

Srdjan Djurovic

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6291125/publications.pdf>

Version: 2024-02-01

280
papers

35,621
citations

11639

70
h-index

4988

167
g-index

332
all docs

332
docs citations

332
times ranked

35919
citing authors

#	ARTICLE	IF	CITATIONS
1	Lower circulating neuron-specific enolase concentrations in adults and adolescents with severe mental illness. <i>Psychological Medicine</i> , 2023, 53, 1479-1488.	2.7	6
2	Systemic Cell Adhesion Molecules in Severe Mental Illness: Potential Role of Intercellular CAM-1 in Linking Peripheral and Neuroinflammation. <i>Biological Psychiatry</i> , 2023, 93, 187-196.	0.7	18
3	Effects of copy number variations on brain structure and risk for psychiatric illness: Large-scale studies from the ENIGMA working groups on CNVs. <i>Human Brain Mapping</i> , 2022, 43, 300-328.	1.9	30
4	Association between complement component 4A expression, cognitive performance and brain imaging measures in UK Biobank. <i>Psychological Medicine</i> , 2022, 52, 3497-3507.	2.7	13
5	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. <i>Biological Psychiatry</i> , 2022, 91, 102-117.	0.7	61
6	Dissecting the shared genetic basis of migraine and mental disorders using novel statistical tools. <i>Brain</i> , 2022, 145, 142-153.	3.7	27
7	Plasma Levels of the Cytokines B Cell-Activating Factor (BAFF) and A Proliferation-Inducing Ligand (APRIL) in Schizophrenia, Bipolar, and Major Depressive Disorder: A Cross Sectional, Multisite Study. <i>Schizophrenia Bulletin</i> , 2022, 48, 37-46.	2.3	10
8	Genome-wide analysis reveals genetic overlap between alcohol use behaviours, schizophrenia and bipolar disorder and identifies novel shared risk loci. <i>Addiction</i> , 2022, 117, 600-610.	1.7	16
9	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. <i>Biological Psychiatry</i> , 2022, 91, 313-327.	0.7	114
10	Dose-dependent transcriptional effects of lithium and adverse effect burden in a psychiatric cohort. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2022, 112, 110408.	2.5	6
11	Cardiometabolic risk factors associated with brain age and accelerated brain ageing. <i>Human Brain Mapping</i> , 2022, 43, 700-720.	1.9	42
12	Increased circulating IL-18 levels in severe mental disorders indicate systemic inflammasome activation. <i>Brain, Behavior, and Immunity</i> , 2022, 99, 299-306.	2.0	33
13	Attitudes among parents of persons with autism spectrum disorder towards information about genetic risk and future health. <i>European Journal of Human Genetics</i> , 2022, 30, 1138-1146.	1.4	7
14	Mapping the expression of an ANK3 isoform associated with bipolar disorder in the human brain. <i>Translational Psychiatry</i> , 2022, 12, 45.	2.4	1
15	Immune marker levels in severe mental disorders: associations with polygenic risk scores of related mental phenotypes and psoriasis. <i>Translational Psychiatry</i> , 2022, 12, 38.	2.4	13
16	Limited association between infections, autoimmune disease and genetic risk and immune activation in severe mental disorders. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2022, 116, 110511.	2.5	4
17	Boosting Schizophrenia Genetics by Utilizing Genetic Overlap With Brain Morphology. <i>Biological Psychiatry</i> , 2022, 92, 291-298.	0.7	20
18	New insights into the genetic etiology of Alzheimer's disease and related dementias. <i>Nature Genetics</i> , 2022, 54, 412-436.	9.4	700

#	ARTICLE	IF	CITATIONS
19	Interleukin-18 signaling system links to agitation in severe mental disorders. <i>Psychoneuroendocrinology</i> , 2022, 140, 105721.	1.3	6
20	Loss-of-function variants in the schizophrenia risk gene SETD1A alter neuronal network activity in human neurons through the cAMP/PKA pathway. <i>Cell Reports</i> , 2022, 39, 110790.	2.9	26
21	Shared genetic loci between depression and cardiometabolic traits. <i>PLoS Genetics</i> , 2022, 18, e1010161.	1.5	18
22	P87. No Signs of Neurodegenerative Effects in 15q11.2 BP1-BP2 Copy Number Variant Carriers. <i>Biological Psychiatry</i> , 2022, 91, S122.	0.7	0
23	Using Polygenic Hazard Scores to Predict Age at Onset of Alzheimer's Disease in Nordic Populations. <i>Journal of Alzheimer's Disease</i> , 2022, 88, 1533-1544.	1.2	3
24	The shared genetic basis of mood instability and psychiatric disorders: A cross-trait genome-wide association analysis. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2022, 189, 207-218.	1.1	10
25	Composite immune marker scores associated with severe mental disorders and illness course. <i>Brain, Behavior, & Immunity - Health</i> , 2022, 24, 100483.	1.3	6
26	Identification of genetic overlap and novel risk loci for attention-deficit/hyperactivity disorder and bipolar disorder. <i>Molecular Psychiatry</i> , 2021, 26, 4055-4065.	4.1	31
27	Genetic control of variability in subcortical and intracranial volumes. <i>Molecular Psychiatry</i> , 2021, 26, 3876-3883.	4.1	6
28	Genome-wide Association Analysis of Parkinson's Disease and Schizophrenia Reveals Shared Genetic Architecture and Identifies Novel Risk Loci. <i>Biological Psychiatry</i> , 2021, 89, 227-235.	0.7	53
29	Using iPSC Models to Understand the Role of Estrogen in Neuron-Glia Interactions in Schizophrenia and Bipolar Disorder. <i>Cells</i> , 2021, 10, 209.	1.8	7
30	Genetic loci shared between major depression and intelligence with mixed directions of effect. <i>Nature Human Behaviour</i> , 2021, 5, 795-801.	6.2	23
31	1q21.1 distal copy number variants are associated with cerebral and cognitive alterations in humans. <i>Translational Psychiatry</i> , 2021, 11, 182.	2.4	24
32	Genetic variants associated with cardiometabolic abnormalities during treatment with selective serotonin reuptake inhibitors: a genome-wide association study. <i>Pharmacogenomics Journal</i> , 2021, 21, 574-585.	0.9	5
33	A Comparison of Ten Polygenic Score Methods for Psychiatric Disorders Applied Across Multiple Cohorts. <i>Biological Psychiatry</i> , 2021, 90, 611-620.	0.7	103
34	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. <i>Nature Genetics</i> , 2021, 53, 817-829.	9.4	629
35	Identifying nootropic drug targets via large-scale cognitive GWAS and transcriptomics. <i>Neuropsychopharmacology</i> , 2021, 46, 1788-1801.	2.8	12
36	Transcriptome analysis reveals disparate expression of inflammation-related miRNAs and their gene targets in iPSC-astrocytes from people with schizophrenia. <i>Brain, Behavior, and Immunity</i> , 2021, 94, 235-244.	2.0	17

