Anders BÃ¸rglum

List of Publications by Year
in descending order

148 papers	22,513 citations	38720 50 h-index	13365 130 g-index
197 all docs	197 docs citations	197 times ranked	22518 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	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. Biological Psychiatry, 2022, 91, 313-327.	0.7	114
4	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
5	Family disadvantage, gender, and the returns to genetic human capital*. Scandinavian Journal of Economics, 2022, 124, 550-578.	0.7	8
6	Comparing Copy Number Variations in a Danish Case Cohort of Individuals With Psychiatric Disorders. JAMA Psychiatry, 2022, 79, 59.	6.0	24
7	Early-Life Injuries and the Development of Attention-Deficit/Hyperactivity Disorder. Journal of Clinical Psychiatry, 2022, 83, .	1.1	0
8	Seasonal variation and risk of febrile seizures; a Danish nationwide cohort study. Neuroepidemiology, 2022, , .	1.1	2
9	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
10	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
11	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	13.7	929
12	Genetic architecture of 11 major psychiatric disorders at biobehavioral, functional genomic and molecular genetic levels of analysis. Nature Genetics, 2022, 54, 548-559.	9.4	101
13	Genetic correlates of phenotypic heterogeneity in autism. Nature Genetics, 2022, 54, 1293-1304.	9.4	51
14	The female protective effect against autism spectrum disorder. Cell Genomics, 2022, 2, 100134.	3.0	30
15	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
16	Genetic predictors of educational attainment and intelligence test performance predict voter turnout. Nature Human Behaviour, 2021, 5, 281-291.	6.2	15
17	Subdividing Y-chromosome haplogroup R1a1 reveals Norse Viking dispersal lineages in Britain. European Journal of Human Genetics, 2021, 29, 512-523.	1.4	9
18	A polygenic resilience score moderates the genetic risk for schizophrenia. Molecular Psychiatry, 2021, 26, 800-815.	4.1	36

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19	No evidence of associations between genetic liability for schizophrenia and development of cannabis use disorder. Psychological Medicine, 2021, 51, 479-484.	2.7	12
20	Investigating causality between liability to ADHD and substance use, and liability to substance use and ADHD risk, using Mendelian randomization. Addiction Biology, 2021, 26, e12849.	1.4	52
21	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
22	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
23	Risk of Early-Onset Depression Associated With Polygenic Liability, Parental Psychiatric History, and Socioeconomic Status. JAMA Psychiatry, 2021, 78, 387.	6.0	33
24	Genetic analyses identify widespread sex-differential participation bias. Nature Genetics, 2021, 53, 663-671.	9.4	124
25	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
26	Pharmacogenetic genotype and phenotype frequencies in a large Danish population-based case-cohort sample. Translational Psychiatry, 2021, 11, 294.	2.4	13
27	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
28	Examining Sex-Differentiated Genetic Effects Across Neuropsychiatric and Behavioral Traits. Biological Psychiatry, 2021, 89, 1127-1137.	0.7	48
29	Polygenic Heterogeneity Across Obsessive-Compulsive Disorder Subgroups Defined by a Comorbid Diagnosis. Frontiers in Genetics, 2021, 12, 711624.	1.1	7
30	Translating polygenic risk scores for clinical use by estimating the confidence bounds of risk prediction. Nature Communications, 2021, 12, 5276.	5.8	12
31	Investigating Shared Genetic Basis Across Tourette Syndrome and Comorbid Neurodevelopmental Disorders Along the Impulsivity-Compulsivity Spectrum. Biological Psychiatry, 2021, 90, 317-327.	0.7	49
32	Genetic, Clinical, and Sociodemographic Factors Associated With Stimulant Treatment Outcomes in ADHD. American Journal of Psychiatry, 2021, 178, 854-864.	4.0	15
33	DNA methylation of the KLK8 gene in depression symptomatology. Clinical Epigenetics, 2021, 13, 200.	1.8	7
34	Discordant associations of educational attainment with ASD and ADHD implicate a polygenic form of pleiotropy. Nature Communications, 2021, 12, 6534.	5.8	3
35	Associations between patterns in comorbid diagnostic trajectories of individuals with schizophrenia and etiological factors. Nature Communications, 2021, 12, 6617.	5.8	9
36	Attention-deficit/hyperactivity disorder and lifetime cannabis use: genetic overlap and causality. Molecular Psychiatry, 2020, 25, 2493-2503.	4.1	59

