

Stephanie H Witt

List of Publications by Year in descending order

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Version: 2024-02-01

153
papers

14,419
citations

76031

42
h-index

32181

105
g-index

173
all docs

173
docs citations

173
times ranked

18895
citing authors

#	ARTICLE	IF	CITATIONS
1	Polygenic risk for schizophrenia and schizotypal traits in non-clinical subjects. <i>Psychological Medicine</i> , 2022, 52, 1069-1079.	2.7	10
2	Ventral Striatum-Hippocampus Coupling During Reward Processing as a Stratification Biomarker for Psychotic Disorders. <i>Biological Psychiatry</i> , 2022, 91, 216-225.	0.7	10
3	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
4	Genome-wide interaction study with major depression identifies novel variants associated with cognitive function. <i>Molecular Psychiatry</i> , 2022, 27, 1111-1119.	4.1	24
5	Epigenome-wide association study of alcohol use disorder in five brain regions. <i>Neuropsychopharmacology</i> , 2022, 47, 832-839.	2.8	16
6	Real-time individual benefit from social interactions before and during the lockdown: the crucial role of personality, neurobiology and genes. <i>Translational Psychiatry</i> , 2022, 12, 28.	2.4	4
7	Urbanicity, behavior problems and HPA axis regulation in preschoolers. <i>Psychoneuroendocrinology</i> , 2022, 137, 105660.	1.3	1
8	Meta-analysis of epigenome-wide associations between DNA methylation at birth and childhood cognitive skills. <i>Molecular Psychiatry</i> , 2022, 27, 2126-2135.	4.1	13
9	Using polygenic scores and clinical data for bipolar disorder patient stratification and lithium response prediction: machine learning approach. <i>British Journal of Psychiatry</i> , 2022, 220, 219-228.	1.7	11
10	Genetic variants associated with longitudinal changes in brain structure across the lifespan. <i>Nature Neuroscience</i> , 2022, 25, 421-432.	7.1	75
11	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. <i>Nature</i> , 2022, 604, 502-508.	13.7	929
12	Epigenetic Signatures of Smoking in Five Brain Regions. <i>Journal of Personalized Medicine</i> , 2022, 12, 566.	1.1	4
13	Adverse childhood experiences and late-life diurnal HPA axis activity: Associations of different childhood adversity types and interaction with timing in a sample of older East Prussian World War II refugees. <i>Psychoneuroendocrinology</i> , 2022, 139, 105717.	1.3	4
14	Borderline personality disorder and the big five: molecular genetic analyses indicate shared genetic architecture with neuroticism and openness. <i>Translational Psychiatry</i> , 2022, 12, 153.	2.4	7
15	Multi-omics signatures of alcohol use disorder in the dorsal and ventral striatum. <i>Translational Psychiatry</i> , 2022, 12, 190.	2.4	11
16	A novel longitudinal clustering approach to psychopathology across diagnostic entities in the hospital-based PsyCourse study. <i>Schizophrenia Research</i> , 2022, 244, 29-38.	1.1	2
17	Epigenetic signatures in antidepressant treatment response: a methylome-wide association study in the EMC trial. <i>Translational Psychiatry</i> , 2022, 12, .	2.4	4
18	Association of polygenic score for major depression with response to lithium in patients with bipolar disorder. <i>Molecular Psychiatry</i> , 2021, 26, 2457-2470.	4.1	44

