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	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
2	Large recurrent microdeletions associated with schizophrenia. Nature, 2008, 455, 232-236.	13.7	1,619
3	Discovery of the first genome-wide significant risk loci for attention deficit/hyperactivity disorder. Nature Genetics, 2019, 51, 63-75.	9.4	1,594
4	Common variants conferring risk of schizophrenia. Nature, 2009, 460, 744-747.	13.7	1,572
5	Identification of common genetic risk variants for autism spectrum disorder. Nature Genetics, 2019, 51, 431-444.	9.4	1,538
6	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
7	Common schizophrenia alleles are enriched in mutation-intolerant genes and in regions under strong background selection. Nature Genetics, 2018, 50, 381-389.	9.4	1,332
8	Genome-wide association study identifies 30 loci associated with bipolar disorder. Nature Genetics, 2019, 51, 793-803.	9.4	1,191
9	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	6.0	1,085
10	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. Cell, 2019, 179, 1469-1482.e11.	13.5	935
11	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	13.7	929
12	Shared molecular neuropathology across major psychiatric disorders parallels polygenic overlap. Science, 2018, 359, 693-697.	6.0	851
13	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
14	Sequence variants affecting eosinophil numbers associate with asthma and myocardial infarction. Nature Genetics, 2009, 41, 342-347.	9.4	709
15	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
16	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
17	Meta-analysis of 375,000 individuals identifies 38 susceptibility loci for migraine. Nature Genetics, 2016, 48, 856-866.	9.4	520
18	Correlation between Genetic and Geographic Structure in Europe. Current Biology, 2008, 18, 1241-1248.	1.8	449

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19	Disruption of the neurexin 1 gene is associated with schizophrenia. Human Molecular Genetics, 2009, 18, 988-996.	1.4	424
20	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
21	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
22	Heritability of Schizophrenia and Schizophrenia Spectrum Based on the Nationwide Danish Twin Register. Biological Psychiatry, 2018, 83, 492-498.	0.7	374
23	Exploring Comorbidity Within Mental Disorders Among a Danish National Population. JAMA Psychiatry, 2019, 76, 259.	6.0	374
24	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
25	Genome-wide association study of migraine implicates a common susceptibility variant on 8q22.1. Nature Genetics, 2010, 42, 869-873.	9.4	332
26	Genomic and drug target evaluation of 90 cardiovascular proteins in 30,931 individuals. Nature Metabolism, 2020, 2, 1135-1148.	5.1	327
27	Genetic risk for autism spectrum disorders and neuropsychiatric variation in the general population. Nature Genetics, 2016, 48, 552-555.	9.4	326
28	Improved Detection of Common Variants Associated with Schizophrenia and Bipolar Disorder Using Pleiotropy-Informed Conditional False Discovery Rate. PLoS Genetics, 2013, 9, e1003455.	1.5	298
29	Impaired glutathione synthesis in schizophrenia: Convergent genetic and functional evidence. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 16621-16626.	3.3	275
30	A rare variant in MYH6 is associated with high risk of sick sinus syndrome. Nature Genetics, 2011, 43, 316-320.	9.4	275
31	Association between Mental Disorders and Subsequent Medical Conditions. New England Journal of Medicine, 2020, 382, 1721-1731.	13.9	258
32	A major role for common genetic variation in anxiety disorders. Molecular Psychiatry, 2020, 25, 3292-3303.	4.1	243
33	Using Electronic Patient Records to Discover Disease Correlations and Stratify Patient Cohorts. PLoS Computational Biology, 2011, 7, e1002141.	1.5	236
34	Brain Expressed microRNAs Implicated in Schizophrenia Etiology. PLoS ONE, 2007, 2, e873.	1.1	235
35	Population genetic differentiation of height and body mass index across Europe. Nature Genetics, 2015, 47, 1357-1362.	9.4	227
36	Common variants on 8p12 and 1q24.2 confer risk of schizophrenia. Nature Genetics, 2011, 43, 1224-1227.	9.4	224