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37	A major role for common genetic variation in anxiety disorders. Molecular Psychiatry, 2020, 25, 3292-3303.	4.1	243
38	Molecular genetic overlap between posttraumatic stress disorder and sleep phenotypes. Sleep, 2020, 43, .	0.6	32
39	Genetic liability to ADHD and substance use disorders in individuals with ADHD. Addiction, 2020, 115, 1368-1377.	1.7	47
40	Polygenic risk score, psychosocial environment and the risk of attention-deficit/hyperactivity disorder. Translational Psychiatry, 2020, 10, 335.	2.4	22
41	A large-scale genome-wide association study meta-analysis of cannabis use disorder. Lancet Psychiatry,the, 2020, 7, 1032-1045.	3.7	200
42	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
43	The genetic architecture of sporadic and multiple consecutive miscarriage. Nature Communications, 2020, 11, 5980.	5.8	52
44	Genetic liability to major depression and risk of childhood asthma. Brain, Behavior, and Immunity, 2020, 89, 433-439.	2.0	5
45	Reduced Brd1 expression leads to reversible depression-like behaviors and gene-expression changes in female mice. Translational Psychiatry, 2020, 10, 239.	2.4	8
46	Common schizophrenia risk variants are enriched in open chromatin regions of human glutamatergic neurons. Nature Communications, 2020, 11, 5581.	5.8	53
47	Neuropsin in mental health. Journal of Physiological Sciences, 2020, 70, 26.	0.9	7
48	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
49	Contribution of Intellectual Disabilityâ€“Related Genes to ADHD Risk and to Locomotor Activity in Drosophila. American Journal of Psychiatry, 2020, 177, 526-536.	4.0	22
50	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
51	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
52	Association between Mental Disorders and Subsequent Medical Conditions. New England Journal of Medicine, 2020, 382, 1721-1731.	13.9	258
53	Shared genetic background between children and adults with attention deficit/hyperactivity disorder. Neuropsychopharmacology, 2020, 45, 1617-1626.	2.8	72
54	FUT2â€“ABO epistasis increases the risk of early childhood asthma and Streptococcus pneumoniae respiratory illnesses. Nature Communications, 2020, 11, 6398.	5.8	21

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55	Genome-wide meta-analysis of problematic alcohol use in 435,563 individuals yields insights into biology and relationships with other traits. Nature Neuroscience, 2020, 23, 809-818.	7.1	242
56	Genomic analysis of the natural history of attention-deficit/hyperactivity disorder using Neanderthal and ancient Homo sapiens samples. Scientific Reports, 2020, 10, 8622.	1.6	18
57	Investigating the association between body fat and depression via Mendelian randomization. Translational Psychiatry, 2019, 9, 184.	2.4	90
58	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
59	Reduced neonatal brain-derived neurotrophic factor is associated with autism spectrum disorders. Translational Psychiatry, 2019, 9, 252.	2.4	56
60	Genetic risk scores for major psychiatric disorders and the risk of postpartum psychiatric disorders. Translational Psychiatry, 2019, 9, 288.	2.4	27
61	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
62	Epigenome-wide association study of depression symptomatology in elderly monozygotic twins. Translational Psychiatry, 2019, 9, 214.	2.4	48
63	Social and non-social autism symptoms and trait domains are genetically dissociable. Communications Biology, 2019, 2, 328.	2.0	57
64	Variants in the fetal genome near pro-inflammatory cytokine genes on 2q13 associate with gestational duration. Nature Communications, 2019, 10, 3927.	5.8	49
65	Schizophrenia polygenic risk scores, urbanicity and treatment-resistant schizophrenia. Schizophrenia Research, 2019, 212, 79-85.	1.1	19
66	Exploring genetic variation that influences brain methylation in attention-deficit/hyperactivity disorder. Translational Psychiatry, 2019, 9, 242.	2.4	21
67	Modeling the cooperativity of schizophrenia risk genes. Nature Genetics, 2019, 51, 1434-1436.	9.4	3
68	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
69	Disentangling polygenic associations between attention-deficit/hyperactivity disorder, educational attainment, literacy and language. Translational Psychiatry, 2019, 9, 35.	2.4	25
70	Genome-wide association study implicates CHRNA2 in cannabis use disorder. Nature Neuroscience, 2019, 22, 1066-1074.	7.1	94
71	Genetic Variants Associated With Anxiety and Stress-Related Disorders. JAMA Psychiatry, 2019, 76, 924.	6.0	140
72	Genome-wide association study identifies 30 loci associated with bipolar disorder. Nature Genetics, 2019, 51, 793-803.	9.4	1,191