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19	Generative network models of altered structural brain connectivity in schizophrenia. <i>NeuroImage</i> , 2021, 225, 117510.	2.1	24
20	Molecular characterization of ulcerative colitis-associated colorectal carcinomas. <i>Modern Pathology</i> , 2021, 34, 1153-1166.	2.9	7
21	Childhood maltreatment and cognitive functioning: the role of depression, parental education, and polygenic predisposition. <i>Neuropsychopharmacology</i> , 2021, 46, 891-899.	2.8	17
22	Bipolar multiplex families have an increased burden of common risk variants for psychiatric disorders. <i>Molecular Psychiatry</i> , 2021, 26, 1286-1298.	4.1	33
23	Interaction of developmental factors and ordinary stressful life events on brain structure in adults. <i>NeuroImage: Clinical</i> , 2021, 30, 102683.	1.4	5
24	DNA methylation signatures of aggression and closely related constructs: A meta-analysis of epigenome-wide studies across the lifespan. <i>Molecular Psychiatry</i> , 2021, 26, 2148-2162.	4.1	21
25	Clinical and genetic differences between bipolar disorder type 1 and 2 in multiplex families. <i>Translational Psychiatry</i> , 2021, 11, 31.	2.4	22
26	“The Heidelberg Five” personality dimensions: Genome-wide associations, polygenic risk for neuroticism, and psychopathology 20 years after assessment. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2021, 186, 77-89.	1.1	6
27	Rhythm of Fetoplacental 11 β -Hydroxysteroid Dehydrogenase Type 2 “Fetal Protection From Morning Maternal Glucocorticoids. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2021, 106, 1630-1636.	1.8	7
28	Effects of polygenic risk for major mental disorders and cross-disorder on cortical complexity. <i>Psychological Medicine</i> , 2021, , 1-12.	2.7	7
29	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
30	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
31	Apolipoprotein E homozygous ϵ 4 allele status: Effects on cortical structure and white matter integrity in a young to mid-age sample. <i>European Neuropsychopharmacology</i> , 2021, 46, 93-104.	0.3	2
32	Childhood trauma and insulin-like growth factors in amniotic fluid. <i>Psychoneuroendocrinology</i> , 2021, 127, 105180.	1.3	2
33	Methylome-wide change associated with response to electroconvulsive therapy in depressed patients. <i>Translational Psychiatry</i> , 2021, 11, 347.	2.4	12
34	Identification of transdiagnostic psychiatric disorder subtypes using unsupervised learning. <i>Neuropsychopharmacology</i> , 2021, 46, 1895-1905.	2.8	24
35	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. <i>JAMA Psychiatry</i> , 2021, 78, 1258.	6.0	88
36	HLA-DRB1 and HLA-DQB1 genetic diversity modulates response to lithium in bipolar affective disorders. <i>Scientific Reports</i> , 2021, 11, 17823.	1.6	10

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37	Microbiome profiles are associated with cognitive functioning in 45-month-old children. <i>Brain, Behavior, and Immunity</i> , 2021, 98, 151-160.	2.0	18
38	Interplay between the genetics of personality traits, severe psychiatric disorders and COVID-19 host genetics in the susceptibility to SARS-CoV-2 infection. <i>BJPsych Open</i> , 2021, 7, e188.	0.3	1
39	Combining schizophrenia and depression polygenic risk scores improves the genetic prediction of lithium response in bipolar disorder patients. <i>Translational Psychiatry</i> , 2021, 11, 606.	2.4	25
40	Polygenic risk scores across the extended psychosis spectrum. <i>Translational Psychiatry</i> , 2021, 11, 600.	2.4	11
41	Cortical surface area alterations shaped by genetic load for neuroticism. <i>Molecular Psychiatry</i> , 2020, 25, 3422-3431.	4.1	20
42	Sex-dependent effects of <i>Cacna1c</i> haploinsufficiency on juvenile social play behavior and pro-social 50kHz ultrasonic communication in rats. <i>Genes, Brain and Behavior</i> , 2020, 19, e12552.	1.1	29
43	Genomics of borderline personality disorder. , 2020, , 227-237.		3
44	Cortical Surfaces Mediate the Relationship Between Polygenic Scores for Intelligence and General Intelligence. <i>Cerebral Cortex</i> , 2020, 30, 2708-2719.	1.6	24
45	Addiction Research Consortium: Losing and regaining control over drug intake (ReCoDe) – From trajectories to mechanisms and interventions. <i>Addiction Biology</i> , 2020, 25, e12866.	1.4	135
46	Genetic correlations and genome-wide associations of cortical structure in general population samples of 22,824 adults. <i>Nature Communications</i> , 2020, 11, 4796.	5.8	61
47	A large-scale genome-wide association study meta-analysis of cannabis use disorder. <i>Lancet Psychiatry</i> , 2020, 7, 1032-1045.	3.7	200
48	Prospects of Genetics and Epigenetics of Alcohol Use Disorder. <i>Current Addiction Reports</i> , 2020, 7, 446-452.	1.6	3
49	Association of Locomotor Activity During Sleep Deprivation Treatment With Response. <i>Frontiers in Psychiatry</i> , 2020, 11, 688.	1.3	2
50	Newly established gastrointestinal cancer cell lines retain the genomic and immunophenotypic landscape of their parental cancers. <i>Scientific Reports</i> , 2020, 10, 17895.	1.6	5
51	Replication of a hippocampus specific effect of the tescalcin regulating variant rs7294919 on gray matter structure. <i>European Neuropsychopharmacology</i> , 2020, 36, 10-17.	0.3	2
52	Leptin predicts cortical and subcortical gray matter volume recovery in alcohol dependent patients: A longitudinal structural magnetic resonance imaging study. <i>Hormones and Behavior</i> , 2020, 124, 104749.	1.0	7
53	The genetic architecture of the human cerebral cortex. <i>Science</i> , 2020, 367, .	6.0	450
54	Advanced paternal age as a risk factor for neurodevelopmental disorders: a translational study. <i>Molecular Autism</i> , 2020, 11, 54.	2.6	20