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37	Schizophrenia and Oxidative Stress: Glutamate Cysteine Ligase Modifier as a Susceptibility Gene. American Journal of Human Genetics, 2006, 79, 586-592.	2.6	209
38	A large-scale genome-wide association study meta-analysis of cannabis use disorder. Lancet Psychiatry,the, 2020, 7, 1032-1045.	3.7	200
39	Common variants at VRK2 and TCF4 conferring risk of schizophrenia. Human Molecular Genetics, 2011, 20, 4076-4081.	1.4	193
40	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
41	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
42	Clozapine-induced agranulocytosis is associated with rare HLA-DQB1 and HLA-B alleles. Nature Communications, 2014, 5, 4757.	5.8	153
43	The Complement Control-Related Genes CSMD1 and CSMD2 Associate to Schizophrenia. Biological Psychiatry, 2011, 70, 35-42.	0.7	149
44	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
45	A Genetic Investigation of Sex Bias in the Prevalence of Attention-Deficit/Hyperactivity Disorder. Biological Psychiatry, 2018, 83, 1044-1053.	0.7	146
46	Population genomics of the Viking world. Nature, 2020, 585, 390-396.	13.7	143
47	Genetic Variants Associated With Anxiety and Stress-Related Disorders. JAMA Psychiatry, 2019, 76, 924.	6.0	140
48	A Comprehensive Family-Based Replication Study of Schizophrenia Genes. JAMA Psychiatry, 2013, 70, 573.	6.0	138
49	Reliability of clinical ICD-10 schizophrenia diagnoses. Nordic Journal of Psychiatry, 2005, 59, 209-212.	0.7	133
50	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
51	At-Risk Variant in TCF7L2 for Type II Diabetes Increases Risk of Schizophrenia. Biological Psychiatry, 2011, 70, 59-63.	0.7	114
52	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
53	Redox Dysregulation in the Pathophysiology of Schizophrenia and Bipolar Disorder: Insights from Animal Models. Antioxidants and Redox Signaling, 2013, 18, 1428-1443.	2.5	102
54	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

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55	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
56	Dose-Specific Adverse Drug Reaction Identification in Electronic Patient Records: Temporal Data Mining in an Inpatient Psychiatric Population. Drug Safety, 2014, 37, 237-247.	1.4	96
57	Maternally Derived Microduplications at 15q11-q13: Implication of Imprinted Genes in Psychotic Illness. American Journal of Psychiatry, 2011, 168, 408-417.	4.0	95
58	A Mouse Model that Recapitulates Cardinal Features of the 15q13.3 Microdeletion Syndrome Including Schizophrenia- and Epilepsy-Related Alterations. Biological Psychiatry, 2014, 76, 128-137.	0.7	95
59	MicroRNAs Show Mutually Exclusive Expression Patterns in the Brain of Adult Male Rats. PLoS ONE, 2009, 4, e7225.	1.1	94
60	Genome-wide association study implicates CHRNA2 in cannabis use disorder. Nature Neuroscience, 2019, 22, 1066-1074.	7.1	94
61	Genetic Markers of Human Evolution Are Enriched in Schizophrenia. Biological Psychiatry, 2016, 80, 284-292.	0.7	92
62	Risk of Psychiatric Disorders Among Individuals With the 22q11.2 Deletion or Duplication. JAMA Psychiatry, 2017, 74, 282.	6.0	89
63	Elevated polygenic burden for autism is associated with differential DNA methylation at birth. Genome Medicine, 2018, 10, 19.	3.6	88
64	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. JAMA Psychiatry, 2021, 78, 1258.	6.0	88
65	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
66	Interaction between genetic polymorphisms and stressful life events in first episode depression. Journal of Affective Disorders, 2009, 119, 107-115.	2.0	84
67	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
68	The Muscarinic M1/M4 Receptor Agonist Xanomeline Exhibits Antipsychotic-Like Activity in Cebus apella Monkeys. Neuropsychopharmacology, 2003, 28, 1168-1175.	2.8	82
69	Genome-wide association study identifies 48 common genetic variants associated with handedness. Nature Human Behaviour, 2021, 5, 59-70.	6.2	79
70	Association Study of Nonsynonymous Single Nucleotide Polymorphisms in Schizophrenia. Biological Psychiatry, 2012, 71, 169-177.	0.7	78
71	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
72	Human Disease Variation in the Light of Population Genomics. Cell, 2019, 177, 115-131.	13.5	75