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73	Genetic Markers of ADHD-Related Variations in Intracranial Volume. American Journal of Psychiatry, 2019, 176, 228-238.	4.0	68
74	Gene expression imputation across multiple brain regions provides insights into schizophrenia risk. Nature Genetics, 2019, 51, 659-674.	9.4	154
75	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
76	Roadmap for a precision-medicine initiative in the Nordic region. Nature Genetics, 2019, 51, 924-930.	9.4	22
77	Identification of common genetic risk variants for autism spectrum disorder. Nature Genetics, 2019, 51, 431-444.	9.4	1,538
78	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
79	Genetic influences on eight psychiatric disorders based on family data of 4 408 646 full and half-siblings, and genetic data of 333 748 cases and controls. Psychological Medicine, 2019, 49, 1166-1173.	2.7	106
80	Discovery of the first genome-wide significant risk loci for attention deficit/hyperactivity disorder. Nature Genetics, 2019, 51, 63-75.	9.4	1,594
81	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
82	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
83	Genome-wide analyses of self-reported empathy: correlations with autism, schizophrenia, and anorexia nervosa. Translational Psychiatry, 2018, 8, 35.	2.4	95
84	ASD and schizophrenia show distinct developmental profiles in common genetic overlap with population-based social communication difficulties. Molecular Psychiatry, 2018, 23, 263-270.	4.1	107
85	Psychiatric Genomics: An Update and an Agenda. American Journal of Psychiatry, 2018, 175, 15-27.	4.0	518
86	A Genetic Investigation of Sex Bias in the Prevalence of Attention-Deficit/Hyperactivity Disorder. Biological Psychiatry, 2018, 83, 1044-1053.	0.7	146
87	Brain volumetric alterations accompanied with loss of striatal medium-sized spiny neurons and cortical parvalbumin expressing interneurons in Brd1+/â^' mice. Scientific Reports, 2018, 8, 16486.	1.6	14
88	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
89	Complex spatio-temporal distribution and genomic ancestry of mitochondrial DNA haplogroups in 24,216 Danes. PLoS ONE, 2018, 13, e0208829.	1.1	5
90	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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91	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	6.0	1,085
92	Elevated polygenic burden for autism is associated with differential DNA methylation at birth. Genome Medicine, 2018, 10, 19.	3.6	88
93	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
94	The iPSYCH2012 caseâ€“cohort sample: new directions for unravelling genetic and environmental architectures of severe mental disorders. Molecular Psychiatry, 2018, 23, 6-14.	4.1	257
95	Genome-wide Association for Major Depression Through Age at Onset Stratification: Major Depressive Disorder Working Group of the Psychiatric Genomics Consortium. Biological Psychiatry, 2017, 81, 325-335.	0.7	175
96	Blood DNA methylation age is not associated with cognitive functioning in middle-aged monozygotic twins. Neurobiology of Aging, 2017, 50, 60-63.	1.5	28
97	Endogenous and Antipsychotic-Related Risks for Diabetes Mellitus in Young People With Schizophrenia: A Danish Population-Based Cohort Study. American Journal of Psychiatry, 2017, 174, 686-694.	4.0	103
98	Mice heterozygous for an inactivated allele of the schizophrenia associated Brd1 gene display selective cognitive deficits with translational relevance to schizophrenia. Neurobiology of Learning and Memory, 2017, 141, 44-52.	1.0	16
99	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
100	Association of the polygenic risk score for schizophrenia with mortality and suicidal behavior - A Danish population-based study. Schizophrenia Research, 2017, 184, 122-127.	1.1	27
101	Epigenome-Wide Association Study of Cognitive Functioning in Middle-Aged Monozygotic Twins. Frontiers in Aging Neuroscience, 2017, 9, 413.	1.7	52
102	Sequencing and de novo assembly of 150 genomes from Denmark as a population reference. Nature, 2017, 548, 87-91.	13.7	130
103	DNA Methylation Analysis of BRD1 Promoter Regions and the Schizophrenia rs138880 Risk Allele. PLoS ONE, 2017, 12, e0170121.	1.1	14
104	Influence of Polygenic Risk Scores on the Association Between Infections and Schizophrenia. Biological Psychiatry, 2016, 80, 609-616.	0.7	38
105	Genome-wide DNA methylation profiling with MeDIP-seq using archived dried blood spots. Clinical Epigenetics, 2016, 8, 81.	1.8	36
106	Identification of the BRD1 interaction network and its impact on mental disorder risk. Genome Medicine, 2016, 8, 53.	3.6	29
107	Genetic risk for autism spectrum disorders and neuropsychiatric variation in the general population. Nature Genetics, 2016, 48, 552-555.	9.4	326
108	High loading of polygenic risk in cases with chronic schizophrenia. Molecular Psychiatry, 2016, 21, 969-974.	4.1	62