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55	Acute alcohol withdrawal and recovery in men lead to profound changes in DNA methylation profiles: a longitudinal clinical study. <i>Addiction</i> , 2020, 115, 2034-2044.	1.7	21
56	Rhythm and blues: Influence of CLOCK T3111C on peripheral electrophysiological indicators of negative affective processing. <i>Physiology and Behavior</i> , 2020, 219, 112831.	1.0	2
57	Whole-exome sequencing of 81 individuals from 27 multiply affected bipolar disorder families. <i>Translational Psychiatry</i> , 2020, 10, 57.	2.4	23
58	Hyperfunctioning of the right posterior superior temporal sulcus in response to neutral facial expressions presents an endophenotype of schizophrenia. <i>Neuropsychopharmacology</i> , 2020, 45, 1346-1352.	2.8	12
59	Genome-wide gene-environment analyses of major depressive disorder and reported lifetime traumatic experiences in UK Biobank. <i>Molecular Psychiatry</i> , 2020, 25, 1430-1446.	4.1	116
60	An Investigation of Psychosis Subgroups With Prognostic Validation and Exploration of Genetic Underpinnings. <i>JAMA Psychiatry</i> , 2020, 77, 523.	6.0	39
61	The role of environmental stress and DNA methylation in the longitudinal course of bipolar disorder. <i>International Journal of Bipolar Disorders</i> , 2020, 8, 9.	0.8	13
62	A longitudinal approach to biological psychiatric research: The PsyCourse study. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019, 180, 89-102.	1.1	47
63	Investigating polygenic burden in age at disease onset in bipolar disorder: Findings from an international multicentric study. <i>Bipolar Disorders</i> , 2019, 21, 68-75.	1.1	20
64	Long-term environmental impact on object recognition, spatial memory, and reversal learning capabilities in <i>Cacna1c</i> haploinsufficient rats. <i>Human Molecular Genetics</i> , 2019, 28, 4113-4131.	1.4	9
65	Effects of Mindfulness-Based Stress Prevention on Serotonin Transporter Gene Methylation. <i>Psychotherapy and Psychosomatics</i> , 2019, 88, 317-319.	4.0	17
66	The genetic relationship between educational attainment and cognitive performance in major psychiatric disorders. <i>Translational Psychiatry</i> , 2019, 9, 210.	2.4	24
67	MAOA ϵ VNTR genotype affects structural and functional connectivity in distributed brain networks. <i>Human Brain Mapping</i> , 2019, 40, 5202-5212.	1.9	14
68	Reproducible grey matter patterns index a multivariate, global alteration of brain structure in schizophrenia and bipolar disorder. <i>Translational Psychiatry</i> , 2019, 9, 12.	2.4	35
69	Apolipoprotein E Homozygous ϵ 4 Allele Status: A Deteriorating Effect on Visuospatial Working Memory and Global Brain Structure. <i>Frontiers in Neurology</i> , 2019, 10, 552.	1.1	10
70	Associations of schizophrenia risk genes ZNF804A and CACNA1C with schizotypy and modulation of attention in healthy subjects. <i>Schizophrenia Research</i> , 2019, 208, 67-75.	1.1	20
71	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019, 51, 793-803.	9.4	1,191
72	Attitudes toward the right to autonomous decision-making in psychiatric genetic testing: Controversial and context-dependent. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019, 180, 555-565.	1.1	6

