Thomas M Werge

List of Publications by Year
in descending order

279 papers	34,597 citations	13068 68 h-index	5364 164 g-index
339 all docs	339 docs citations	339 times ranked	34389 citing authors

#	Article	IF	Citations
1	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. Biological Psychiatry, 2022, 91, 102-117.	0.7	61
2	Identifying the Common Genetic Basis of Antidepressant Response. Biological Psychiatry Global Open Science, 2022, 2, 115-126.	1.0	31
3	Common Genetic Variation and Age of Onset of Anorexia Nervosa. Biological Psychiatry Global Open Science, 2022, 2, 368-378.	1.0	10
4	Enhancing Discovery of Genetic Variants for Posttraumatic Stress Disorder Through Integration of Quantitative Phenotypes and Trauma Exposure Information. Biological Psychiatry, 2022, 91, 626-636.	0.7	21
5	Cohort Profile: COVIDMENT: COVID-19 cohorts on mental health across six nations. International Journal of Epidemiology, 2022, 51, e108-e122.	0.9	16
6	Evaluating the interrelations between the autism polygenic score and psychiatric family history in risk for autism. Autism Research, 2022, 15, 171-182.	2.1	7
7	Family disadvantage, gender, and the returns to genetic human capital*. Scandinavian Journal of Economics, 2022, 124, 550-578.	0.7	8
8	Life-time Actionable Pharmacogenetic Drug Use: A Population-based Cohort Study in 86â€‰040 Young People With and Without Mental Disorders in Denmark. Pharmacopsychiatry, 2022, 55, 95-107.	1.7	3
9	Comparing Copy Number Variations in a Danish Case Cohort of Individuals With Psychiatric Disorders. JAMA Psychiatry, 2022, 79, 59.	6.0	24
10	Infection Polygenic Factors Account for a Small Proportion of the Relationship Between Infections and Mental Disorders. Biological Psychiatry, 2022, 92, 283-290.	0.7	5
11	Rare SLC13A1 variants associate with intervertebral disc disorder highlighting role of sulfate in disc pathology. Nature Communications, 2022, 13, 634.	5.8	21
12	A comprehensive map of genetic relationships among diagnostic categories based on 48.6 million relative pairs from the Danish genealogy. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	3.3	11
13	Accounting for age of onset and family history improves power in genome-wide association studies. American Journal of Human Genetics, 2022, 109, 417-432.	2.6	16
14	A genome-wide meta-analysis identifies 50 genetic loci associated with carpal tunnel syndrome. Nature Communications, 2022, 13, 1598.	5.8	8
15	Genome-wide study of early and severe childhood asthma identifies interaction between CDHR3 and GSDMB. Journal of Allergy and Clinical Immunology, 2022, 150, 622-630.	1.5	8
16	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	13.7	929
17	Genetic correlates of phenotypic heterogeneity in autism. Nature Genetics, 2022, 54, 1293-1304.	9.4	51
18	Comprehensive genome-wide association study of different forms of hernia identifies more than 80 associated loci. Nature Communications, 2022, 13, .	5.8	9

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19	Deep learningâ€“based integration of genetics with registry data for stratification of schizophrenia and depression. Science Advances, 2022, 8, .	4.7	6
20	Genome-wide association study of panic disorder reveals genetic overlap with neuroticism and depression. Molecular Psychiatry, 2021, 26, 4179-4190.	4.1	58
21	Copy Number Variants and Polygenic Risk Scores Predict Need of Care in Autism and/or ADHD Families. Journal of Autism and Developmental Disorders, 2021, 51, 276-285.	1.7	9
22	Genetic factors underlying the bidirectional relationship between autoimmune and mental disorders â€“ Findings from a Danish population-based study. Brain, Behavior, and Immunity, 2021, 91, 10-23.	2.0	8
23	Shared genetic risk between eating disorderâ€•and substanceâ€•useâ€•related phenotypes: Evidence from genomeâ€•wide association studies. Addiction Biology, 2021, 26, e12880.	1.4	28
24	Genetic predictors of educational attainment and intelligence test performance predict voter turnout. Nature Human Behaviour, 2021, 5, 281-291.	6.2	15
