David M Hougaard

List of Publications by Year in descending order

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154 papers 27,253 citations

26610 56 h-index 146 g-index

197 all docs

197 docs citations

197 times ranked

28564 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	Discovery of the first genome-wide significant risk loci for attention deficit/hyperactivity disorder. Nature Genetics, 2019, 51, 63-75.	9.4	1,594
3	Common variants conferring risk of schizophrenia. Nature, 2009, 460, 744-747.	13.7	1,572
4	Identification of common genetic risk variants for autism spectrum disorder. Nature Genetics, 2019, 51, 431-444.	9.4	1,538
5	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
6	Genome-wide association analysis identifies 13 new risk loci for schizophrenia. Nature Genetics, 2013, 45, 1150-1159.	9.4	1,395
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	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
14	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. Cell, 2018, 173, 1705-1715.e16.	13.5	623
15	A genome-wide association study identifies CDHR3 as a susceptibility locus for early childhood asthma with severe exacerbations. Nature Genetics, 2014, 46, 51-55.	9.4	497
16	Genome-wide associations for birth weight and correlations with adult disease. Nature, 2016, 538, 248-252.	13.7	406
17	Maternal and fetal genetic effects on birth weight and their relevance to cardio-metabolic risk factors. Nature Genetics, 2019, 51, 804-814.	9.4	402
18	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

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19	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
20	Genetic risk for autism spectrum disorders and neuropsychiatric variation in the general population. Nature Genetics, 2016, 48, 552-555.	9.4	326
21	Neonatal Vitamin D Status and Risk of Schizophrenia. Archives of General Psychiatry, 2010, 67, 889.	13.8	315
22	New loci associated with birth weight identify genetic links between intrauterine growth and adult height and metabolism. Nature Genetics, 2013, 45, 76-82.	9.4	293
23	Simultaneous Measurement of 25 Inflammatory Markers and Neurotrophins in Neonatal Dried Blood Spots by Immunoassay with xMAP Technology. Clinical Chemistry, 2005, 51, 1854-1866.	1.5	252
24	A major role for common genetic variation in anxiety disorders. Molecular Psychiatry, 2020, 25, 3292-3303.	4.1	243
25	Polygenic Risk Score, Parental Socioeconomic Status, Family History of Psychiatric Disorders, and the Risk for Schizophrenia. JAMA Psychiatry, 2015, 72, 635.	6.0	242
26	Danish premature birth rates during the COVID-19 lockdown. Archives of Disease in Childhood: Fetal and Neonatal Edition, 2021, 106, 93-95.	1.4	223
27	A large-scale genome-wide association study meta-analysis of cannabis use disorder. Lancet Psychiatry,the, 2020, 7, 1032-1045.	3.7	200
28	An epigenetic clock for gestational age at birth based on blood methylation data. Genome Biology, 2016, 17, 206.	3.8	193
29	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
30	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
31	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
32	A Genetic Investigation of Sex Bias in the Prevalence of Attention-Deficit/Hyperactivity Disorder. Biological Psychiatry, 2018, 83, 1044-1053.	0.7	146
33	Genetic Variants Associated With Anxiety and Stress-Related Disorders. JAMA Psychiatry, 2019, 76, 924.	6.0	140
34	Foetal oestrogens and autism. Molecular Psychiatry, 2020, 25, 2970-2978.	4.1	132
35	Common variants associated with general and MMR vaccine–related febrile seizures. Nature Genetics, 2014, 46, 1274-1282.	9.4	128
36	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