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73	Cover Image, Volume 180B, Number 2, March 2019. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, i.	1.1	0
74	Prenatal maternal stress is associated with lower cortisol and cortisone levels in the first morning urine of 45-month-old children. Psychoneuroendocrinology, 2019, 103, 219-224.	1.3	19
75	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. Cell, 2019, 179, 1469-1482.e11.	13.5	935
76	Longitudinal transcriptome-wide gene expression analysis of sleep deprivation treatment shows involvement of circadian genes and immune pathways. Translational Psychiatry, 2019, 9, 343.	2.4	21
77	The influence of religious activity and polygenic schizophrenia risk on religious delusions in schizophrenia. Schizophrenia Research, 2019, 210, 255-261.	1.1	9
78	Stress reactivity in preschool-aged children: Evaluation of a social stress paradigm and investigation of the impact of prenatal maternal stress. Psychoneuroendocrinology, 2019, 101, 223-231.	1.3	11
79	Effects of BDNF Val66Met genotype and schizophrenia familial risk on a neural functional network for cognitive control in humans. Neuropsychopharmacology, 2019, 44, 590-597.	2.8	19
80	Evidence for increased genetic risk load for major depression in patients assigned to electroconvulsive therapy. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, 35-45.	1.1	18
81	Efficient region-based test strategy uncovers genetic risk factors for functional outcome in bipolar disorder. European Neuropsychopharmacology, 2019, 29, 156-170.	0.3	7
82	Effects of leptin and ghrelin on neural cue-reactivity in alcohol addiction: Two streams merge to one river?. Psychoneuroendocrinology, 2019, 100, 1-9.	1.3	28
83	Impact of FAAH genetic variation on fronto-amygdala function during emotional processing. European Archives of Psychiatry and Clinical Neuroscience, 2019, 269, 209-221.	1.8	26
84	Neurobiology of the major psychoses: a translational perspective on brain structure and function—the FOR2107 consortium. European Archives of Psychiatry and Clinical Neuroscience, 2019, 269, 949-962.	1.8	103
85	Association of Polygenic Score for Schizophrenia and HLA Antigen and Inflammation Genes With Response to Lithium in Bipolar Affective Disorder. JAMA Psychiatry, 2018, 75, 65-74.	6.0	102
86	Polygenic risk for schizophrenia affects working memory and its neural correlates in healthy subjects. Schizophrenia Research, 2018, 197, 315-320.	1.1	11
87	Outgroup emotion processing in the vACC is modulated by childhood trauma and CACNA1C risk variant. Social Cognitive and Affective Neuroscience, 2018, 13, 341-348.	1.5	13
88	Fast sleep spindle density is associated with rs4680 (Val108/158Met) genotype of catechol-O-methyltransferase (COMT). Sleep, 2018, 41, .	0.6	13
89	Neurogenetic Approaches to Stress and Fear in Humans as Pathophysiological Mechanisms for Posttraumatic Stress Disorder. Biological Psychiatry, 2018, 83, 810-820.	0.7	21
90	Impact on birth weight of maternal smoking throughout pregnancy mediated by DNA methylation. BMC Genomics, 2018, 19, 290.	1.2	41