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109	EWS and FUS bind a subset of transcribed genes encoding proteins enriched in RNA regulatory functions. BMC Genomics, 2015, 16, 929.	1.2	21
110	Polygenic Risk Score, Parental Socioeconomic Status, Family History of Psychiatric Disorders, and the Risk for Schizophrenia. JAMA Psychiatry, 2015, 72, 635.	6.0	242
111	Systematic Integration of Brain eQTL and GWAS IdentifiesZNF323as a Novel Schizophrenia Risk Gene and Suggests Recent Positive Selection Based on Compensatory Advantage on Pulmonary Function. Schizophrenia Bulletin, 2015, 41, 1294-1308.	2.3	48
112	Experimental validation of methods for differential gene expression analysis and sample pooling in RNA-seq. BMC Genomics, 2015, 16, 548.	1.2	139
113	Genome-wide study of association and interaction with maternal cytomegalovirus infection suggests new schizophrenia loci. Molecular Psychiatry, 2014, 19, 325-333.	4.1	163
114	GWAS, Cytomegalovirus Infection, and Schizophrenia. Current Behavioral Neuroscience Reports, 2014, 1, 215-223.	0.6	9
115	Cuba: Exploring the History of Admixture and the Genetic Basis of Pigmentation Using Autosomal and Uniparental Markers. PLoS Genetics, 2014, 10, e1004488.	1.5	57
116	Investigation of the involvement of MIR185 and its target genes in the development of schizophrenia. Journal of Psychiatry and Neuroscience, 2014, 39, 386-396.	1.4	23
117	Integrated Pathway-Based Approach Identifies Association between Genomic Regions at CTCF and CACNB2 and Schizophrenia. PLoS Genetics, 2014, 10, e1004345.	1.5	44
118	Common variant at 16p11.2 conferring risk of psychosis. Molecular Psychiatry, 2014, 19, 108-114.	4.1	85
119	The Schizophrenia and Bipolar Disorder associated BRD1 gene is regulated upon chronic restraint stress. European Neuropsychopharmacology, 2012, 22, 651-656.	0.3	22
120	Electroconvulsive seizures regulates the Brd1 gene in the frontal cortex and hippocampus of the adult rat. Neuroscience Letters, 2012, 516, 110-113.	1.0	17
121	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
122	Robustness of genome-wide scanning using archived dried blood spot samples as a DNA source. BMC Genetics, 2011, 12, 58.	2.7	79
123	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
124	Support of association between BRD1 and both schizophrenia and bipolar affective disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 582-591.	1.1	47
125	Genome-wide scans using archived neonatal dried blood spot samples. BMC Genomics, 2009, 10, 297.	1.2	80
126	Further immunohistochemical characterization of BRD1 a new susceptibility gene for schizophrenia and bipolar affective disorder. Brain Structure and Function, 2009, 214, 37-47.	1.2	14