#	ARTICLE	IF	CITATIONS
37	All-Optical Electrophysiology in hiPSC-Derived Neurons With Synthetic Voltage Sensors. <i>Frontiers in Cellular Neuroscience</i> , 2021, 15, 671549.	1.8	3
38	Genetic Overlap Between Schizophrenia and Brain Morphology. <i>Biological Psychiatry</i> , 2021, 89, S85-S86.	0.7	0
39	Common variants in Alzheimer's disease and risk stratification by polygenic risk scores. <i>Nature Communications</i> , 2021, 12, 3417.	5.8	140
40	Lithium increases mitochondrial respiration in iPSC-derived neural precursor cells from lithium responders. <i>Molecular Psychiatry</i> , 2021, 26, 6789-6805.	4.1	29
41	Shared genetic architecture between neuroticism, coronary artery disease and cardiovascular risk factors. <i>Translational Psychiatry</i> , 2021, 11, 368.	2.4	10
42	Genome-wide association identifies the first risk loci for psychosis in Alzheimer disease. <i>Molecular Psychiatry</i> , 2021, 26, 5797-5811.	4.1	30
43	Identification of pleiotropy at the gene level between psychiatric disorders and related traits. <i>Translational Psychiatry</i> , 2021, 11, 410.	2.4	7
44	Characterizing the Genetic Overlap Between Psychiatric Disorders and Sleep-Related Phenotypes. <i>Biological Psychiatry</i> , 2021, 90, 621-631.	0.7	24
45	Extensive bidirectional genetic overlap between bipolar disorder and cardiovascular disease phenotypes. <i>Translational Psychiatry</i> , 2021, 11, 407.	2.4	16
46	Characterisation of age and polarity at onset in bipolar disorder. <i>British Journal of Psychiatry</i> , 2021, 219, 659-669.	1.7	20
47	Derivation and Molecular Characterization of a Morphological Subpopulation of Human iPSC Astrocytes Reveal a Potential Role in Schizophrenia and Clozapine Response. <i>Schizophrenia Bulletin</i> , 2021, , .	2.3	14
48	A genome-wide association study with 1,126,563 individuals identifies new risk loci for Alzheimer's disease. <i>Nature Genetics</i> , 2021, 53, 1276-1282.	9.4	430
49	Characterising the shared genetic determinants of bipolar disorder, schizophrenia and risk-taking. <i>Translational Psychiatry</i> , 2021, 11, 466.	2.4	15
50	Genetic Association Between Schizophrenia and Cortical Brain Surface Area and Thickness. <i>JAMA Psychiatry</i> , 2021, 78, 1020.	6.0	43
51	Telomeres are shorter and associated with number of suicide attempts in affective disorders. <i>Journal of Affective Disorders</i> , 2021, 295, 1032-1039.	2.0	13
52	Polygenic overlap and shared genetic loci between loneliness, severe mental disorders, and cardiovascular disease risk factors suggest shared molecular mechanisms. <i>Translational Psychiatry</i> , 2021, 11, 3.	2.4	29
53	A human iPSC-astroglia neurodevelopmental model reveals divergent transcriptomic patterns in schizophrenia. <i>Translational Psychiatry</i> , 2021, 11, 554.	2.4	19
54	The Relationship Between Polygenic Risk Scores and Cognition in Schizophrenia. <i>Schizophrenia Bulletin</i> , 2020, 46, 336-344.	2.3	60

#	ARTICLE	IF	CITATIONS
55	Brain scans from 21,297 individuals reveal the genetic architecture of hippocampal subfield volumes. <i>Molecular Psychiatry</i> , 2020, 25, 3053-3065.	4.1	80
56	Dose response of the 16p11.2 distal copy number variant on intracranial volume and basal ganglia. <i>Molecular Psychiatry</i> , 2020, 25, 584-602.	4.1	49
57	Genome-wide analysis reveals extensive genetic overlap between schizophrenia, bipolar disorder, and intelligence. <i>Molecular Psychiatry</i> , 2020, 25, 844-853.	4.1	156
58	Metabolic dysfunctions in the kynurenine pathway, noradrenergic and purine metabolism in schizophrenia and bipolar disorders. <i>Psychological Medicine</i> , 2020, 50, 595-606.	2.7	23
59	Discovery of shared genomic loci using the conditional false discovery rate approach. <i>Human Genetics</i> , 2020, 139, 85-94.	1.8	109
60	Exploring lithium's transcriptional mechanisms of action in bipolar disorder: a multi-step study. <i>Neuropsychopharmacology</i> , 2020, 45, 947-955.	2.8	24
61	The Genetics of the Mood Disorder Spectrum: Genome-wide Association Analyses of More Than 185,000 Cases and 439,000 Controls. <i>Biological Psychiatry</i> , 2020, 88, 169-184.	0.7	137
62	Association of Copy Number Variation of the 15q11.2 BP1-BP2 Region With Cortical and Subcortical Morphology and Cognition. <i>JAMA Psychiatry</i> , 2020, 77, 420.	6.0	54
63	Shared Genetic Loci Between Body Mass Index and Major Psychiatric Disorders. <i>JAMA Psychiatry</i> , 2020, 77, 503.	6.0	82
64	Identification of Genetic Loci Shared Between Attention-Deficit/Hyperactivity Disorder, Intelligence, and Educational Attainment. <i>Biological Psychiatry</i> , 2020, 87, 1052-1062.	0.7	13
65	The genetic architecture of human brainstem structures and their involvement in common brain disorders. <i>Nature Communications</i> , 2020, 11, 4016.	5.8	26
66	Cannabis Use Is Associated With Increased Levels of Soluble gp130 in Schizophrenia but Not in Bipolar Disorder. <i>Frontiers in Psychiatry</i> , 2020, 11, 642.	1.3	5
67	Atherogenic Lipid Ratios Related to Myeloperoxidase and C-Reactive Protein Levels in Psychotic Disorders. <i>Frontiers in Psychiatry</i> , 2020, 11, 672.	1.3	15
68	Computationally efficient familywise error rate control in genome-wide association studies using score tests for generalized linear models. <i>Scandinavian Journal of Statistics</i> , 2020, 47, 1090-1113.	0.9	2
69	Polygenic scores for schizophrenia and general cognitive ability: associations with six cognitive domains, premorbid intelligence, and cognitive composite score in individuals with a psychotic disorder and in healthy controls. <i>Translational Psychiatry</i> , 2020, 10, 416.	2.4	16
70	Epigenetic Differences in Patients With Psychosis Using Cannabis and After Discontinuation. <i>Biological Psychiatry</i> , 2020, 87, S283.	0.7	0
71	Runaway multi-allelic copy number variation at the β -defensin locus in African and Asian populations. <i>Scientific Reports</i> , 2020, 10, 9101.	1.6	3
72	Phenotype-specific differences in polygenicity and effect size distribution across functional annotation categories revealed by AI-MiXeR. <i>Bioinformatics</i> , 2020, 36, 4749-4756.	1.8	6