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37	Cytokines associated with necrotizing enterocolitis in extremely-low-birth-weight infants. Pediatric Research, 2014, 76, 100-108.	1.1	120
38	Genome-wide meta-analysis associates HLA-DQA1/DRB1 and LPA and lifestyle factors with human longevity. Nature Communications, 2017, 8, 910.	5.8	118
39	Neonatal Screening in Europe Revisited: An ISNS Perspective on the Current State and Developments Since 2010. International Journal of Neonatal Screening, 2021, 7, 15.	1.2	118
40	Genome-wide gene-environment analyses of major depressive disorder and reported lifetime traumatic experiences in UK Biobank. Molecular Psychiatry, 2020, 25, 1430-1446.	4.1	116
41	Effects of blood sample handling procedures on measurable inflammatory markers in plasma, serum and dried blood spot samples. Journal of Immunological Methods, 2008, 336, 78-84.	0.6	115
42	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. Biological Psychiatry, 2022, 91, 313-327.	0.7	114
43	A Comparison of Ten Polygenic Score Methods for Psychiatric Disorders Applied Across Multiple Cohorts. Biological Psychiatry, 2021, 90, 611-620.	0.7	103
44	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
45	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
46	Genetic predisposition to hypertension is associated with preeclampsia in European and Central Asian women. Nature Communications, 2020, 11, 5976.	5.8	102
47	Genome-wide association analyses identify variants in developmental genes associated with hypospadias. Nature Genetics, 2014, 46, 957-963.	9.4	97
48	Genome-wide association study implicates CHRNA2 in cannabis use disorder. Nature Neuroscience, 2019, 22, 1066-1074.	7.1	94
49	Elevated polygenic burden for autism is associated with differential DNA methylation at birth. Genome Medicine, 2018, 10, 19.	3.6	88
50	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
51	Genome-wide scans using archived neonatal dried blood spot samples. BMC Genomics, 2009, 10, 297.	1.2	80
52	Robustness of genome-wide scanning using archived dried blood spot samples as a DNA source. BMC Genetics, 2011, 12, 58.	2.7	79
53	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
54	The association between neonatal vitamin D status and risk of schizophrenia. Scientific Reports, 2018, 8, 17692.	1.6	73

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55	The utility of neonatal dried blood spots for the assessment of neonatal vitamin D status. Paediatric and Perinatal Epidemiology, 2010, 24, 303-308.	0.8	69
56	Resveratrol reduces the levels of circulating androgen precursors but has no effect on, testosterone, dihydrotestosterone, PSA levels or prostate volume. A 4-month randomised trial in middle-aged men. Prostate, 2015, 75, 1255-1263.	1.2	63
57	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. Biological Psychiatry, 2022, 91, 102-117.	0.7	61
58	Archived neonatal dried blood spot samples can be used for accurate whole genome and exome-targeted next-generation sequencing. Molecular Genetics and Metabolism, 2013, 110, 65-72.	0.5	60
59	Blood Cytokine Profiles Associated with Distinct Patterns of Bronchopulmonary Dysplasia among Extremely Low Birth Weight Infants. Journal of Pediatrics, 2016, 174, 45-51.e5.	0.9	60
60	Genome-wide association study of panic disorder reveals genetic overlap with neuroticism and depression. Molecular Psychiatry, 2021, 26, 4179-4190.	4.1	58
61	Reduced neonatal brain-derived neurotrophic factor is associated with autism spectrum disorders. Translational Psychiatry, 2019, 9, 252.	2.4	56
62	Biochemical screening of 504,049 newborns in Denmark, the Faroe Islands and Greenland $\hat{a}\in$ " Experience and development of a routine program for expanded newborn screening. Molecular Genetics and Metabolism, 2012, 107, 281-293.	0.5	55
63	Association of Preterm Birth With Sustained Postnatal Inflammatory Response. Obstetrics and Gynecology, 2008, 111, 1118-1128.	1.2	52
64	The genetic architecture of sporadic and multiple consecutive miscarriage. Nature Communications, 2020, 11, 5980.	5.8	52
65	Genetic correlates of phenotypic heterogeneity in autism. Nature Genetics, 2022, 54, 1293-1304.	9.4	51
66	Variants in the fetal genome near pro-inflammatory cytokine genes on 2q13 associate with gestational duration. Nature Communications, 2019, 10, 3927.	5.8	49
67	Adipose tissue, estradiol levels, and bone health in obese men with metabolic syndrome. European Journal of Endocrinology, 2015, 172, 205-216.	1.9	48
68	Genetic liability to ADHD and substance use disorders in individuals with ADHD. Addiction, 2020, 115, 1368-1377.	1.7	47
69	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
70	Interaction Testing and Polygenic Risk Scoring to Estimate the Association of Common Genetic Variants With Treatment Resistance in Schizophrenia. JAMA Psychiatry, 2022, 79, 260.	6.0	44
71	Mid-pregnancy circulating immune biomarkers in women with preeclampsia and normotensive controls. Pregnancy Hypertension, 2016, 6, 72-78.	0.6	43
72	First and second trimester immune biomarkers in preeclamptic and normotensive women. Pregnancy Hypertension, 2016, 6, 388-393.	0.6	41