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127	Common variants conferring risk of schizophrenia. Nature, 2009, 460, 744-747.	13.7	1,572
128	No signature of Y chromosomal resemblance between possible descendants of the Cimbri in Denmark and Northern Italy. American Journal of Physical Anthropology, 2007, 132, 278-284.	2.1	2
129	Evidence implicating BRD1 with brain development and susceptibility to both schizophrenia and bipolar affective disorder. Molecular Psychiatry, 2006, 11, 1126-1138.	4.1	77
130	The origin of the isolated population of the Faroe Islands investigated using Y chromosomal markers. Human Genetics, 2004, 115, 19-28.	1.8	43
131	Genomic structure of the human mitochondrial chaperonin genes: HSP60 and HSP10 are localised head to head on chromosomeÂ2 separated by a bidirectional promoter. Human Genetics, 2003, 112, 436-436.	1.8	1
132	Possible parent-of-origin effect of Dopa decarboxylase in susceptibility to bipolar affective disorder. , 2003, 117B, 18-22.		39
133	Search for common haplotypes on chromosome 22q in patients with schizophrenia or bipolar disorder from the Faroe Islands. American Journal of Medical Genetics Part A, 2002, 114, 245-252.	2.4	40
134	A new locus for Seckel syndrome on chromosome 18p11.31-q11.2. European Journal of Human Genetics, 2001, 9, 753-757.	1.4	54
135	Dopa decarboxylase genotypes may influence age at onset of schizophrenia. Molecular Psychiatry, 2001, 6, 712-717.	4.1	31
136	Human and mouse mitochondrial orthologs of bacterial ClpX. Mammalian Genome, 2000, 11, 899-905.	1.0	36
137	Two novel variants in the DOPA decarboxylase gene: association with bipolar affective disorder. Molecular Psychiatry, 1999, 4, 545-551.	4.1	49
138	Mapping of 34 minisatellite loci resolved by two-dimensional DNA typing. Cytogenetic and Genome Research, 1997, 79, 248-256.	0.6	0
139	Refined localization of the pyruvate dehydrogenase E1Î± gene (PDHA1) by linkage analysis. Human Genetics, 1996, 99, 80-82.	1.8	6
140	Assignment of the human tryptophanyl-tRNA synthetase gene (WARS) to chromosome 14q32.2→q32.32. Cytogenetic and Genome Research, 1996, 73, 99-103.	0.6	5
141	Refined mapping of the psoriasin gene S100A7 to chromosome 1cen-q21. Human Genetics, 1995, 96, 592-6.	1.8	14
142	Two-dimensional DNA typing as a genetic marker system in humans. Cytogenetic and Genome Research, 1995, 71, 260-265.	0.6	0
143	Charcotâ€Marieâ€Tooth disease type 1A: the parental origin of a de novo 17p11.2â€p12 duplication. Clinical Genetics, 1994, 46, 291-294.	1.0	15
144	Two PstI polymorphisms for the urokinase-type plasminogen activator receptor gene (PLAUR). Human Genetics, 1992, 89, 584.	1.8	2

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145	Assignment of the urokinase-type plasminogen activator receptor gene (PLAUR) to chromosome 19q13.1-q13.2. American Journal of Human Genetics, 1992, 50, 492-7.	2.6	36
146	An EcoRI polymorphism for the PLAUR gene. Nucleic Acids Research, 1991, 19, 6661-6661.	6.5	5
147	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
148	Genetic liability to posttraumatic stress disorder and its association with postpartum depression. Psychological Medicine, 0, , 1-8.	2.7	0