#	ARTICLE	IF	CITATIONS
73	Copy number variants (CNVs): a powerful tool for iPSC-based modelling of ASD. <i>Molecular Autism</i> , 2020, 11, 42.	2.6	14
74	Identification of a novel polymorphism associated with reduced clozapine concentration in schizophrenia patients—a genome-wide association study adjusting for smoking habits. <i>Translational Psychiatry</i> , 2020, 10, 198.	2.4	32
75	The genetic architecture of the human cerebral cortex. <i>Science</i> , 2020, 367, .	6.0	450
76	Decreased IL-1 β -induced CCL20 response in human iPSC-astrocytes in schizophrenia: Potential attenuating effects on recruitment of regulatory T cells. <i>Brain, Behavior, and Immunity</i> , 2020, 87, 634-644.	2.0	49
77	Indicated association between polygenic risk score and treatment-resistance in a naturalistic sample of patients with schizophrenia spectrum disorders. <i>Schizophrenia Research</i> , 2020, 218, 55-62.	1.1	26
78	GWASinlps: non-local prior based iterative SNP selection tool for genome-wide association studies. <i>Bioinformatics</i> , 2019, 35, 1-11.	1.8	26
79	Pleiotropic Meta-Analysis of Cognition, Education, and Schizophrenia Differentiates Roles of Early Neurodevelopmental and Adult Synaptic Pathways. <i>American Journal of Human Genetics</i> , 2019, 105, 334-350.	2.6	86
80	Biophysical Psychiatry—How Computational Neuroscience Can Help to Understand the Complex Mechanisms of Mental Disorders. <i>Frontiers in Psychiatry</i> , 2019, 10, 534.	1.3	19
81	Examining the association between genetic liability for schizophrenia and psychotic symptoms in Alzheimer's disease. <i>Translational Psychiatry</i> , 2019, 9, 273.	2.4	36
82	Attention-deficit hyperactivity disorder shares copy number variant risk with schizophrenia and autism spectrum disorder. <i>Translational Psychiatry</i> , 2019, 9, 258.	2.4	75
83	O1.6. TELOMERE LENGTH IS ASSOCIATED WITH CHILDHOOD TRAUMA IN PATIENTS WITH SEVERE MENTAL DISORDERS. <i>Schizophrenia Bulletin</i> , 2019, 45, S160-S161.	2.3	0
84	Common brain disorders are associated with heritable patterns of apparent aging of the brain. <i>Nature Neuroscience</i> , 2019, 22, 1617-1623.	7.1	358
85	Inflammatory markers are altered in severe mental disorders independent of comorbid cardiometabolic disease risk factors. <i>Psychological Medicine</i> , 2019, 49, 1749-1757.	2.7	40
86	Bivariate causal mixture model quantifies polygenic overlap between complex traits beyond genetic correlation. <i>Nature Communications</i> , 2019, 10, 2417.	5.8	190
87	GWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. <i>American Journal of Psychiatry</i> , 2019, 176, 651-660.	4.0	186
88	GBA and APOE ϵ 4 associate with sporadic dementia with Lewy bodies in European genome wide association study. <i>Scientific Reports</i> , 2019, 9, 7013.	1.6	53
89	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019, 51, 793-803.	9.4	1,191
90	Telomere length is associated with childhood trauma in patients with severe mental disorders. <i>Translational Psychiatry</i> , 2019, 9, 97.	2.4	41