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73	The Anorexia Nervosa Genetics Initiative (ANGI): Overview and methods. Contemporary Clinical Trials, 2018, 74, 61-69.	0.8	73
74	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
75	Polymorphisms of Serotonin Receptor 2A and 2C Genes and COMT in Relation to Obesity and Type 2 Diabetes. PLoS ONE, 2009, 4, e6696.	1.1	65
76	Kynurenine 3-monooxygenase polymorphisms: relevance for kynurenic acid synthesis in patients with schizophrenia and healthy controls. Journal of Psychiatry and Neuroscience, 2012, 37, 53-57.	1.4	65
77	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
78	Ancient genomes from Iceland reveal the making of a human population. Science, 2018, 360, 1028-1032.	6.0	62
79	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. Biological Psychiatry, 2022, 91, 102-117.	0.7	61
80	Genome-wide association study of panic disorder reveals genetic overlap with neuroticism and depression. Molecular Psychiatry, 2021, 26, 4179-4190.	4.1	58
81	Reduced neonatal brain-derived neurotrophic factor is associated with autism spectrum disorders. Translational Psychiatry, 2019, 9, 252.	2.4	56
82	Enhanced Maternal Origin of the 22q11.2 Deletion in Velocardiofacial and DiGeorge Syndromes. American Journal of Human Genetics, 2013, 92, 439-447.	2.6	53
83	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
84	Prenatal stress may increase vulnerability to life events: Comparison with the effects of prenatal dexamethasone. Developmental Brain Research, 2005, 159, 55-63.	2.1	52
85	The genetic architecture of sporadic and multiple consecutive miscarriage. Nature Communications, 2020, 11, 5980.	5.8	52
86	CYP2D6 Genotype Predicts Antipsychotic Side Effects in Schizophrenia Inpatients: A Retrospective Matched Case-Control Study. Neuropsychobiology, 2009, 59, 222-226.	0.9	51
87	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
88	Parental Origin of Interstitial Duplications at 15q11.2-q13.3 in Schizophrenia and Neurodevelopmental Disorders. PLoS Genetics, 2016, 12, e1005993.	1.5	51
89	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
90	Genetic correlates of phenotypic heterogeneity in autism. Nature Genetics, 2022, 54, 1293-1304.	9.4	51

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91	Effects of prenatal exposure to chronic mild stress and toluene in rats. Neurotoxicology and Teratology, 2005, 27, 153-167.	1.2	50
92	Association Between the CCR5 32-bp Deletion Allele and Late Onset of Schizophrenia. American Journal of Psychiatry, 2006, 163, 507-511.	4.0	49
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	Examining Sex-Differentiated Genetic Effects Across Neuropsychiatric and Behavioral Traits. Biological Psychiatry, 2021, 89, 1127-1137.	0.7	48
95	Promoter variants in IL18 are associated with onset of depression in patients previously exposed to stressful-life events. Journal of Affective Disorders, 2012, 136, 134-138.	2.0	47
96	Genetic liability to ADHD and substance use disorders in individuals with ADHD. Addiction, 2020, 115, 1368-1377.	1.7	47
97	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
98	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
99	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
100	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
101	The tryptophan hydroxylase 1 (TPH1) gene, schizophrenia susceptibility, and suicidal behavior: A multiâ€centre caseâ€“control study and metaâ€analysis. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 387-396.	1.1	45
102	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
103	Genome-Wide Association Study Identifies Four Loci Associated with Eruption of Permanent Teeth. PLoS Genetics, 2011, 7, e1002275.	1.5	42
104	Association of AADAC Deletion and Gilles de la Tourette Syndrome in a Large European Cohort. Biological Psychiatry, 2016, 79, 383-391.	0.7	41
105	Gene-Based Analysis of Regionally Enriched Cortical Genes in GWAS Data Sets of Cognitive Traits and Psychiatric Disorders. PLoS ONE, 2012, 7, e31687.	1.1	40
106	Estimating Effect Sizes and Expected Replication Probabilities from GWAS Summary Statistics. Frontiers in Genetics, 2016, 7, 15.	1.1	40
107	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
108	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