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91	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. <i>Nature Genetics</i> , 2018, 50, 668-681.	9.4	2,224
92	The 5-HTTLPR Polymorphism Affects Network-Based Functional Connectivity in the Visual-Limbic System in Healthy Adults. <i>Neuropsychopharmacology</i> , 2018, 43, 406-414.	2.8	22
93	The influence of MIR137 on white matter fractional anisotropy and cortical surface area in individuals with familial risk for psychosis. <i>Schizophrenia Research</i> , 2018, 195, 190-196.	1.1	6
94	Gene set enrichment analysis and expression pattern exploration implicate an involvement of neurodevelopmental processes in bipolar disorder. <i>Journal of Affective Disorders</i> , 2018, 228, 20-25.	2.0	14
95	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. <i>Nature Neuroscience</i> , 2018, 21, 1656-1669.	7.1	490
96	Exome sequencing in large, multiplex bipolar disorder families from Cuba. <i>PLoS ONE</i> , 2018, 13, e0205895.	1.1	13
97	Maternal hypothalamus-pituitary-adrenal (HPA) system activity and stress during pregnancy: Effects on gestational age and infant's anthropometric measures at birth. <i>Psychoneuroendocrinology</i> , 2018, 94, 152-161.	1.3	66
98	Analysis of the Influence of microRNAs in Lithium Response in Bipolar Disorder. <i>Frontiers in Psychiatry</i> , 2018, 9, 207.	1.3	28
99	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , 2018, 360, .	6.0	1,085
100	Response to the letter by Esteves et al.. <i>Neuropsychopharmacology</i> , 2018, 43, 2164-2164.	2.8	0
101	Risk-seeking for losses is associated with 5-HTTLPR, but not with transient changes in 5-HT levels. <i>Psychopharmacology</i> , 2018, 235, 2151-2165.	1.5	13
102	<i>Cacna1c</i> haploinsufficiency leads to pro-social 50-kHz ultrasonic communication deficits in rats. <i>DMM Disease Models and Mechanisms</i> , 2018, 11, .	1.2	51
103	Response to therapeutic sleep deprivation: a naturalistic study of clinical and genetic factors and post-treatment depressive symptom trajectory. <i>Neuropsychopharmacology</i> , 2018, 43, 2572-2577.	2.8	17
104	Shared genetic etiology between alcohol dependence and major depressive disorder. <i>Psychiatric Genetics</i> , 2018, 28, 66-70.	0.6	19
105	Neurocognitive Endophenotypes of Schizophrenia and Bipolar Disorder and Possible Associations with FKBP Variant rs3800373. <i>Medicinski Arhiv = Medical Archives = Archives De Médecine</i> , 2018, 72, 352.	0.4	4
106	Functional neuroimaging effects of recently discovered genetic risk loci for schizophrenia and polygenic risk profile in five RDoC subdomains. <i>Translational Psychiatry</i> , 2017, 7, e997-e997.	2.4	31
107	Telomere Length in Newborns is Related to Maternal Stress During Pregnancy. <i>Neuropsychopharmacology</i> , 2017, 42, 2407-2413.	2.8	83
108	Associations of the Intellectual Disability Gene MYT1L with Helix-Loop-Helix Gene Expression, Hippocampus Volume and Hippocampus Activation During Memory Retrieval. <i>Neuropsychopharmacology</i> , 2017, 42, 2516-2526.	2.8	20