#	ARTICLE	IF	CITATIONS
91	Genetic Overlap Between Alzheimer's Disease and Bipolar Disorder Implicates the MARK2 and VAC14 Genes. <i>Frontiers in Neuroscience</i> , 2019, 13, 220.	1.4	42
92	Brain Heterogeneity in Schizophrenia and Its Association With Polygenic Risk. <i>JAMA Psychiatry</i> , 2019, 76, 739.	6.0	195
93	S18. THE RELATIONSHIP BETWEEN PHYSICAL ACTIVITY, CLINICAL AND COGNITIVE CHARACTERISTICS AND BDNF MRNA LEVELS IN PATIENTS WITH SEVERE MENTAL DISORDERS. <i>Schizophrenia Bulletin</i> , 2019, 45, S312-S312.	2.3	0
94	Identification of common genetic risk variants for autism spectrum disorder. <i>Nature Genetics</i> , 2019, 51, 431-444.	9.4	1,538
95	188. ENIGMA-CNV: Unraveling the Effects of Rare Copy Number Variants on Brain Structure. <i>Biological Psychiatry</i> , 2019, 85, S78.	0.7	3
96	Genetic architecture of subcortical brain structures in 38,851 individuals. <i>Nature Genetics</i> , 2019, 51, 1624-1636.	9.4	192
97	The relationship between physical activity, clinical and cognitive characteristics and BDNF mRNA levels in patients with severe mental disorders. <i>World Journal of Biological Psychiatry</i> , 2019, 20, 567-576.	1.3	15
98	Genome-wide meta-analysis identifies new loci and functional pathways influencing Alzheimer's disease risk. <i>Nature Genetics</i> , 2019, 51, 404-413.	9.4	1,625
99	Chronotype and cellular circadian rhythms predict the clinical response to lithium maintenance treatment in patients with bipolar disorder. <i>Neuropsychopharmacology</i> , 2019, 44, 620-628.	2.8	80
100	Alterations in Schizophrenia-Associated Genes Can Lead to Increased Power in Delta Oscillations. <i>Cerebral Cortex</i> , 2019, 29, 875-891.	1.6	30
101	Exploring the Wnt signaling pathway in schizophrenia and bipolar disorder. <i>Translational Psychiatry</i> , 2018, 8, 55.	2.4	94
102	Common schizophrenia alleles are enriched in mutation-intolerant genes and in regions under strong background selection. <i>Nature Genetics</i> , 2018, 50, 381-389.	9.4	1,332
103	Genetic Overlap Between Schizophrenia and Volumes of Hippocampus, Putamen, and Intracranial Volume Indicates Shared Molecular Genetic Mechanisms. <i>Schizophrenia Bulletin</i> , 2018, 44, 854-864.	2.3	85
104	Novel Loci Associated With Attention-Deficit/Hyperactivity Disorder Are Revealed by Leveraging Polygenic Overlap With Educational Attainment. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2018, 57, 86-95.	0.3	30
105	Identification of shared genetic variants between schizophrenia and lung cancer. <i>Scientific Reports</i> , 2018, 8, 674.	1.6	33
106	Genetic variation in 117 myelination-related genes in schizophrenia: Replication of association to lipid biosynthesis genes. <i>Scientific Reports</i> , 2018, 8, 6915.	1.6	10
107	A molecule-based genetic association approach implicates a range of voltage-gated calcium channels associated with schizophrenia. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018, 177, 454-467.	1.1	12
108	Attenuated Notch signaling in schizophrenia and bipolar disorder. <i>Scientific Reports</i> , 2018, 8, 5349.	1.6	37

#	ARTICLE	IF	CITATIONS
109	Genetic factors influencing prostate cancer risk in Norwegian men. <i>Prostate</i> , 2018, 78, 186-192.	1.2	11
110	Meta-analysis of Alzheimer's disease on 9,751 samples from Norway and IGAP study identifies four risk loci. <i>Scientific Reports</i> , 2018, 8, 18088.	1.6	47
111	<i>In Vivo</i> Two-Photon Voltage Imaging with Sulfonated Rhodamine Dyes. <i>ACS Central Science</i> , 2018, 4, 1371-1378.	5.3	41
112	Elevated expression of a minor isoform of ANK3 is a risk factor for bipolar disorder. <i>Translational Psychiatry</i> , 2018, 8, 210.	2.4	24
113	F50. Genetic Architecture of Hippocampal Subfield Volumes: Shared and Specific Influences. <i>Biological Psychiatry</i> , 2018, 83, S257.	0.7	0
114	Cross-tissue eQTL enrichment of associations in schizophrenia. <i>PLoS ONE</i> , 2018, 13, e0202812.	1.1	6
115	Enrichment of genetic markers of recent human evolution in educational and cognitive traits. <i>Scientific Reports</i> , 2018, 8, 12585.	1.6	9
116	Study of 300,486 individuals identifies 148 independent genetic loci influencing general cognitive function. <i>Nature Communications</i> , 2018, 9, 2098.	5.8	484
117	Stability of the Brain Functional Connectome Fingerprint in Individuals With Schizophrenia. <i>JAMA Psychiatry</i> , 2018, 75, 749.	6.0	28
118	Deep 2-photon imaging and artifact-free optogenetics through transparent graphene microelectrode arrays. <i>Nature Communications</i> , 2018, 9, 2035.	5.8	143
119	Estimation of Genetic Correlation via Linkage Disequilibrium Score Regression and Genomic Restricted Maximum Likelihood. <i>American Journal of Human Genetics</i> , 2018, 102, 1185-1194.	2.6	119
120	Genome-wide association meta-analysis in 269,867 individuals identifies new genetic and functional links to intelligence. <i>Nature Genetics</i> , 2018, 50, 912-919.	9.4	893
121	Multi-Trait Analysis of GWAS and Biological Insights Into Cognition: A Response to Hill (2018). <i>Twin Research and Human Genetics</i> , 2018, 21, 394-397.	0.3	3
122	Expression of TCN1 in Blood is Negatively Associated with Verbal Declarative Memory Performance. <i>Scientific Reports</i> , 2018, 8, 12654.	1.6	14
123	Vitamin D levels, brain volume, and genetic architecture in patients with psychosis. <i>PLoS ONE</i> , 2018, 13, e0200250.	1.1	11
124	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. <i>Cell</i> , 2018, 173, 1705-1715.e16.	13.5	623
125	Identification of Gene Loci That Overlap Between Schizophrenia and Educational Attainment. <i>Schizophrenia Bulletin</i> , 2017, 43, sbw085.	2.3	56
126	Novel genetic loci associated with hippocampal volume. <i>Nature Communications</i> , 2017, 8, 13624.	5.8	250