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109	Association between a disrupted-in-schizophrenia 1 (DISC1) single nucleotide polymorphism and schizophrenia in a combined Scandinavian caseâ€“control sample. Schizophrenia Research, 2008, 106, 237-241.	1.1	39
110	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
111	A mouse model of the schizophrenia-associated 1q21.1 microdeletion syndrome exhibits altered mesolimbic dopamine transmission. Translational Psychiatry, 2017, 7, 1261.	2.4	37
112	A polygenic resilience score moderates the genetic risk for schizophrenia. Molecular Psychiatry, 2021, 26, 800-815.	4.1	36
113	Estrogen receptor alpha and risk for cognitive impairment in postmenopausal women. Psychiatric Genetics, 2006, 16, 85-88.	0.6	35
114	An exploratory model for GÃ—E interaction on hippocampal volume in schizophrenia; obstetric complications and hypoxia-related genes. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2010, 34, 1259-1265.	2.5	35
115	Leveraging Genomic Annotations and Pleiotropic Enrichment for Improved Replication Rates in Schizophrenia GWAS. PLoS Genetics, 2016, 12, e1005803.	1.5	34
116	The influence of genetic constitution on migraine drug responses. Cephalalgia, 2016, 36, 624-639.	1.8	34
117	Can Animal Models of Copy Number Variants That Predispose to Schizophrenia Elucidate Underlying Biology?. Biological Psychiatry, 2019, 85, 13-24.	0.7	34
118	<i>DTNBP1, NRG1, DAOA</i>, <i>DAO</i> and <i>GRM3</i> Polymorphisms and Schizophrenia: An Association Study. Neuropsychobiology, 2009, 59, 142-150.	0.9	33
119	Genetic susceptibility factors for multiple chemical sensitivity revisited. International Journal of Hygiene and Environmental Health, 2010, 213, 131-139.	2.1	33
120	22q11.2 Deletion Syndrome Is Associated With Impaired Auditory Steady-State Gamma Response. Schizophrenia Bulletin, 2018, 44, 388-397.	2.3	33
121	Risk of Early-Onset Depression Associated With Polygenic Liability, Parental Psychiatric History, and Socioeconomic Status. JAMA Psychiatry, 2021, 78, 387.	6.0	33
122	Association between methylenetetrahydrofolate reductase (MTHFR) C677T polymorphism and age of onset in schizophrenia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 610-618.	1.1	32
123	Molecular genetic overlap between posttraumatic stress disorder and sleep phenotypes. Sleep, 2020, 43, .	0.6	32
124	No interactions between genetic polymorphisms and stressful life events on outcome of antidepressant treatment. European Neuropsychopharmacology, 2010, 20, 327-335.	0.3	31
125	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
126	Identifying the Common Genetic Basis of Antidepressant Response. Biological Psychiatry Global Open Science, 2022, 2, 115-126.	1.0	31

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127	DCLK1 Variants Are Associated across Schizophrenia and Attention Deficit/Hyperactivity Disorder. PLoS ONE, 2012, 7, e35424.	1.1	30
128	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
129	Antidepressive-drug-induced bodyweight gain is associated with polymorphisms in genes coding for COMT and TPH1. International Clinical Psychopharmacology, 2009, 24, 199-203.	0.9	28
130	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
131	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
132	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
133	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
134	Association Between Childhood Green Space, Genetic Liability, and the Incidence of Schizophrenia. Schizophrenia Bulletin, 2020, 46, 1629-1637.	2.3	28
135	Genetic risk scores for major psychiatric disorders and the risk of postpartum psychiatric disorders. Translational Psychiatry, 2019, 9, 288.	2.4	27
136	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. Biological Psychiatry, 2020, 87, 419-430.	0.7	27
137	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
138	Point mutation increases a form of the NK1receptor with high affinity for neurokinin A and B and septide. British Journal of Pharmacology, 1998, 125, 393-401.	2.7	26
139	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
140	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
141	Variation in the purinergic P2RX7 receptor gene and schizophrenia. Schizophrenia Research, 2008, 104, 146-152.	1.1	24
142	Association of MCTP2 gene variants with schizophrenia in three independent samples of Scandinavian origin (SCOPE). Psychiatry Research, 2009, 168, 256-258.	1.7	24
143	The association between candidate migraine susceptibility loci and severe migraine phenotype in a clinical sample. Cephalalgia, 2016, 36, 615-623.	1.8	24
144	Altered auditory processing and effective connectivity in 22q11.2 deletion syndrome. Schizophrenia Research, 2018, 197, 328-336.	1.1	24