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109	Polygenic risk for depression and the neural correlates of working memory in healthy subjects. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2017, 79, 67-76.	2.5	41
110	Genome-wide association study of borderline personality disorder reveals genetic overlap with bipolar disorder, major depression and schizophrenia. <i>Translational Psychiatry</i> , 2017, 7, e1155-e1155.	2.4	150
111	Genetic effects influencing risk for major depressive disorder in China and Europe. <i>Translational Psychiatry</i> , 2017, 7, e1074-e1074.	2.4	64
112	Associations between SNPs and immune-related circulating proteins in schizophrenia. <i>Scientific Reports</i> , 2017, 7, 12586.	1.6	21
113	Drinking water to reduce alcohol craving? A randomized controlled study on the impact of ghrelin in mediating the effects of forced water intake in alcohol addiction. <i>Psychoneuroendocrinology</i> , 2017, 85, 56-62.	1.3	15
114	Hair Cortisol in Twins: Heritability and Genetic Overlap with Psychological Variables and Stress-System Genes. <i>Scientific Reports</i> , 2017, 7, 15351.	1.6	50
115	Epigenome-wide DNA methylation analysis in siblings and monozygotic twins discordant for sporadic Parkinson's disease revealed different epigenetic patterns in peripheral blood mononuclear cells. <i>Neurogenetics</i> , 2017, 18, 7-22.	0.7	47
116	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
117	Genetic Contribution to Alcohol Dependence: Investigation of a Heterogeneous German Sample of Individuals with Alcohol Dependence, Chronic Alcoholic Pancreatitis, and Alcohol-Related Cirrhosis. <i>Genes</i> , 2017, 8, 183.	1.0	11
118	Identification of shared risk loci and pathways for bipolar disorder and schizophrenia. <i>PLoS ONE</i> , 2017, 12, e0171595.	1.1	77
119	Analysis of Rare Variants in the Alcohol Dependence Candidate Gene GATA 4. <i>Alcoholism: Clinical and Experimental Research</i> , 2016, 40, 1627-1632.	1.4	1
120	Hair Cortisol and Its Association With Psychological Risk Factors for Psychiatric Disorders: A Pilot Study in Adolescent Twins. <i>Twin Research and Human Genetics</i> , 2016, 19, 438-446.	0.3	31
121	Perceived stress and hair cortisol: Differences in bipolar disorder and schizophrenia. <i>Psychoneuroendocrinology</i> , 2016, 69, 26-34.	1.3	48
122	Altered Functional Subnetwork During Emotional Face Processing. <i>JAMA Psychiatry</i> , 2016, 73, 598.	6.0	59
123	Mutational hierarchies in myelodysplastic syndromes dynamically adapt and evolve upon therapy response and failure. <i>Blood</i> , 2016, 128, 1246-1259.	0.6	111
124	Genome-wide association study of 40,000 individuals identifies two novel loci associated with bipolar disorder. <i>Human Molecular Genetics</i> , 2016, 25, 3383-3394.	1.4	182
125	Interacting effect of MAOA genotype and maternal prenatal smoking on aggressive behavior in young adulthood. <i>Journal of Neural Transmission</i> , 2016, 123, 885-894.	1.4	10
126	Genetic variants associated with response to lithium treatment in bipolar disorder: a genome-wide association study. <i>Lancet</i> , 2016, 387, 1085-1093.	6.3	306