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73	Systemic and Local Inflammatory Response in Women with Preterm Prelabor Rupture of Membranes. PLoS ONE, 2014, 9, e85277.	1.1	40
74	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
75	Influence of Polygenic Risk Scores on the Association Between Infections and Schizophrenia. Biological Psychiatry, 2016, 80, 609-616.	0.7	38
76	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
77	High-Quality Exome Sequencing of Whole-Genome Amplified Neonatal Dried Blood Spot DNA. PLoS ONE, 2016, 11, e0153253.	1.1	38
78	A polygenic resilience score moderates the genetic risk for schizophrenia. Molecular Psychiatry, 2021, 26, 800-815.	4.1	36
79	Modelling the contribution of family history and variation in single nucleotide polymorphisms to risk of schizophrenia: A Danish national birth cohort-based study. Schizophrenia Research, 2012, 134, 246-252.	1.1	33
80	MCAD deficiency in Denmark. Molecular Genetics and Metabolism, 2012, 106, 175-188.	0.5	33
81	Amniotic Fluid INSL3 Measured During the Critical Time Window in Human Pregnancy Relates to Cryptorchidism, Hypospadias, and Phthalate Load: A Large Case–Control Study. Frontiers in Physiology, 2018, 9, 406.	1.3	33
82	Risk of Early-Onset Depression Associated With Polygenic Liability, Parental Psychiatric History, and Socioeconomic Status. JAMA Psychiatry, 2021, 78, 387.	6.0	33
83	Development of a Multiplex real-time PCR Assay for the Newborn Screening of SCID, SMA, and XLA. International Journal of Neonatal Screening, 2019, 5, 39.	1.2	32
84	Molecular genetic overlap between posttraumatic stress disorder and sleep phenotypes. Sleep, 2020, 43, .	0.6	32
85	Exploring Cuba's population structure and demographic history using genome-wide data. Scientific Reports, 2018, 8, 11422.	1.6	31
86	Identifying the Common Genetic Basis of Antidepressant Response. Biological Psychiatry Global Open Science, 2022, 2, 115-126.	1.0	31
87	The female protective effect against autism spectrum disorder. Cell Genomics, 2022, 2, 100134.	3.0	30
88	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
89	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
90	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

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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	Cystic fibrosis newborn screening in Denmark: Experience from the first 2 years. Pediatric Pulmonology, 2020, 55, 549-555.	1.0	26
93	Whole Genome Amplification and Genetic Analysis after Extraction of Proteins from Dried Blood Spots. Clinical Chemistry, 2007, 53, 1161-1162.	1.5	25
94	Comparing Copy Number Variations in a Danish Case Cohort of Individuals With Psychiatric Disorders. JAMA Psychiatry, 2022, 79, 59.	6.0	24
95	Decrease in Vitamin D Status in the Greenlandic Adult Population from 1987–2010. PLoS ONE, 2014, 9, e112949.	1.1	22
96	Roadmap for a precision-medicine initiative in the Nordic region. Nature Genetics, 2019, 51, 924-930.	9.4	22
97	Polygenic risk score, psychosocial environment and the risk of attention-deficit/hyperactivity disorder. Translational Psychiatry, 2020, 10, 335.	2.4	22
98	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
99	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
100	FUT2 $\hat{a}\in\text{``ABO}$ epistasis increases the risk of early childhood asthma and Streptococcus pneumoniae respiratory illnesses. Nature Communications, 2020, 11, 6398.	5.8	21
101	1,25-Dihydroxyvitamin D modulates L-type voltage-gated calcium channels in a subset of neurons in the developing mouse prefrontal cortex. Translational Psychiatry, 2019, 9, 281.	2.4	20
102	Neonatal metabolome of caesarean section and risk of childhood asthma. European Respiratory Journal, 2022, 59, 2102406.	3.1	20
103	Gene expression profiling of archived dried blood spot samples from the Danish Neonatal Screening Biobank. Molecular Genetics and Metabolism, 2015, 116, 119-124.	0.5	19
104	Neonatal Levels of Inflammatory Markers and Later Risk of Schizophrenia. Biological Psychiatry, 2015, 77, 548-555.	0.7	19
105	Mannose-Binding Lectin Gene, MBL2, Polymorphisms Are Not Associated With Susceptibility to Invasive Pneumococcal Disease in Children. Clinical Infectious Diseases, 2014, 59, e66-e71.	2.9	18
106	Genome-wide meta-analysis identifies <i>BARX1</i> and <i>EML4-MTA3</i> as new loci associated with infantile hypertrophic pyloric stenosis. Human Molecular Genetics, 2019, 28, 332-340.	1.4	18
107	Fetal Gender and Several Cytokines Are Associated with the Number of Fetal Cells in Maternal Blood – An Observational Study. PLoS ONE, 2014, 9, e106934.	1.1	17
108	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