25	Genome-wide association study identifies 48 common genetic variants associated with handedness. Nature Human Behaviour, 2021, 5, 59-70.	6.2	79
26	Gestational age-dependent development of the neonatal metabolome. Pediatric Research, 2021, 89, 1396-1404.	1.1	16
27	Genome-wide association study across pediatric central nervous system tumors implicates shared predisposition and points to 1q25.2 (PAPPA2) and 11p12 (LRRC4C) as novel candidate susceptibility loci. Child's Nervous System, 2021, 37, 819-830.	0.6	9
28	A polygenic resilience score moderates the genetic risk for schizophrenia. Molecular Psychiatry, 2021, 26, 800-815.	4.1	36
29	Transcriptomic networks implicate neuronal energetic abnormalities in three mouse models harboring autism and schizophrenia-associated mutations. Molecular Psychiatry, 2021, 26, 1520-1534.	4.1	28
30	No evidence of associations between genetic liability for schizophrenia and development of cannabis use disorder. Psychological Medicine, 2021, 51, 479-484.	2.7	12
31	Risk variants and polygenic architecture of disruptive behavior disorders in the context of attention-deficit/hyperactivity disorder. Nature Communications, 2021, 12, 576.	5.8	28
32	Identification of genetic loci associated with nocturnal enuresis: a genome-wide association study. The Lancet Child and Adolescent Health, 2021, 5, 201-209.	2.7	27
33	Risk of Early-Onset Depression Associated With Polygenic Liability, Parental Psychiatric History, and Socioeconomic Status. JAMA Psychiatry, 2021, 78, 387.	6.0	33
34	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
35	Pharmacogenetic genotype and phenotype frequencies in a large Danish population-based case-cohort sample. Translational Psychiatry, 2021, 11, 294.	2.4	13
36	A large-scale investigation into the role of classical HLA loci in multiple types of severe infections, with a focus on overlaps with autoimmune and mental disorders. Journal of Translational Medicine, 2021, 19, 230.	1.8	5

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37	The genetic structure of Norway. European Journal of Human Genetics, 2021, 29, 1710-1718.	1.4	10
38	Leveraging both individual-level genetic data and GWAS summary statistics increases polygenic prediction. American Journal of Human Genetics, 2021, 108, 1001-1011.	2.6	22
39	Examining Sex-Differentiated Genetic Effects Across Neuropsychiatric and Behavioral Traits. Biological Psychiatry, 2021, 89, 1127-1137.	0.7	48
40	Polygenic Heterogeneity Across Obsessive-Compulsive Disorder Subgroups Defined by a Comorbid Diagnosis. Frontiers in Genetics, 2021, 12, 711624.	1.1	7
41	Translating polygenic risk scores for clinical use by estimating the confidence bounds of risk prediction. Nature Communications, 2021, 12, 5276.	5.8	12
42	Identifying dominant-negative actions of a dopamine transporter variant in patients with parkinsonism and neuropsychiatric disease. JCI Insight, 2021, 6, .	2.3	11
43	Anorexia nervosa and inflammatory bowel diseasesâ€"Diagnostic and genetic associations. JCPP Advances, 2021, 1, e12036.	1.4	9
44	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. JAMA Psychiatry, 2021, 78, 1258.	6.0	88
45	Genetic, Clinical, and Sociodemographic Factors Associated With Stimulant Treatment Outcomes in ADHD. American Journal of Psychiatry, 2021, 178, 854-864.	4.0	15
46	CYP2D6 Genotyping and Antipsychotic-Associated Extrapyramidal Adverse Effects in a Randomized Trial of Aripiprazole Versus Quetiapine Extended Release in Children and Adolescents, Aged 12â€"17 Years, With First Episode Psychosis. Journal of Clinical Psychopharmacology, 2021, 41, 667-672.	0.7	2
47	Associations between patterns in comorbid diagnostic trajectories of individuals with schizophrenia and etiological factors. Nature Communications, 2021, 12, 6617.	5.8	9
48	Pleiotropy between language impairment and broader behavioral disordersâ€"an investigation of both common and rare genetic variants. Journal of Neurodevelopmental Disorders, 2021, 13, 54.	1.5	5