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145	Elevated expression of a minor isoform of ANK3 is a risk factor for bipolar disorder. Translational Psychiatry, 2018, 8, 210.	2.4	24
146	Comorbidity of migraine with ADHD in adults. BMC Neurology, 2018, 18, 147.	0.8	24
147	Comparing Copy Number Variations in a Danish Case Cohort of Individuals With Psychiatric Disorders. JAMA Psychiatry, 2022, 79, 59.	6.0	24
148	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
149	Restless legs syndrome is associated with major comorbidities in a population of Danish blood donors. Sleep Medicine, 2018, 45, 124-131.	0.8	23
150	No significant association of the 5â€² end of neuregulin 1 and schizophrenia in a large Danish sample. Schizophrenia Research, 2006, 83, 1-5.	1.1	22
151	Association of a dopamine beta-hydroxylase gene variant with depression in elderly women possibly reflecting noradrenergic dysfunction. Journal of Affective Disorders, 2008, 106, 169-172.	2.0	22
152	An Empirical Bayes Mixture Model for Effect Size Distributions in Genome-Wide Association Studies. PLoS Genetics, 2015, 11, e1005717.	1.5	22
153	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
154	Roadmap for a precision-medicine initiative in the Nordic region. Nature Genetics, 2019, 51, 924-930.	9.4	22
155	Polygenic risk score, psychosocial environment and the risk of attention-deficit/hyperactivity disorder. Translational Psychiatry, 2020, 10, 335.	2.4	22
156	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
157	Intracellular single chain Fv antibody inhibits Ras activity in T-cell antigen receptor stimulated Jurkat cells. FEBS Letters, 1994, 351, 393-396.	1.3	21
158	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
159	FUT2â€“ABO epistasis increases the risk of early childhood asthma and Streptococcus pneumoniae respiratory illnesses. Nature Communications, 2020, 11, 6398.	5.8	21
160	Rare SLC13A1 variants associate with intervertebral disc disorder highlighting role of sulfate in disc pathology. Nature Communications, 2022, 13, 634.	5.8	21
161	A Mutation Changes Ligand Selectivity and Transmembrane Signaling Preference of the Neurokinin-1 Receptor. Journal of Biological Chemistry, 1997, 272, 7646-7655.	1.6	20
162	The Acetylcholinesterase Inhibitor Galantamine Inhibits d-Amphetamine-Induced Psychotic-Like Behavior in Cebus Monkeys. Journal of Pharmacology and Experimental Therapeutics, 2007, 321, 1179-1182.	1.3	20

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163	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
164	Combinations of SNPs Related to Signal Transduction in Bipolar Disorder. PLoS ONE, 2011, 6, e23812.	1.1	20
165	A Genetic Deconstruction of Neurocognitive Traits in Schizophrenia and Bipolar Disorder. PLoS ONE, 2013, 8, e81052.	1.1	20
166	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
167	Effects of the cannabinoid CB1 receptor agonist CP55,940 and antagonist SR141716A on d-amphetamine-induced behaviours in Cebusmonkeys. Journal of Psychopharmacology, 2006, 20, 622-628.	2.0	19
168	Novel procedure for genotyping of the human serotonin transporter gene-linked polymorphic region (5-HTTLPR) â€“ a region with a high level of allele diversity. Psychiatric Genetics, 2007, 17, 287-291.	0.6	19
169	Kynurenine 3-monooxygenase (KMO) polymorphisms in schizophrenia: An association study. Schizophrenia Research, 2011, 127, 270-272.	1.1	19
170	Individualization of treatments with drugs metabolized by CES1: combining genetics and metabolomics. Pharmacogenomics, 2015, 16, 649-665.	0.6	19
171	Schizophrenia polygenic risk scores, urbanicity and treatment-resistant schizophrenia. Schizophrenia Research, 2019, 212, 79-85.	1.1	19
172	Genetic stratification of depression in UK Biobank. Translational Psychiatry, 2020, 10, 163.	2.4	19
173	No association between DGKH and bipolar disorder in a Scandinavian caseâ€“control sample. Psychiatric Genetics, 2009, 19, 269-272.	0.6	18
174	Habitual sleep disturbances and migraine: a Mendelian randomization study. Annals of Clinical and Translational Neurology, 2020, 7, 2370-2380.	1.7	18
175	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
176	Soluble Urokinase-Type Plasminogen Activator Receptor Levels in Patients With Schizophrenia. Schizophrenia Bulletin, 2015, 41, 764-771.	2.3	16
177	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
178	Gestational age-dependent development of the neonatal metabolome. Pediatric Research, 2021, 89, 1396-1404.	1.1	16
179	Cohort Profile: COVIDMENT: COVID-19 cohorts on mental health across six nations. International Journal of Epidemiology, 2022, 51, e108-e122.	0.9	16
180	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