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109	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
110	Genetic predictors of educational attainment and intelligence test performance predict voter turnout. Nature Human Behaviour, 2021, 5, 281-291.	6.2	15
111	Genetic, Clinical, and Sociodemographic Factors Associated With Stimulant Treatment Outcomes in ADHD. American Journal of Psychiatry, 2021, 178, 854-864.	4.0	15
112	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
113	Neonatal levels of adiponectin, interleukin- 10 and interleukin- 12 are associated with the risk of developing type 1 diabetes in childhood and adolescence: A nationwide Danish case-control study. Clinical Immunology, 2017 , 174 , 18 - 23 .	1.4	13
114	Genome-wide association study identifies locus at chromosome 2q32.1 associated with syncope and collapse. Cardiovascular Research, 2020, 116, 138-148.	1.8	13
115	Pharmacogenetic genotype and phenotype frequencies in a large Danish population-based case-cohort sample. Translational Psychiatry, 2021, 11, 294.	2.4	13
116	Novel feto-specific mRNA species suitable for identification of fetal cells from the maternal circulation. Prenatal Diagnosis, 2001, 21, 806-812.	1.1	12
117	Post-traumatic stress following military deployment: Genetic associations and cross-disorder genetic correlations. Journal of Affective Disorders, 2019, 252, 350-357.	2.0	12
118	No evidence of associations between genetic liability for schizophrenia and development of cannabis use disorder. Psychological Medicine, 2021, 51, 479-484.	2.7	12
119	Use of Molecular Genetic Analyses in Danish Routine Newborn Screening. International Journal of Neonatal Screening, 2021, 7, 50.	1.2	12
120	Translating polygenic risk scores for clinical use by estimating the confidence bounds of risk prediction. Nature Communications, 2021, 12, 5276.	5.8	12
121	Genome-wide association study identifies 16 genomic regions associated with circulating cytokines at birth. PLoS Genetics, 2020, 16 , e 1009163 .	1.5	12
122	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
123	Immunocytochemical Evidence Suggesting that Diamine Oxidase Catalyzes Biosynthesis of Î ³ -Aminobutyric Acid in Antropyloric Gastrin Cells. Journal of Histochemistry and Cytochemistry, 2000, 48, 839-846.	1.3	10
124	Neonatal Screening for Primary Carnitine Deficiency: Lessons Learned from the Faroe Islands. International Journal of Neonatal Screening, 2017, 3, 1.	1.2	10
125	Women With Turner Syndrome Are Both Estrogen and Androgen Deficient: The Impact of Hormone Replacement Therapy. Journal of Clinical Endocrinology and Metabolism, 2022, 107, 1983-1993.	1.8	10
126	Normal Levels of Plasma Free Carnitine and Acylcarnitines in Follow-Up Samples from a Presymptomatic Case of Carnitine Palmitoyl Transferase 1 (CPT1) Deficiency Detected Through Newborn Screening in Denmark. JIMD Reports, 2011, 3, 11-15.	0.7	9