49	Genetics of suicide attempts in individuals with and without mental disorders: a population-based genome-wide association study. Molecular Psychiatry, 2020, 25, 2410-2421.	4.1	124
50	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. Biological Psychiatry, 2020, 87, 419-430.	0.7	27
51	Language deficits in specific language impairment, attention deficit/hyperactivity disorder, and autism spectrum disorder: An analysis of polygenic risk. Autism Research, 2020, 13, 369-381.	2.1	17
52	A major role for common genetic variation in anxiety disorders. Molecular Psychiatry, 2020, 25, 3292-3303.	4.1	243
53	Molecular genetic overlap between posttraumatic stress disorder and sleep phenotypes. Sleep, 2020, 43, .	0.6	32
54	Genetic liability to ADHD and substance use disorders in individuals with ADHD. Addiction, 2020, 115, 1368-1377.	1.7	47

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55	Polygenic risk score, psychosocial environment and the risk of attention-deficit/hyperactivity disorder. Translational Psychiatry, 2020, 10, 335.	2.4	22
56	Genomic and drug target evaluation of 90 cardiovascular proteins in 30,931 individuals. Nature Metabolism, 2020, 2, 1135-1148.	5.1	327
57	A large-scale genome-wide association study meta-analysis of cannabis use disorder. Lancet Psychiatry,the, 2020, 7, 1032-1045.	3.7	200
58	The Duffy-null genotype and risk of infection. Human Molecular Genetics, 2020, 29, 3341-3349.	1.4	11
59	Polygenic Risk and Progression to Bipolar or Psychotic Disorders Among Individuals Diagnosed With Unipolar Depression in Early Life. American Journal of Psychiatry, 2020, 177, 936-943.	4.0	40
60	PCM1 is necessary for focal ciliary integrity and is a candidate for severe schizophrenia. Nature Communications, 2020, 11, 5903.	5.8	13
61	The genetic architecture of sporadic and multiple consecutive miscarriage. Nature Communications, 2020, 11, 5980.	5.8	52
62	Genetic liability to major depression and risk of childhood asthma. Brain, Behavior, and Immunity, 2020, 89, 433-439.	2.0	5
63	Population genomics of the Viking world. Nature, 2020, 585, 390-396.	13.7	143
64	Habitual sleep disturbances and migraine: a Mendelian randomization study. Annals of Clinical and Translational Neurology, 2020, 7, 2370-2380.	1.7	18
65	Transethnic Genome-Wide Association Study Provides Insights in the Genetic Architecture and Heritability of Long QT Syndrome. Circulation, 2020, 142, 324-338.	1.6	83
66	Genetic stratification of depression in UK Biobank. Translational Psychiatry, 2020, 10, 163.	2.4	19
67	A large population-based investigation into the genetics of susceptibility to gastrointestinal infections and the link between gastrointestinal infections and mental illness. Human Genetics, 2020, 139, 593-604.	1.8	14
68	Association between GLP-1 receptor gene polymorphisms with reward learning, anhedonia and depression diagnosis. Acta Neuropsychiatrica, 2020, 32, 218-225.	1.0	8
69	Impute.me: An Open-Source, Non-profit Tool for Using Data From Direct-to-Consumer Genetic Testing to Calculate and Interpret Polygenic Risk Scores. Frontiers in Genetics, 2020, 11, 578.	1.1	47
70	Quantitative genome-wide association analyses of receptive language in the Danish High Risk and Resilience Study. BMC Neuroscience, 2020, 21, 30.	0.8	7
71	Large-Scale Exome Sequencing Study Implicates Both Developmental and Functional Changes in the Neurobiology of Autism. Cell, 2020, 180, 568-584.e23.	13.5	1,422
72	Adolescent residential mobility, genetic liability and risk of schizophrenia, bipolar disorder and major depression. British Journal of Psychiatry, 2020, 217, 390-396.	1.7	11

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73	Analysis of genes within the schizophrenia-linked 22q11.2 deletion identifies interaction of night owl/LZTR1 and NF1 in GABAergic sleep control. PLoS Genetics, 2020, 16, e1008727.	1.5	20