#	ARTICLE	IF	CITATIONS
127	A Study of TNF Pathway Activation in Schizophrenia and Bipolar Disorder in Plasma and Brain Tissue. <i>Schizophrenia Bulletin</i> , 2017, 43, sbw183.	2.3	47
128	A genetic association study of CSMD1 and CSMD2 with cognitive function. <i>Brain, Behavior, and Immunity</i> , 2017, 61, 209-216.	2.0	49
129	Genome-wide Pleiotropy Between Parkinson Disease and Autoimmune Diseases. <i>JAMA Neurology</i> , 2017, 74, 780.	4.5	245
130	Identification of genetic loci shared between schizophrenia and the Big Five personality traits. <i>Scientific Reports</i> , 2017, 7, 2222.	1.6	79
131	Task modulations and clinical manifestations in the brain functional connectome in 1615 fMRI datasets. <i>NeuroImage</i> , 2017, 147, 243-252.	2.1	41
132	Identification of Genetic Loci Jointly Influencing Schizophrenia Risk and the Cognitive Traits of Verbal-Numerical Reasoning, Reaction Time, and General Cognitive Function. <i>JAMA Psychiatry</i> , 2017, 74, 1065.	6.0	123
133	Fourteen sequence variants that associate with multiple sclerosis discovered by meta-analysis informed by genetic correlations. <i>Npj Genomic Medicine</i> , 2017, 2, 24.	1.7	16
134	Distinct multivariate brain morphological patterns and their added predictive value with cognitive and polygenic risk scores in mental disorders. <i>NeuroImage: Clinical</i> , 2017, 15, 719-731.	1.4	89
135	Large-Scale Cognitive GWAS Meta-Analysis Reveals Tissue-Specific Neural Expression and Potential Nootropic Drug Targets. <i>Cell Reports</i> , 2017, 21, 2597-2613.	2.9	103
136	Leveraging genome characteristics to improve gene discovery for putamen subcortical brain structure. <i>Scientific Reports</i> , 2017, 7, 15736.	1.6	15
137	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. <i>Nature Genetics</i> , 2017, 49, 27-35.	9.4	838
138	Analysis of the joint effect of SNPs to identify independent loci and allelic heterogeneity in schizophrenia GWAS data. <i>Translational Psychiatry</i> , 2017, 7, 1289.	2.4	4
139	Parents' Attitudes toward Clinical Genetic Testing for Autism Spectrum Disorder—Data from a Norwegian Sample. <i>International Journal of Molecular Sciences</i> , 2017, 18, 1078.	1.8	28
140	Probing the Association between Early Evolutionary Markers and Schizophrenia. <i>PLoS ONE</i> , 2017, 12, e0169227.	1.1	17
141	Combinations of genetic variants associated with bipolar disorder. <i>PLoS ONE</i> , 2017, 12, e0189739.	1.1	6
142	Contribution of oxytocin receptor polymorphisms to amygdala activation in schizophrenia spectrum disorders. <i>BJPsych Open</i> , 2016, 2, 353-358.	0.3	11
143	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. <i>Frontiers in Neuroscience</i> , 2016, 10, 423.	1.4	64
144	Cell type specificity of neurovascular coupling in cerebral cortex. <i>ELife</i> , 2016, 5, .	2.8	176

#	ARTICLE	IF	CITATIONS
145	Leveraging Genomic Annotations and Pleiotropic Enrichment for Improved Replication Rates in Schizophrenia GWAS. <i>PLoS Genetics</i> , 2016, 12, e1005803.	1.5	34
146	Parents' attitudes toward genetic research in autism spectrum disorder. <i>Psychiatric Genetics</i> , 2016, 26, 74-80.	0.6	13
147	Novel genetic loci underlying human intracranial volume identified through genome-wide association. <i>Nature Neuroscience</i> , 2016, 19, 1569-1582.	7.1	213
148	<i>VRK2</i> gene expression in schizophrenia, bipolar disorder and healthy controls. <i>British Journal of Psychiatry</i> , 2016, 209, 114-120.	1.7	17
149	Pleiotropic Analysis of Lung Cancer and Blood Triglycerides. <i>Journal of the National Cancer Institute</i> , 2016, 108, djw167.	3.0	17
150	The roadmap for estimation of cell-type-specific neuronal activity from non-invasive measurements. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2016, 371, 20150356.	1.8	41
151	Evidence for Genetic Overlap Between Schizophrenia and Age at First Birth in Women. <i>JAMA Psychiatry</i> , 2016, 73, 497.	6.0	51
152	Functional Effects of Schizophrenia-Linked Genetic Variants on Intrinsic Single-Neuron Excitability: A Modeling Study. <i>Biological Psychiatry: Cognitive Neuroscience and Neuroimaging</i> , 2016, 1, 49-59.	1.1	21
153	Genetic overlap between multiple sclerosis and several cardiovascular disease risk factors. <i>Multiple Sclerosis Journal</i> , 2016, 22, 1783-1793.	1.4	25
154	Inflammatory evidence for the psychosis continuum model. <i>Psychoneuroendocrinology</i> , 2016, 67, 189-197.	1.3	39
155	Identification of rare high-risk copy number variants affecting the dopamine transporter gene in mental disorders. <i>Nordic Journal of Psychiatry</i> , 2016, 70, 276-279.	0.7	2
156	A Loss-of-Function Variant in a Minor Isoform of ANK3 Protects Against Bipolar Disorder and Schizophrenia. <i>Biological Psychiatry</i> , 2016, 80, 323-330.	0.7	31
157	Genetic Markers of Human Evolution Are Enriched in Schizophrenia. <i>Biological Psychiatry</i> , 2016, 80, 284-292.	0.7	92
158	Identifying Novel Gene Variants in Coronary Artery Disease and Shared Genes With Several Cardiovascular Risk Factors. <i>Circulation Research</i> , 2016, 118, 83-94.	2.0	52
159	Conservation of Distinct Genetically-Mediated Human Cortical Pattern. <i>PLoS Genetics</i> , 2016, 12, e1006143.	1.5	15
160	Independent evidence for an association between general cognitive ability and a genetic locus for educational attainment. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2015, 168, 363-373.	1.1	25
161	Altered Brain Activation during Emotional Face Processing in Relation to Both Diagnosis and Polygenic Risk of Bipolar Disorder. <i>PLoS ONE</i> , 2015, 10, e0134202.	1.1	54
162	Combinations of Genetic Data Present in Bipolar Patients, but Absent in Control Persons. <i>PLoS ONE</i> , 2015, 10, e0143432.	1.1	4