#	Article	IF	Citations
181	Analysis of coding-polymorphisms in NOTCH-related genes reveals NUMBL poly-glutamine repeat to be associated with schizophrenia in Brazilian and Danish subjects. Schizophrenia Research, 2006, 88, 275-282.	1.1	15
182	Replication Study and Meta-Analysis in European Samples Supports Association of the 3p21.1 Locus with Bipolar Disorder. Biological Psychiatry, 2012, 72, 645-650.	0.7	15
183	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
184	Genetic predictors of educational attainment and intelligence test performance predict voter turnout. Nature Human Behaviour, 2021, 5, 281-291.	6.2	15
185	Genetic, Clinical, and Sociodemographic Factors Associated With Stimulant Treatment Outcomes in ADHD. American Journal of Psychiatry, 2021, 178, 854-864.	4.0	15
186	Effects of postnatal anoxia on striatal dopamine metabolism and prepulse inhibition in rats. Pharmacology Biochemistry and Behavior, 2004, 77, 767-774.	1.3	14
187	The Danish 22q11 research initiative. BMC Psychiatry, 2015, 15, 220.	1.1	14
188	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
189	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
190	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
191	Diagnostic stability among chronic patients with functional psychoses: an epidemiological and clinical study. BMC Psychiatry, 2007, 7, 41.	1.1	13
192	Tryptophan hydroxylase gene 1 (TPH1) variants associated with cerebrospinal fluid 5-hydroxyindole acetic acid and homovanillic acid concentrations in healthy volunteers. Psychiatry Research, 2010, 180, 63-67.	1.7	13
193	PCM1 is necessary for focal ciliary integrity and is a candidate for severe schizophrenia. Nature Communications, 2020, 11, 5903.	5.8	13
194	Pharmacogenetic genotype and phenotype frequencies in a large Danish population-based case-cohort sample. Translational Psychiatry, 2021, 11, 294.	2.4	13
195	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
196	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
197	Post-traumatic stress following military deployment: Genetic associations and cross-disorder genetic correlations. Journal of Affective Disorders, 2019, 252, 350-357.	2.0	12
198	No evidence of associations between genetic liability for schizophrenia and development of cannabis use disorder. Psychological Medicine, 2021, 51, 479-484.	2.7	12

#	Article	IF	Citations
199	Translating polygenic risk scores for clinical use by estimating the confidence bounds of risk prediction. Nature Communications, 2021, 12, 5276.	5.8	12
200	Genome-wide association study identifies 16 genomic regions associated with circulating cytokines at birth. PLoS Genetics, 2020, 16, e1009163.	1.5	12
201	The gene encoding the melanin-concentrating hormone receptor 1 is associated with schizophrenia in a Danish caseâ€“control sample. Psychiatric Genetics, 2012, 22, 62-69.	0.6	11
202	d-amino acid oxidase activator gene (DAOA) variation affects cerebrospinal fluid homovanillic acid concentrations in healthy Caucasians. European Archives of Psychiatry and Clinical Neuroscience, 2012, 262, 549-556.	1.8	11
203	The Duffy-null genotype and risk of infection. Human Molecular Genetics, 2020, 29, 3341-3349.	1.4	11
204	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
205	Identifying dominant-negative actions of a dopamine transporter variant in patients with parkinsonism and neuropsychiatric disease. JCI Insight, 2021, 6, .	2.3	11
206	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
207	Mutation screening of the glutamate cysteine ligase modifier (GCLM) gene in patients with schizophrenia. Psychiatric Genetics, 2009, 19, 201-208.	0.6	10
208	No association of polymorphisms in human endogenous retrovirus K18 and CD48 with schizophrenia. Psychiatric Genetics, 2012, 22, 146-148.	0.6	10
209	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
210	The genetic structure of Norway. European Journal of Human Genetics, 2021, 29, 1710-1718.	1.4	10
211	Common Genetic Variation and Age of Onset of Anorexia Nervosa. Biological Psychiatry Global Open Science, 2022, 2, 368-378.	1.0	10
212	Genome-Wide Association Study of Genetic Variants in LPS-Stimulated IL-6, IL-8, IL-10, IL-1ra and TNF-Î± Cytokine Response in a Danish Cohort. PLoS ONE, 2013, 8, e66262.	1.1	10
213	Identifying a putative common binding site shared by substance P receptor and an anti-substance P monoclonal antibody. Protein Engineering, Design and Selection, 1995, 8, 403-408.	1.0	9
214	Does the Medication Pattern Reflect the CYP2D6 Genotype in Patients With Diagnoses Within the Schizophrenic Spectrum?. Journal of Clinical Psychopharmacology, 2012, 32, 100-105.	0.7	9
215	Lack of association between the regulator of G-protein signaling 4 (RGS4) rs951436 polymorphism and schizophrenia. Psychiatric Genetics, 2012, 22, 263-264.	0.6	9
216	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