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127	Evaluation of whole genome amplified DNA to decrease material expenditure and increase quality. Molecular Genetics and Metabolism Reports, 2017 , 11 , $36-45$.	0.4	9
128	Anorexia nervosa and inflammatory bowel diseasesâ€"Diagnostic and genetic associations. JCPP Advances, 2021, 1, e12036.	1.4	9
129	Associations between patterns in comorbid diagnostic trajectories of individuals with schizophrenia and etiological factors. Nature Communications, 2021, 12, 6617.	5.8	9
130	Vertical Transfer of Metabolites Detectable from Newborn's Dried Blood Spot Samples Using UPLC-MS: A Chemometric Study. Metabolites, 2022, 12, 94.	1.3	9
131	Genetic factors underlying the bidirectional relationship between autoimmune and mental disorders – Findings from a Danish population-based study. Brain, Behavior, and Immunity, 2021, 91, 10-23.	2.0	8
132	Co-occurring hydrocephalus in autism spectrum disorder: a Danish population-based cohort study. Journal of Neurodevelopmental Disorders, 2021, 13, 19.	1.5	8
133	Family disadvantage, gender, and the returns to genetic human capital*. Scandinavian Journal of Economics, 2022, 124, 550-578.	0.7	8
134	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
135	Carboxypeptidase�E in rat antropyloric mucosa: distribution in progenitor and mature endocrine cell types. Histochemistry and Cell Biology, 2004, 121, 55-61.	0.8	7
136	Abnormal Newborn Screening in a Healthy Infant of a Mother with Undiagnosed Medium-Chain Acyl-CoA Dehydrogenase Deficiency. JIMD Reports, 2015, 23, 67-70.	0.7	7
137	Early Elevation in Interleukin-6 is Associated with Reduced Growth in Extremely Low Birth Weight Infants. American Journal of Perinatology, 2017, 34, 240-247.	0.6	7
138	Including Classical Galactosaemia in the Expanded Newborn Screening Panel Using Tandem Mass Spectrometry for Galactose-1-Phosphate. International Journal of Neonatal Screening, 2019, 5, 19.	1.2	7
139	Polygenic liability, stressful life events and risk for secondary-treated depression in early life: a nationwide register-based case-cohort study. Psychological Medicine, 2023, 53, 217-226.	2.7	7
140	Polygenic Heterogeneity Across Obsessive-Compulsive Disorder Subgroups Defined by a Comorbid Diagnosis. Frontiers in Genetics, 2021, 12, 711624.	1.1	7
141	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
142	Differential effects of strength training and testosterone treatment on soluble CD36 in aging men: Possible relation to changes in body composition. Scandinavian Journal of Clinical and Laboratory Investigation, 2015, 75, 659-66.	0.6	6
143	Associations of 25 Hydroxyvitamin D and High Sensitivity C-reactive Protein Levels in Early Life. Nutrients, 2022, 14, 15.	1.7	6
144	Deep learning–based integration of genetics with registry data for stratification of schizophrenia and depression. Science Advances, 2022, 8, .	4.7	6

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145	Fetal sexual dimorphism in systemic soluble fmsâ€like tyrosine kinase 1 among normotensive and preeclamptic women. American Journal of Reproductive Immunology, 2018, 80, e13034.	1.2	5
146	Genetic liability to major depression and risk of childhood asthma. Brain, Behavior, and Immunity, 2020, 89, 433-439.	2.0	5
147	Implementation of SCID Screening in Denmark. International Journal of Neonatal Screening, 2021, 7, 54.	1.2	5
148	Raising Awareness of False Positive Newborn Screening Results Arising from Pivalate-Containing Creams and Antibiotics in Europe When Screening for Isovaleric Acidaemia. International Journal of Neonatal Screening, 2018, 4, 8.	1.2	4
149	Genome-wide by Environment Interaction Study of Stressful Life Events and Hospital-Treated Depression in the iPSYCH2012 Sample. Biological Psychiatry Global Open Science, 2022, 2, 400-410.	1.0	2
150	Danish expanded newborn screening is a successful preventive public health programme. Danish Medical Journal, 2020, 67, .	0.5	2
151	Protein Concentrations of Thrombospondin-1, MIP- $1\hat{l}^2$, and S100A8 Suggest the Reflection of a Pregnancy Clock in Mid-Trimester Amniotic Fluid. Reproductive Sciences, 2020, 27, 2146-2157.	1.1	1
152	Genetic regulation of spermine oxidase activity and cancer risk: a Mendelian randomization study. Scientific Reports, 2021, 11, 17463.	1.6	1
153	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
154	Genetic liability to posttraumatic stress disorder and its association with postpartum depression. Psychological Medicine, 0, , 1-8.	2.7	0