74	Association between Mental Disorders and Subsequent Medical Conditions. New England Journal of Medicine, 2020, 382, 1721-1731.	13.9	258
75	Effect of Routine Cytochrome P450 2D6 and 2C19 Genotyping on Antipsychotic Drug Persistence in Patients With Schizophrenia. JAMA Network Open, 2020, 3, e2027909.	2.8	14
76	FUT2â€“ABO epistasis increases the risk of early childhood asthma and Streptococcus pneumoniae respiratory illnesses. Nature Communications, 2020, 11, 6398.	5.8	21
77	Association Between Childhood Green Space, Genetic Liability, and the Incidence of Schizophrenia. Schizophrenia Bulletin, 2020, 46, 1629-1637.	2.3	28
78	Genome-wide association study identifies 16 genomic regions associated with circulating cytokines at birth. PLoS Genetics, 2020, 16, e1009163.	1.5	12
79	Title is missing!. , 2020, 16, e1009163.		0
80	Title is missing!. , 2020, 16, e1009163.		0
81	Title is missing!. , 2020, 16, e1009163.		0
82	Title is missing!. , 2020, 16, e1009163.		0
83	Can Animal Models of Copy Number Variants That Predispose to Schizophrenia Elucidate Underlying Biology?. Biological Psychiatry, 2019, 85, 13-24.	0.7	34
84	A CROSS-MENTAL-DISORDERS GWAS IN THE DANISH NATIONAL HEALTH REGISTER. European Neuropsychopharmacology, 2019, 29, S996.	0.3	0
85	Genome-wide association study identifies eight risk loci and implicates metabo-psychiatric origins for anorexia nervosa. Nature Genetics, 2019, 51, 1207-1214.	9.4	641
86	Herpes Simplex Virus Type 1 infection is associated with suicidal behavior and first registered psychiatric diagnosis in a healthy population. Psychoneuroendocrinology, 2019, 108, 150-154.	1.3	10
87	9IDENTICAL BY DESCENT SEGMENTS ASSOCIATES WITH SCHIZOPHRENIA. European Neuropsychopharmacology, 2019, 29, S1071.	0.3	0
88	International meta-analysis of PTSD genome-wide association studies identifies sex- and ancestry-specific genetic risk loci. Nature Communications, 2019, 10, 4558.	5.8	363
89	Reduced neonatal brain-derived neurotrophic factor is associated with autism spectrum disorders. Translational Psychiatry, 2019, 9, 252.	2.4	56
90	Association of Childhood Exposure to Nitrogen Dioxide and Polygenic Risk Score for Schizophrenia With the Risk of Developing Schizophrenia. JAMA Network Open, 2019, 2, e1914401.	2.8	29

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91	Genetic risk scores for major psychiatric disorders and the risk of postpartum psychiatric disorders. Translational Psychiatry, 2019, 9, 288.	2.4	27
92	A large-scale genomic investigation of susceptibility to infection and its association with mental disorders in the Danish population. Translational Psychiatry, 2019, 9, 283.	2.4	46
93	Variants in the fetal genome near pro-inflammatory cytokine genes on 2q13 associate with gestational duration. Nature Communications, 2019, 10, 3927.	5.8	49
94	Schizophrenia polygenic risk scores, urbanicity and treatment-resistant schizophrenia. Schizophrenia Research, 2019, 212, 79-85.	1.1	19
95	F28SINGLE-CELL ENRICHMENT ANALYSIS FOR THE IDENTIFICATION OF TRAIT-RELEVANT GENES AND BRAIN CELL TYPES IN PSYCHIATRIC DISORDERS. European Neuropsychopharmacology, 2019, 29, S1124.	0.3	0
96	Large-scale study of Toxoplasma and Cytomegalovirus shows an association between infection and serious psychiatric disorders. Brain, Behavior, and Immunity, 2019, 79, 152-158.	2.0	107
97	A genome-wide association study of shared risk across psychiatric disorders implicates gene regulation during fetal neurodevelopment. Nature Neuroscience, 2019, 22, 353-361.	7.1	173
98	Association of Polygenic Liabilities for Major Depression, Bipolar Disorder, and Schizophrenia With Risk for Depression in the Danish Population. JAMA Psychiatry, 2019, 76, 516.	6.0	78
99	Genome-wide association study implicates CHRNA2 in cannabis use disorder. Nature Neuroscience, 2019, 22, 1066-1074.	7.1	94
100	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