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127	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
128	Theory of mind network activity is altered in subjects with familial liability for schizophrenia. <i>Social Cognitive and Affective Neuroscience</i> , 2016, 11, 299-307.	1.5	18
129	No Reliable Association between Runs of Homozygosity and Schizophrenia in a Well-Powered Replication Study. <i>PLoS Genetics</i> , 2016, 12, e1006343.	1.5	24
130	New-born females show higher stress- and genotype-independent methylation of SLC6A4 than males. <i>Borderline Personality Disorder and Emotion Dysregulation</i> , 2015, 2, 8.	1.1	38
131	Identification of increased genetic risk scores for schizophrenia in treatment-resistant patients. <i>Molecular Psychiatry</i> , 2015, 20, 150-151.	4.1	98
132	XRCC5 as a Risk Gene for Alcohol Dependence: Evidence from a Genome-Wide Gene-Set-Based Analysis and Follow-up Studies in <i>Drosophila</i> and Humans. <i>Neuropsychopharmacology</i> , 2015, 40, 361-371.	2.8	12
133	Neural Mechanism of a Sex-Specific Risk Variant for Posttraumatic Stress Disorder in the Type I Receptor of the Pituitary Adenylate Cyclase Activating Polypeptide. <i>Biological Psychiatry</i> , 2015, 78, 840-847.	0.7	47
134	MORC1 exhibits cross-species differential methylation in association with early life stress as well as genome-wide association with MDD. <i>Translational Psychiatry</i> , 2014, 4, e429-e429.	2.4	82
135	Striatal Response to Reward Anticipation. <i>JAMA Psychiatry</i> , 2014, 71, 531.	6.0	96
136	Identification of gene ontologies linked to prefrontalâ€“hippocampal functional coupling in the human brain. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 9657-9662.	3.3	9
137	Further Evidence for the Impact of a Genome-Wide-Supported Psychosis Risk Variant in ZNF804A on the Theory of Mind Network. <i>Neuropsychopharmacology</i> , 2014, 39, 1196-1205.	2.8	42
138	Analysis of genome-wide significant bipolar disorder genes in borderline personality disorder. <i>Psychiatric Genetics</i> , 2014, 24, 262-265.	0.6	26
139	Epigenetic alteration of the dopamine transporter gene in alcoholâ€“dependent patients is associated with age. <i>Addiction Biology</i> , 2014, 19, 305-311.	1.4	44
140	A genome-wide supported variant in CACNA1C influences hippocampal activation during episodic memory encoding and retrieval. <i>European Archives of Psychiatry and Clinical Neuroscience</i> , 2014, 264, 103-110.	1.8	33
141	Genome-wide association study reveals two new risk loci for bipolar disorder. <i>Nature Communications</i> , 2014, 5, 3339.	5.8	294
142	Genetic variation in CACNA1C affects neural processing in major depression. <i>Journal of Psychiatric Research</i> , 2014, 53, 38-46.	1.5	42
143	Replication of brain function effects of a genome-wide supported psychiatric risk variant in the CACNA1C gene and new multi-locus effects. <i>NeuroImage</i> , 2014, 94, 147-154.	2.1	32
144	The impact of a CACNA1C gene polymorphism on learning and hippocampal formation in healthy individuals: A diffusion tensor imaging study. <i>NeuroImage</i> , 2014, 89, 256-261.	2.1	32

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145	Hippocampal and Frontolimbic Function as Intermediate Phenotype for Psychosis: Evidence from Healthy Relatives and a Common Risk Variant in CACNA1C. <i>Biological Psychiatry</i> , 2014, 76, 466-475.	0.7	57
146	Moderating role of FKBP5 genotype in the impact of childhood adversity on cortisol stress response during adulthood. <i>European Neuropsychopharmacology</i> , 2014, 24, 837-845.	0.3	78
147	Longer telomere length in patients with schizophrenia. <i>Schizophrenia Research</i> , 2013, 149, 116-120.	1.1	57
148	Comparison of gene expression profiles in the blood, hippocampus and prefrontal cortex of rats. In <i>Silico Pharmacology</i> , 2013, 1, 15.	1.8	39
149	The Effect of Neurogranin on Neural Correlates of Episodic Memory Encoding and Retrieval. <i>Schizophrenia Bulletin</i> , 2013, 39, 141-150.	2.3	33
150	Genome-wide investigation of rare structural variants identifies <i>VIPR2</i> as a new candidate gene for schizophrenia. <i>Expert Review of Neurotherapeutics</i> , 2011, 11, 937-941.	1.4	8
151	Effects of a genome-wide supported psychosis risk variant on neural activation during a theory-of-mind task. <i>Molecular Psychiatry</i> , 2011, 16, 462-470.	4.1	133
152	Neural Mechanisms of a Genome-Wide Supported Psychosis Variant. <i>Science</i> , 2009, 324, 605-605.	6.0	375
153	Relationship Between Parental Big Five And Children's Ability To Delay Gratification. <i>Journal of Child and Family Studies</i> , 0, , 1.	0.7	1