#	Article	IF	Citations
217	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
218	Anorexia nervosa and inflammatory bowel diseasesâ€"Diagnostic and genetic associations. JCPP Advances, 2021, 1, e12036.	1.4	9
219	Cognitive impairment in elderly women: the relative importance of selected genes, lifestyle factors, and comorbidities. Neuropsychiatric Disease and Treatment, 2006, 2, 227-233.	1.0	9
220	Associations between patterns in comorbid diagnostic trajectories of individuals with schizophrenia and etiological factors. Nature Communications, 2021, 12, 6617.	5.8	9
221	Comprehensive genome-wide association study of different forms of hernia identifies more than 80 associated loci. Nature Communications, 2022, 13, .	5.8	9
222	Gly166in the NK1receptor regulates tachykinin selectivity and receptor conformation1. FEBS Letters, 1997, 416, 335-338.	1.3	8
223	An evaluation of the genetic-matched pair study design using genome-wide SNP data from the European population. European Journal of Human Genetics, 2009, 17, 967-975.	1.4	8
224	Novel variant of CYP2D6*6 is undetected by a commonly used genotyping procedure. Pharmacological Reports, 2011, 63, 1264-1266.	1.5	8
225	Dual association of a TRKA polymorphism with schizophrenia. Psychiatric Genetics, 2011, 21, 125-131.	0.6	8
226	Interactive effects between hemizygous 15q13.3 microdeletion and peripubertal stress on adult behavioral functions. Neuropsychopharmacology, 2019, 44, 703-710.	2.8	8
227	Association between GLP-1 receptor gene polymorphisms with reward learning, anhedonia and depression diagnosis. Acta Neuropsychiatrica, 2020, 32, 218-225.	1.0	8
228	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
229	Family disadvantage, gender, and the returns to genetic human capital*. Scandinavian Journal of Economics, 2022, 124, 550-578.	0.7	8
230	A genome-wide meta-analysis identifies 50 genetic loci associated with carpal tunnel syndrome. Nature Communications, 2022, 13, 1598.	5.8	8
231	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
232	The Impact of CYP2D6 and CYP2C19 Polymorphisms on Suicidal Behavior and Substance Abuse Disorder Among Patients With Schizophrenia: A Retrospective Study. Therapeutic Drug Monitoring, 2008, 30, 265-270.	1.0	7
233	Quantitative genome-wide association analyses of receptive language in the Danish High Risk and Resilience Study. BMC Neuroscience, 2020, 21, 30.	0.8	7
234	Polygenic Heterogeneity Across Obsessive-Compulsive Disorder Subgroups Defined by a Comorbid Diagnosis. Frontiers in Genetics, 2021, 12, 711624.	1.1	7

#	Article	IF	Citations
235	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
236	Identification and Characterization of a Tandem Repeat in Exon III of the Dopamine Receptor D4 (DRD4) Gene in Cetaceans. Journal of Heredity, 2006, 97, 279-284.	1.0	6
237	Effects of cannabinoid CB1 receptor agonism and antagonism on SKF81297-induced dyskinesia and haloperidol-induced dystonia in Cebus apella monkeys. Neuropharmacology, 2011, 60, 418-422.	2.0	6
238	Deep learningâ€"based integration of genetics with registry data for stratification of schizophrenia and depression. Science Advances, 2022, 8, .	4.7	6
239	Association of the 120-bp duplication in the dopamine D4 receptor gene and schizophrenia in a sample of Danish subjects. Schizophrenia Research, 2005, 73, 133-135.	1.1	5
240	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
241	Complex spatio-temporal distribution and genomic ancestry of mitochondrial DNA haplogroups in 24,216 Danes. PLoS ONE, 2018, 13, e0208829.	1.1	5
242	Genetic liability to major depression and risk of childhood asthma. Brain, Behavior, and Immunity, 2020, 89, 433-439.	2.0	5
243	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
244	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
245	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
246	Response to Boot et al. Letter. American Journal of Psychiatry, 2012, 169, 97-97.	4.0	4
247	Association of the leucine-7 to proline-7 variation in the signal sequence of neuropeptide Y with major depression. Acta Neuropsychiatrica, 2012, 24, 81-90.	1.0	4
248	Combinations of Genetic Data Present in Bipolar Patients, but Absent in Control Persons. PLoS ONE, 2015, 10, e0143432.	1.1	4
249	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
250	Misclassification of Allele CYP2C19*10 as CY2C19*2 by a Commonly Used PCR-RFLP Procedure. Genetic Testing and Molecular Biomarkers, 2008, 12, 57-58.	1.7	3
251	Tyrosine hydroxylase Val81Met polymorphism: lack of association with schizophrenia. Psychiatric Genetics, 2009, 19, 273-274.	0.6	3
252	Lack of association between two dopamine D2 receptor gene polymorphisms and schizophrenia. Psychiatric Genetics, 2011, 21, 214-215.	0.6	3