#	ARTICLE	IF	CITATIONS
163	Genetic Sharing with Cardiovascular Disease Risk Factors and Diabetes Reveals Novel Bone Mineral Density Loci. PLoS ONE, 2015, 10, e0144531.	1.1	14
164	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
165	Polygenic risk scores in bipolar disorder subgroups. Journal of Affective Disorders, 2015, 183, 310-314.	2.0	24
166	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
167	Common genetic variants influence human subcortical brain structures. Nature, 2015, 520, 224-229.	13.7	772
168	Large-scale genomics unveil polygenic architecture of human cortical surface area. Nature Communications, 2015, 6, 7549.	5.8	30
169	Loss-of-function variants in ABCA7 confer risk of Alzheimer's disease. Nature Genetics, 2015, 47, 445-447.	9.4	283
170	Genome-wide association study identifies common variants associated with pharmacokinetics of psychotropic drugs. Journal of Psychopharmacology, 2015, 29, 884-891.	2.0	12
171	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
172	Polygenic Overlap Between C-Reactive Protein, Plasma Lipids, and Alzheimer Disease. Circulation, 2015, 131, 2061-2069.	1.6	145
173	Modeling Linkage Disequilibrium Increases Accuracy of Polygenic Risk Scores. American Journal of Human Genetics, 2015, 97, 576-592.	2.6	1,098
174	MicroRNAs enrichment in GWAS of complex human phenotypes. BMC Genomics, 2015, 16, 304.	1.2	24
175	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
176	Investigation of the genetic interaction between <i>BDNF</i> and <i>DRD3</i> genes in suicidal behaviour in psychiatric disorders. World Journal of Biological Psychiatry, 2015, 16, 171-179.	1.3	14
177	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
178	Association between altered brain morphology and elevated peripheral endothelial markers " Implications for psychotic disorders. Schizophrenia Research, 2015, 161, 222-228.	1.1	23
179	Genome-Wide Analysis of Attention Deficit Hyperactivity Disorder in Norway. PLoS ONE, 2015, 10, e0122501.	1.1	71
180	Abundant Genetic Overlap between Blood Lipids and Immune-Mediated Diseases Indicates Shared Molecular Genetic Mechanisms. PLoS ONE, 2015, 10, e0123057.	1.1	40

#	ARTICLE	IF	CITATIONS
181	Microarray Analysis of Copy Number Variants on the Human Y Chromosome Reveals Novel and Frequent Duplications Overrepresented in Specific Haplogroups. PLoS ONE, 2015, 10, e0137223.	1.1	17
182	Shared common variants in prostate cancer and blood lipids. International Journal of Epidemiology, 2014, 43, 1205-1214.	0.9	45
183	An Attempt to Identify Single Nucleotide Polymorphisms Contributing to Possible Relationships between Personality Traits and Oxytocin-Related Genes. Neuropsychobiology, 2014, 69, 25-30.	0.9	15
184	Identifying Common Genetic Variants in Blood Pressure Due to Polygenic Pleiotropy With Associated Phenotypes. Hypertension, 2014, 63, 819-826.	1.3	83
185	The ENIGMA Consortium: large-scale collaborative analyses of neuroimaging and genetic data. Brain Imaging and Behavior, 2014, 8, 153-182.	1.1	696
186	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
187	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
188	Polygenic dissection of diagnosis and clinical dimensions of bipolar disorder and schizophrenia. Molecular Psychiatry, 2014, 19, 1017-1024.	4.1	333
189	<i>ANKK1</i> gene expression in bipolar disorder and schizophrenia. British Journal of Psychiatry, 2014, 205, 244-245.	1.7	32
190	Association analysis between suicidal behaviour and candidate genes of bipolar disorder and schizophrenia. Journal of Affective Disorders, 2014, 163, 110-114.	2.0	28
191	Pathway analysis of genetic markers associated with a functional MRI faces paradigm implicates polymorphisms in calcium responsive pathways. NeuroImage, 2013, 70, 143-149.	2.1	13
192	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. Nature Genetics, 2013, 45, 984-994.	9.4	2,067
193	Variant of <i>TREM2</i> Associated with the Risk of Alzheimer's Disease. New England Journal of Medicine, 2013, 368, 107-116.	13.9	2,085
194	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
195	ZNF804A and cortical thickness in schizophrenia and bipolar disorder. Psychiatry Research - Neuroimaging, 2013, 212, 154-157.	0.9	17
196	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
197	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
198	A Comprehensive Family-Based Replication Study of Schizophrenia Genes. JAMA Psychiatry, 2013, 70, 573.	6.0	138

