 2018, 8, 674.	1.6	33
134	Increased circulating IL-18 levels in severe mental disorders indicate systemic inflammasome activation. Brain, Behavior, and Immunity, 2022, 99, 299-306.	2.0	33
135	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
136	ANK3 gene expression in bipolar disorder and schizophrenia. British Journal of Psychiatry, 2014, 205, 244-245.	1.7	32
137	Identification of a novel polymorphism associated with reduced clozapine concentration in schizophrenia patientsâ€”a genome-wide association study adjusting for smoking habits. Translational Psychiatry, 2020, 10, 198.	2.4	32
138	Meta-analysis of heterogeneous data sources for genome-scale identification of risk genes in complex phenotypes. Genetic Epidemiology, 2011, 35, 318-332.	0.6	31
139	A Loss-of-Function Variant in a Minor Isoform of ANK3 Protects Against Bipolar Disorder and Schizophrenia. Biological Psychiatry, 2016, 80, 323-330.	0.7	31
140	Identification of genetic overlap and novel risk loci for attention-deficit/hyperactivity disorder and bipolar disorder. Molecular Psychiatry, 2021, 26, 4055-4065.	4.1	31
141	Large-scale genomics unveil polygenic architecture of human cortical surface area. Nature Communications, 2015, 6, 7549.	5.8	30
142	Novel Loci Associated With Attention-Deficit/Hyperactivity Disorder Are Revealed by Leveraging Polygenic Overlap With Educational Attainment. Journal of the American Academy of Child and Adolescent Psychiatry, 2018, 57, 86-95.	0.3	30
143	Alterations in Schizophrenia-Associated Genes Can Lead to Increased Power in Delta Oscillations. Cerebral Cortex, 2019, 29, 875-891.	1.6	30
144	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

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145	Genome-wide association identifies the first risk loci for psychosis in Alzheimer disease. Molecular Psychiatry, 2021, 26, 5797-5811.	4.1	30
146	DCLK1 Variants Are Associated across Schizophrenia and Attention Deficit/Hyperactivity Disorder. PLoS ONE, 2012, 7, e35424.	1.1	30
147	Lithium increases mitochondrial respiration in iPSC-derived neural precursor cells from lithium responders. Molecular Psychiatry, 2021, 26, 6789-6805.	4.1	29
148	Polygenic overlap and shared genetic loci between loneliness, severe mental disorders, and cardiovascular disease risk factors suggest shared molecular mechanisms. Translational Psychiatry, 2021, 11, 3.	2.4	29
149	The effect of red wine on plasma leptin levels and vasoactive factors from adipose tissue: A randomized crossover trial. Alcohol and Alcoholism, 2007, 42, 525-528.	0.9	28
150	Common Sequence Variants in the Major Histocompatibility Complex Region Associate with Cerebral Ventricular Size in Schizophrenia. Biological Psychiatry, 2011, 70, 696-698.	0.7	28
151	Association analysis between suicidal behaviour and candidate genes of bipolar disorder and schizophrenia. Journal of Affective Disorders, 2014, 163, 110-114.	2.0	28
152	Parentsâ€™ Attitudes toward Clinical Genetic Testing for Autism Spectrum Disorderâ€”Data from a Norwegian Sample. International Journal of Molecular Sciences, 2017, 18, 1078.	1.8	28
153	Stability of the Brain Functional Connectome Fingerprint in Individuals With Schizophrenia. JAMA Psychiatry, 2018, 75, 749.	6.0	28
154	Dissecting the shared genetic basis of migraine and mental disorders using novel statistical tools. Brain, 2022, 145, 142-153.	3.7	27
155	GWASinlps: non-local prior based iterative SNP selection tool for genome-wide association studies. Bioinformatics, 2019, 35, 1-11.	1.8	26
156	The genetic architecture of human brainstem structures and their involvement in common brain disorders. Nature Communications, 2020, 11, 4016.	5.8	26
157	Indicated association between polygenic risk score and treatment-resistance in a naturalistic sample of patients with schizophrenia spectrum disorders. Schizophrenia Research, 2020, 218, 55-62.	1.1	26
158	Loss-of-function variants in the schizophrenia risk gene SETD1A alter neuronal network activity in human neurons through the cAMP/PKA pathway. Cell Reports, 2022, 39, 110790.	2.9	26
159	Spontaneous atherosclerosis in the proximal aorta of LPA transgenic mice on a normal diet. Atherosclerosis, 2002, 163, 99-104.	0.4	25
160	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
161	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
162	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.	1.1	25

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163	Genetic overlap between multiple sclerosis and several cardiovascular disease risk factors. Multiple Sclerosis Journal, 2016, 22, 1783-1793.	1.4	25
164	Association of MCTP2 gene variants with schizophrenia in three independent samples of Scandinavian origin (SCOPE). Psychiatry Research, 2009, 168, 256-258.	1.7	24
165	Polygenic risk scores in bipolar disorder subgroups. Journal of Affective Disorders, 2015, 183, 310-314.	2.0	24
166	MicroRNAs enrichment in GWAS of complex human phenotypes. BMC Genomics, 2015, 16, 304.	1.2	24
167	Elevated expression of a minor isoform of ANK3 is a risk factor for bipolar disorder. Translational Psychiatry, 2018, 8, 210.	2.4	24
168	Exploring lithiumâ€™s transcriptional mechanisms of action in bipolar disorder: a multi-step study. Neuropsychopharmacology, 2020, 45, 947-955.	2.8	24
169	1q21.1 distal copy number variants are associated with cerebral and cognitive alterations in humans. Translational Psychiatry, 2021, 11, 182.	2.4	24
170	Characterizing the Genetic Overlap Between Psychiatric Disorders and Sleep-Related Phenotypes. Biological Psychiatry, 2021, 90, 621-631.	0.7	24
171	Associations Between Variants Near a Monoaminergic Pathways Gene (PHOX2B) and Amygdala Reactivity: A Genome-Wide Functional Imaging Study. Twin Research and Human Genetics, 2012, 15, 273-285.	0.3	23
172	Association between altered brain morphology and elevated peripheral endothelial markers â€" Implications for psychotic disorders. Schizophrenia Research, 2015, 161, 222-228.	1.1	23
173	Metabolic dysfunctions in the kynurenine pathway, noradrenergic and purine metabolism in schizophrenia and bipolar disorders. Psychological Medicine, 2020, 50, 595-606.	2.7	23
174	Genetic loci shared between major depression and intelligence with mixed directions of effect. Nature Human Behaviour, 2021, 5, 795-801.	6.2	23
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