101	DBDS Genomic Cohort, a prospective and comprehensive resource for integrative and temporal analysis of genetic, environmental and lifestyle factors affecting health of blood donors. BMJ Open, 2019, 9, e028401.	0.8	68
102	Genetic Variants Associated With Anxiety and Stress-Related Disorders. JAMA Psychiatry, 2019, 76, 924.	6.0	140
103	GENOME-WIDE METHYLOMIC ANALYSIS OF NEONATAL BLOOD FROM DANISH TWINS DISCORDANT FOR MENTAL ILLNESS. European Neuropsychopharmacology, 2019, 29, S794-S795.	0.3	0
104	ESTIMATED DNA METHYLATION GESTATIONAL AGE IN NEWBORN MONOZYGOTIC TWINS ASSOCIATE WITH LATER PSYCHIATRIC DISORDERS BETWEEN CON/DISCORDANT PAIRS. European Neuropsychopharmacology, 2019, 29, S795.	0.3	0
105	Post-traumatic stress following military deployment: Genetic associations and cross-disorder genetic correlations. Journal of Affective Disorders, 2019, 252, 350-357.	2.0	12
106	Genome-wide association study identifies 30 loci associated with bipolar disorder. Nature Genetics, 2019, 51, 793-803.	9.4	1,191
107	Human Disease Variation in the Light of Population Genomics. Cell, 2019, 177, 115-131.	13.5	75
108	Variable DNA methylation in neonates mediates the association between prenatal smoking and birth weight. Philosophical Transactions of the Royal Society B: Biological Sciences, 2019, 374, 20180120.	1.8	46

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109	Immunity and mental illness: findings from a Danish population-based immunogenetic study of seven psychiatric and neurodevelopmental disorders. European Journal of Human Genetics, 2019, 27, 1445-1455.	1.4	38
110	Roadmap for a precision-medicine initiative in the Nordic region. Nature Genetics, 2019, 51, 924-930.	9.4	22
111	Identification of common genetic risk variants for autism spectrum disorder. Nature Genetics, 2019, 51, 431-444.	9.4	1,538
112	THE IMPACT OF CNVs ON ASD/ADHD RISK IN MULTIPLEX FAMILIES. European Neuropsychopharmacology, 2019, 29, S895.	0.3	0
113	SA133GENETIC PREDICTION OF TRANSITION FROM CHILDHOOD AND ADOLESCENT PSYCHIATRIC DISORDER TO SCHIZOPHRENIA. European Neuropsychopharmacology, 2019, 29, S1262.	0.3	0
114	Autism spectrum disorder and attention deficit hyperactivity disorder have a similar burden of rare protein-truncating variants. Nature Neuroscience, 2019, 22, 1961-1965.	7.1	148
115	Clinical association to FKBP5 rs1360780 in patients with depression. Psychiatric Genetics, 2019, 29, 220-225.	0.6	3
116	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. Cell, 2019, 179, 1469-1482.e11.	13.5	935
117	Interactive effects between hemizygous 15q13.3 microdeletion and peripubertal stress on adult behavioral functions. Neuropsychopharmacology, 2019, 44, 703-710.	2.8	8
118	Association of Whole-Genome and NETRIN1 Signaling Pathwayâ€"Derived Polygenic Risk Scores for Major Depressive Disorder and White Matter Microstructure in the UK Biobank. Biological Psychiatry: Cognitive Neuroscience and Neuroimaging, 2019, 4, 91-100.	1.1	16
119	Discovery of the first genome-wide significant risk loci for attention deficit/hyperactivity disorder. Nature Genetics, 2019, 51, 63-75.	9.4	1,594
120	Exploring Comorbidity Within Mental Disorders Among a Danish National Population. JAMA Psychiatry, 2019, 76, 259.	6.0	374
121	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
122	Altered auditory processing and effective connectivity in 22q11.2 deletion syndrome. Schizophrenia Research, 2018, 197, 328-336.	1.1	24
123	Shared molecular neuropathology across major psychiatric disorders parallels polygenic overlap. Science, 2018, 359, 693-697.	6.0	851
124	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. Nature Genetics, 2018, 50, 668-681.	9.4	2,224
125	Restless legs syndrome is associated with major comorbidities in a population of Danish blood donors. Sleep Medicine, 2018, 45, 124-131.	0.8	23
126	22q11.2 Deletion Syndrome Is Associated With Impaired Auditory Steady-State Gamma Response. Schizophrenia Bulletin, 2018, 44, 388-397.	2.3	33