#	ARTICLE	IF	CITATIONS
199	Improved Detection of Common Variants Associated with Schizophrenia and Bipolar Disorder Using Pleiotropy-Informed Conditional False Discovery Rate. <i>PLoS Genetics</i> , 2013, 9, e1003455.	1.5	298
200	A Genetic Deconstruction of Neurocognitive Traits in Schizophrenia and Bipolar Disorder. <i>PLoS ONE</i> , 2013, 8, e81052.	1.1	20
201	CACNA1C Risk Variant and Amygdala Activity in Bipolar Disorder, Schizophrenia and Healthy Controls. <i>PLoS ONE</i> , 2013, 8, e56970.	1.1	76
202	Genetic Clustering on the Hippocampal Surface for Genome-Wide Association Studies. <i>Lecture Notes in Computer Science</i> , 2013, 16, 690-697.	1.0	7
203	Associations Between Variants Near a Monoaminergic Pathways Gene (PHOX2B) and Amygdala Reactivity: A Genome-Wide Functional Imaging Study. <i>Twin Research and Human Genetics</i> , 2012, 15, 273-285.	0.3	23
204	Up-Regulation of <i>NOTCH4</i> Gene Expression in Bipolar Disorder. <i>American Journal of Psychiatry</i> , 2012, 169, 1292-1300.	4.0	44
205	Association of common genetic variants in GPCPD1 with scaling of visual cortical surface area in humans. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 3985-3990.	3.3	50
206	Serotonin Transporter Gene Polymorphism, Childhood Trauma, and Cognition in Patients With Psychotic Disorders. <i>Schizophrenia Bulletin</i> , 2012, 38, 15-22.	2.3	58
207	Lack of association between the regulator of G-protein signaling 4 (RGS4) rs951436 polymorphism and schizophrenia. <i>Psychiatric Genetics</i> , 2012, 22, 263-264.	0.6	9
208	Genome-wide association study identifies genetic loci associated with body mass index and high density lipoprotein-cholesterol levels during psychopharmacological treatment – a cross-sectional naturalistic study. <i>Psychiatry Research</i> , 2012, 197, 327-336.	1.7	9
209	Identification of common variants associated with human hippocampal and intracranial volumes. <i>Nature Genetics</i> , 2012, 44, 552-561.	9.4	594
210	Association between a genetic variant in the serotonin transporter gene (SLC6A4) and suicidal behavior in patients with schizophrenia. <i>Behavioral and Brain Functions</i> , 2012, 8, 24.	1.4	15
211	Replication Study and Meta-Analysis in European Samples Supports Association of the 3p21.1 Locus with Bipolar Disorder. <i>Biological Psychiatry</i> , 2012, 72, 645-650.	0.7	15
212	Gene-Based Analysis of Regionally Enriched Cortical Genes in GWAS Data Sets of Cognitive Traits and Psychiatric Disorders. <i>PLoS ONE</i> , 2012, 7, e31687.	1.1	40
213	Connection between Genetic and Clinical Data in Bipolar Disorder. <i>PLoS ONE</i> , 2012, 7, e44623.	1.1	8
214	Effect of <i>DISC1</i> SNPs on brain structure in healthy controls and patients with a history of psychosis. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2012, 159B, 722-730.	1.1	14
215	Linkage-Disequilibrium-Based Binning Affects the Interpretation of GWASs. <i>American Journal of Human Genetics</i> , 2012, 90, 727-733.	2.6	44
216	Genome-wide study identifies PTPRO and WDR72 and FOXQ1-SUMO1P1 interaction associated with neurocognitive function. <i>Journal of Psychiatric Research</i> , 2012, 46, 271-278.	1.5	36

#	ARTICLE	IF	CITATIONS
217	DCLK1 Variants Are Associated across Schizophrenia and Attention Deficit/Hyperactivity Disorder. PLoS ONE, 2012, 7, e35424.	1.1	30
218	Common variants at VRK2 and TCF4 conferring risk of schizophrenia. Human Molecular Genetics, 2011, 20, 4076-4081.	1.4	193
219	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
220	The Complement Control-Related Genes CSMD1 and CSMD2 Associate to Schizophrenia. Biological Psychiatry, 2011, 70, 35-42.	0.7	149
221	At-Risk Variant in TCF7L2 for Type II Diabetes Increases Risk of Schizophrenia. Biological Psychiatry, 2011, 70, 59-63.	0.7	114
222	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
223	Kynurenine 3-monooxygenase (KMO) polymorphisms in schizophrenia: An association study. Schizophrenia Research, 2011, 127, 270-272.	1.1	19
224	Intron 12 in NTRK3 is associated with bipolar disorder. Psychiatry Research, 2011, 185, 358-362.	1.7	21
225	Combinations of SNPs Related to Signal Transduction in Bipolar Disorder. PLoS ONE, 2011, 6, e23812.	1.1	20
226	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
227	Dual association of a TRKA polymorphism with schizophrenia. Psychiatric Genetics, 2011, 21, 125-131.	0.6	8
228	Lack of association between two dopamine D2 receptor gene polymorphisms and schizophrenia. Psychiatric Genetics, 2011, 21, 214-215.	0.6	3
229	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
230	Association analysis of ANK3 gene variants in nordic bipolar disorder and schizophrenia case-control samples. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 969-974.	1.1	37
231	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
232	Maternally Derived Microduplications at 15q11-q13: Implication of Imprinted Genes in Psychotic Illness. American Journal of Psychiatry, 2011, 168, 408-417.	4.0	95
233	Copy number variations in affective disorders and meta-analysis. Psychiatric Genetics, 2011, 21, 319-322.	0.6	3
234	Association of Genetic Variants on 15q12 With Cortical Thickness and Cognition in Schizophrenia. Archives of General Psychiatry, 2011, 68, 781.	13.8	22

#	ARTICLE	IF	CITATIONS
235	Association study of <i>PDE4B</i> 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
236	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
237	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
238	The tryptophan hydroxylase 1 (<i>TPH1</i>) 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
239	Association between methylenetetrahydrofolate reductase (<i>MTHFR</i>) C677T polymorphism and age of onset in schizophrenia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 610-618.	1.1	32
240	Association analysis of <i>PALB2</i> and <i>BRCA2</i> 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
241	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
242	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
243	Catechol O-methyltransferase variants and cognitive performance in schizophrenia and bipolar disorder versus controls. Schizophrenia Research, 2010, 122, 31-37.	1.1	47
244	SRD5A2 is associated with increased cortisol metabolism in schizophrenia spectrum disorders. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2010, 34, 1500-1506.	2.5	19
245	Osteoprotegerin levels in patients with severe mental disorders. Journal of Psychiatry and Neuroscience, 2010, 35, 304-310.	1.4	21
246	Dysbindin and <i>D</i>-Amino-Acid-Oxidase Gene Polymorphisms Associated with Positive and Negative Symptoms in Schizophrenia. Neuropsychobiology, 2009, 60, 31-36.	0.9	20
247	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
248	Disruption of the neurexin 1 gene is associated with schizophrenia. Human Molecular Genetics, 2009, 18, 988-996.	1.4	424
249	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
250	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
251	Common variants conferring risk of schizophrenia. Nature, 2009, 460, 744-747.	13.7	1,572
252	<i>DTNBP1</i>, <i>NRG1</i>, <i>DAO</i>, <i>DAO</i> and <i>GRM3</i> Polymorphisms and Schizophrenia: An Association Study. Neuropsychobiology, 2009, 59, 142-150.	0.9	33