#	Article	IF	Citations
253	Copy number variations in affective disorders and meta-analysis. Psychiatric Genetics, 2011, 21, 319-322.	0.6	3
254	Clinical association to FKBP5 rs1360780 in patients with depression. Psychiatric Genetics, 2019, 29, 220-225.	0.6	3
255	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
256	A closed-tube assay for genotyping of the 32-bp deletion polymorphism in the chemokine receptor 5 (CCR5) gene: Dissociation analysis of amplified fragments of DNA. Molecular and Cellular Probes, 2007, 21, 8-11.	0.9	2
257	Dystrobrevin-binding protein 1 gene (DTNBP1) variants associated with cerebrospinal fluid homovanillic acid and 5-hydroxyindoleacetic acid concentrations in healthy volunteers. European Neuropsychopharmacology, 2011, 21, 700-704.	0.3	2
258	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
259	Lessons to be Learned From 22q2.11 Syndromesâ€”Reply. JAMA Psychiatry, 2017, 74, 757.	6.0	2
260	Antipsychotic-Like Effect of the Muscarinic Acetylcholine Receptor Agonist BuTAC in Non-Human Primates. PLoS ONE, 2015, 10, e0122722.	1.1	2
261	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
262	Neuropeptide Y genes and suicidal behaviour among schizophrenic patients. Psychiatric Genetics, 2013, 23, 139-140.	0.6	1
263	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
264	Multi-State Survival Analysis Of Multi-Psychiatry Disorders. European Neuropsychopharmacology, 2017, 27, S368-S369.	0.3	0
265	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
266	A CROSS-MENTAL-DISORDERS GWAS IN THE DANISH NATIONAL HEALTH REGISTER. European Neuropsychopharmacology, 2019, 29, S996.	0.3	0
267	9IDENTICAL BY DESCENT SEGMENTS ASSOCIATES WITH SCHIZOPHRENIA. European Neuropsychopharmacology, 2019, 29, S1071.	0.3	0
268	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
269	GENOME-WIDE METHYLOMIC ANALYSIS OF NEONATAL BLOOD FROM DANISH TWINS DISCORDANT FOR MENTAL ILLNESS. European Neuropsychopharmacology, 2019, 29, S794-S795.	0.3	0
270	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

#	Article	IF	Citations
271	THE IMPACT OF CNVs ON ASD/ADHD RISK IN MULTIPLEX FAMILIES. European Neuropsychopharmacology, 2019, 29, S895.	0.3	0
272	SA133GENETIC PREDICTION OF TRANSITION FROM CHILDHOOD AND ADOLESCENT PSYCHIATRIC DISORDER TO SCHIZOPHRENIA. European Neuropsychopharmacology, 2019, 29, S1262.	0.3	0
273	Accumulation of disadvantages across multiple domains amongst subgroups of children of parents with schizophrenia or bipolar disorder. Clustering data from the Danish High Risk and Resilience Study VIA 7. Schizophrenia Bulletin Open, 0, , .	0.9	0
274	Title is missing!. , 2020, 16, e1009163.		0
275	Title is missing!. , 2020, 16, e1009163.		0
276	Title is missing!. , 2020, 16, e1009163.		0
277	Title is missing!. , 2020, 16, e1009163.		0
278	School performance and genetic propensities for educational attainment and depression in the etiology of self-harm: a Danish population-based study. Nordic Journal of Psychiatry, 0, , 1-9.	0.7	0
279	Genetic liability to posttraumatic stress disorder and its association with postpartum depression. Psychological Medicine, 0, , 1-8.	2.7	0