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127	Does Childhood Trauma Moderate Polygenic Risk for Depression? A Meta-analysis of 5765 Subjects From the Psychiatric Genomics Consortium. Biological Psychiatry, 2018, 84, 138-147.	0.7	87
128	Heritability of Schizophrenia and Schizophrenia Spectrum Based on the Nationwide Danish Twin Register. Biological Psychiatry, 2018, 83, 492-498.	0.7	374
129	A Genetic Investigation of Sex Bias in the Prevalence of Attention-Deficit/Hyperactivity Disorder. Biological Psychiatry, 2018, 83, 1044-1053.	0.7	146
130	Schizophrenia-associated mt-DNA SNPs exhibit highly variable haplogroup affiliation and nuclear ancestry: Bi-genomic dependence raises major concerns for link to disease. PLoS ONE, 2018, 13, e0208828.	1.1	15
131	Complex spatio-temporal distribution and genomic ancestry of mitochondrial DNA haplogroups in 24,216 Danes. PLoS ONE, 2018, 13, e0208829.	1.1	5
132	AMPK signaling linked to the schizophrenia-associated 1q21.1 deletion is required for neuronal and sleep maintenance. PLoS Genetics, 2018, 14, e1007623.	1.5	25
133	Elevated expression of a minor isoform of ANK3 is a risk factor for bipolar disorder. Translational Psychiatry, 2018, 8, 210.	2.4	24
134	The Anorexia Nervosa Genetics Initiative (ANGI): Overview and methods. Contemporary Clinical Trials, 2018, 74, 61-69.	0.8	73
135	Comorbidity of migraine with ADHD in adults. BMC Neurology, 2018, 18, 147.	0.8	24
136	Quantifying the Impact of Rare and Ultra-rare Coding Variation across the Phenotypic Spectrum. American Journal of Human Genetics, 2018, 102, 1204-1211.	2.6	102
137	Ancient genomes from Iceland reveal the making of a human population. Science, 2018, 360, 1028-1032.	6.0	62
138	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	6.0	1,085
139	Elevated polygenic burden for autism is associated with differential DNA methylation at birth. Genome Medicine, 2018, 10, 19.	3.6	88
140	Prevalence of rearrangements in the 22q11.2 region and population-based risk of neuropsychiatric and developmental disorders in a Danish population: a case-cohort study. Lancet Psychiatry,the, 2018, 5, 573-580.	3.7	102
141	Risk of Psychiatric Disorders Among Individuals With the 22q11.2 Deletion or Duplication. JAMA Psychiatry, 2017, 74, 282.	6.0	89
142	Is an Early Age at Illness Onset in Schizophrenia Associated With Increased Genetic Susceptibility? Analysis of Data From the Nationwide Danish Twin Register. EBioMedicine, 2017, 18, 320-326.	2.7	22
143	Polygenic transmission disequilibrium confirms that common and rare variation act additively to create risk for autism spectrum disorders. Nature Genetics, 2017, 49, 978-985.	9.4	401
144	Quetiapine extended release versus aripiprazole in children and adolescents with first-episode psychosis: the multicentre, double-blind, randomised tolerability and efficacy of antipsychotics (TEA) trial. Lancet Psychiatry,the, 2017, 4, 605-618.	3.7	40

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145	Multi-State Survival Analysis Of Multi-Psychiatry Disorders. European Neuropsychopharmacology, 2017, 27, S368-S369.	0.3	0
146	5. Mitochondrial DNA Haplogroups are Associated with Psychiatric Disease: A Nation-Wide Study of 74,763 Danes. Biological Psychiatry, 2017, 81, S3.	0.7	0
147	Severe Parkinsonism and Creatine Kinase Increase After Low-Dose Aripiprazole Treatment in a Patient of African Descent. Journal of Clinical Psychopharmacology, 2017, 37, 630-631.	0.7	4
148	A mouse model of the schizophrenia-associated 1q21.1 microdeletion syndrome exhibits altered mesolimbic dopamine transmission. Translational Psychiatry, 2017, 7, 1261.	2.4	37
149	Prevalence of restless legs syndrome and associated factors in an otherwise healthy population: results from the Danish Blood Donor Study. Sleep Medicine, 2017, 36, 55-61.	0.8	51
150	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
151	Lessons to be Learned From 22q2.11 Syndromesâ€"Reply. JAMA Psychiatry, 2017, 74, 757.	6.0	2