#	ARTICLE	IF	CITATIONS
253	Association of MCTP2 gene variants with schizophrenia in three independent samples of Scandinavian origin (SCOPE). <i>Psychiatry Research</i> , 2009, 168, 256-258.	1.7	24
254	A possible association between schizophrenia and GRIK3 polymorphisms in a multicenter sample of Scandinavian origin (SCOPE). <i>Schizophrenia Research</i> , 2009, 107, 242-248.	1.1	17
255	No association between DGKH and bipolar disorder in a Scandinavian case-control sample. <i>Psychiatric Genetics</i> , 2009, 19, 269-272.	0.6	18
256	Tyrosine hydroxylase Val81Met polymorphism: lack of association with schizophrenia. <i>Psychiatric Genetics</i> , 2009, 19, 273-274.	0.6	3
257	A novel myogenic cell line with phenotypic properties of muscle progenitors. <i>Journal of Molecular Medicine</i> , 2008, 86, 105-115.	1.7	4
258	Association analysis of schizophrenia on 18 genes involved in neuronal migration: <i>MDGA1</i> as a new susceptibility gene. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008, 147B, 1089-1100.	1.1	101
259	Large recurrent microdeletions associated with schizophrenia. <i>Nature</i> , 2008, 455, 232-236.	13.7	1,619
260	Association between a disrupted-in-schizophrenia 1 (DISC1) single nucleotide polymorphism and schizophrenia in a combined Scandinavian case-control sample. <i>Schizophrenia Research</i> , 2008, 106, 237-241.	1.1	39
261	The effect of red wine on plasma leptin levels and vasoactive factors from adipose tissue: A randomized crossover trial. <i>Alcohol and Alcoholism</i> , 2007, 42, 525-528.	0.9	28
262	Brain Expressed microRNAs Implicated in Schizophrenia Etiology. <i>PLoS ONE</i> , 2007, 2, e873.	1.1	235
263	Electroporation by nucleofector is the best nonviral transfection technique in human endothelial and smooth muscle cells. <i>Genetic Vaccines and Therapy</i> , 2005, 3, 2.	1.5	33
264	Activity of peroxisomal enzymes, and levels of polyamines in LPA-transgenic mice on two different diets. <i>Lipids in Health and Disease</i> , 2005, 4, 23.	1.2	1
265	Variations in Transfection Efficiency of VEGF ₁₆₅ and VEGF ₁₂₁ -cDNA: Its Effects on Proliferation and Migration of Human Endothelial Cells. <i>Molecular Biotechnology</i> , 2004, 26, 7-16.	1.3	4
266	Comparison of Nonviral Transfection and Adeno-Associated Viral Transduction on Cardiomyocytes. <i>Molecular Biotechnology</i> , 2004, 28, 21-32.	1.3	64
267	Human apoB contributes to increased serum total apo(a) level in LPA transgenic mice. <i>Lipids in Health and Disease</i> , 2004, 3, 8.	1.2	4
268	Missense mutation W86R in exon 3 of the lipoprotein lipase gene in a boy with chylomicronemia. <i>Clinica Chimica Acta</i> , 2004, 343, 179-184.	0.5	14
269	The decrease in urinary albumin excretion after irbesartan treatment in normotensive diabetes type 1 patients with incipient nephropathy is related to blood pressure, and not to changes in soluble cytokines and growth factors. <i>American Journal of Hypertension</i> , 2004, 17, S119.	1.0	0
270	Circulating Transforming Growth Factor- β 21, Lipoprotein(a) and Cellular Adhesion Molecules in Angiographically Assessed Coronary Artery Disease. <i>Clinical Chemistry and Laboratory Medicine</i> , 2003, 41, 893-8.	1.4	12

#	ARTICLE	IF	CITATIONS
271	Spontaneous atherosclerosis in the proximal aorta of LPA transgenic mice on a normal diet. <i>Atherosclerosis</i> , 2002, 163, 99-104.	0.4	25
272	Absence of enhanced systemic inflammatory response at 18 weeks of gestation in women with subsequent pre-eclampsia. <i>BJOG: an International Journal of Obstetrics and Gynaecology</i> , 2002, 109, 759-764.	1.1	73
273	Absence of enhanced systemic inflammatory response at 18 weeks of gestation in women with subsequent pre-eclampsia. <i>BJOG: an International Journal of Obstetrics and Gynaecology</i> , 2002, 109, 759-764.	1.1	1
274	Dyslipidemia in early second trimester is mainly a feature of women with early onset pre-eclampsia. <i>British Journal of Obstetrics and Gynaecology</i> , 2001, 108, 1081-1087.	0.9	61
275	Plasma N-terminal Pro-atrial Natriuretic Peptide Predicts Death after Premature Myocardial Infarction, but not as well as Radionuclide Ejection Fraction. A Ten-year Follow-up Study. <i>Scandinavian Cardiovascular Journal</i> , 2001, 35, 373-378.	0.4	3
276	Dyslipidemia in early second trimester is mainly a feature of women with early onset pre-eclampsia. <i>BJOG: an International Journal of Obstetrics and Gynaecology</i> , 2001, 108, 1081-1087.	1.1	55
277	Altered circulating levels of adhesion molecules at 18 weeks gestation among women with eventual preeclampsia: Indicators of disturbed placentation in absence of evidence of endothelial dysfunction?. <i>American Journal of Obstetrics and Gynecology</i> , 2000, 182, 321-325.	0.7	56
278	Increased levels of intercellular adhesion molecules and vascular cell adhesion molecules in pre-eclampsia. <i>BJOG: an International Journal of Obstetrics and Gynaecology</i> , 1997, 104, 466-470.	1.1	40
279	Review: Epidemiology of Lp(a) lipoprotein: its role in atherosclerotic/thrombotic disease. <i>Clinical Genetics</i> , 1997, 52, 281-292.	1.0	61
280	Plasma concentrations of Lp(a) lipoprotein and TGF β ₁ are altered in preeclampsia. <i>Clinical Genetics</i> , 1997, 52, 371-376.	1.0	71