152	Persistent gating deficit and increased sensitivity to NMDA receptor antagonism after puberty in a new mouse model of the human 22q11.2 microdeletion syndrome: a study in male mice. Journal of Psychiatry and Neuroscience, 2017, 42, 48-58.	1.4	63
153	Estimating Effect Sizes and Expected Replication Probabilities from GWAS Summary Statistics. Frontiers in Genetics, 2016, 7, 15.	1.1	40
154	Investigation of SNP rs2060546 Immediately Upstream to NTN4 in a Danish Gilles de la Tourette Syndrome Cohort. Frontiers in Neuroscience, 2016, 10, 531.	1.4	5
155	Leveraging Genomic Annotations and Pleiotropic Enrichment for Improved Replication Rates in Schizophrenia GWAS. PLoS Genetics, 2016, 12, e1005803.	1.5	34
156	Parental Origin of Interstitial Duplications at 15q11.2-q13.3 in Schizophrenia and Neurodevelopmental Disorders. PLoS Genetics, 2016, 12, e1005993.	1.5	51
157	Meta-analysis of 375,000 individuals identifies 38 susceptibility loci for migraine. Nature Genetics, 2016, 48, 856-866.	9.4	520
158	Genomeâ€wide association study reveals greater polygenic loading for schizophrenia in cases with a family history of illness. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2016, 171, 276-289.	1.1	28
159	Genetic risk for autism spectrum disorders and neuropsychiatric variation in the general population. Nature Genetics, 2016, 48, 552-555.	9.4	326
160	Schizophrenia Spectrum Disorders in a Danish 22q11.2 Deletion Syndrome Cohort Compared to the Total Danish Populationâ€"A Nationwide Register Study. Schizophrenia Bulletin, 2016, 42, 824-831.	2.3	14
161	Identification of rare high-risk copy number variants affecting the dopamine transporter gene in mental disorders. Nordic Journal of Psychiatry, 2016, 70, 276-279.	0.7	2
162	Association of AADAC Deletion and Gilles de la Tourette Syndrome in a Large European Cohort. Biological Psychiatry, 2016, 79, 383-391.	0.7	41

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163	The influence of genetic constitution on migraine drug responses. Cephalalgia, 2016, 36, 624-639.	1.8	34
164	Genetic Markers of Human Evolution Are Enriched in Schizophrenia. Biological Psychiatry, 2016, 80, 284-292.	0.7	92
165	The association between candidate migraine susceptibility loci and severe migraine phenotype in a clinical sample. Cephalalgia, 2016, 36, 615-623.	1.8	24
166	Two rare deletions upstream of the NRXN1 gene (2p16.3) affecting the non-coding mRNA AK127244 segregate with diverse psychopathological phenotypes in a family. European Journal of Medical Genetics, 2015, 58, 650-653.	0.7	12
167	The Danish 22q11 research initiative. BMC Psychiatry, 2015, 15, 220.	1.1	14
168	An Empirical Bayes Mixture Model for Effect Size Distributions in Genome-Wide Association Studies. PLoS Genetics, 2015, 11, e1005717.	1.5	22
169	Combinations of Genetic Data Present in Bipolar Patients, but Absent in Control Persons. PLoS ONE, 2015, 10, e0143432.	1.1	4
170	Linkage and whole genome sequencing identify a locus on 6q25â€“26 for formal thought disorder and implicate MEF2A regulation. Schizophrenia Research, 2015, 169, 441-446.	1.1	12
171	Individualization of treatments with drugs metabolized by CES1: combining genetics and metabolomics. Pharmacogenomics, 2015, 16, 649-665.	0.6	19
172	Association Study of CHRNA7 Promoter Variants with Sensory and Sensorimotor Gating in Schizophrenia Patients and Healthy Controls: A Danish Caseâ€“Control Study. NeuroMolecular Medicine, 2015, 17, 423-430.	1.8	23
173	Usefulness of the SNP microarray technology to identify rare mutations in the case of perinatal death. Case Reports in Perinatal Medicine, 2015, 4, .	0.1	1
174	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
175	Population genetic differentiation of height and body mass index across Europe. Nature Genetics, 2015, 47, 1357-1362.	9.4	227
176	Soluble Urokinase-Type Plasminogen Activator Receptor Levels in Patients With Schizophrenia. Schizophrenia Bulletin, 2015, 41, 764-771.	2.3	16
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