

Johan Hilge Thygesen

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

34 papers	2,550 citations	12 h-index	39 g-index
39 ext. papers	3,046 ext. citations	8.5 avg, IF	5.97 L-index

#	Paper	IF	Citations
34	Genome-wide association study identifies five new schizophrenia loci. <i>Nature Genetics</i> , 2011 , 43, 969-76	36.3	1508
33	CNVs conferring risk of autism or schizophrenia affect cognition in controls. <i>Nature</i> , 2014 , 505, 361-6	50.4	444
32	At-risk variant in TCF7L2 for type II diabetes increases risk of schizophrenia. <i>Biological Psychiatry</i> , 2011 , 70, 59-63	7.9	101
31	Runs of homozygosity implicate autozygosity as a schizophrenia risk factor. <i>PLoS Genetics</i> , 2012 , 8, e1002656	26.56	91
30	Genetic schizophrenia risk variants jointly modulate total brain and white matter volume. <i>Biological Psychiatry</i> , 2013 , 73, 525-31	7.9	87
29	Genome-wide meta-analysis of problematic alcohol use in 435,563 individuals yields insights into biology and relationships with other traits. <i>Nature Neuroscience</i> , 2020 , 23, 809-818	25.5	69
28	A polygenic risk score analysis of psychosis endophenotypes across brain functional, structural, and cognitive domains. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018 , 177, 21-34	3.5	39
27	Schizophrenia genetic variants are not associated with intelligence. <i>Psychological Medicine</i> , 2013 , 43, 2563-70	6.9	34
26	Variability in working memory performance explained by epistasis vs polygenic scores in the ZNF804A pathway. <i>JAMA Psychiatry</i> , 2014 , 71, 778-785	14.5	24
25	Chromosomal microarray testing in adults with intellectual disability presenting with comorbid psychiatric disorders. <i>European Journal of Human Genetics</i> , 2016 , 25, 66-72	5.3	24
24	Neurodevelopmental risk copy number variants in adults with intellectual disabilities and comorbid psychiatric disorders. <i>British Journal of Psychiatry</i> , 2018 , 212, 287-294	5.4	21
23	Use of schizophrenia and bipolar disorder polygenic risk scores to identify psychotic disorders. <i>British Journal of Psychiatry</i> , 2018 , 213, 535-541	5.4	21
22	Association between a genetic variant in the serotonin transporter gene (SLC6A4) and suicidal behavior in patients with schizophrenia. <i>Behavioral and Brain Functions</i> , 2012 , 8, 24	4.1	11
21	Sequence analysis of 17 NRXN1 deletions. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2014 , 165B, 52-61	3.5	10
20	Two rare deletions upstream of the NRXN1 gene (2p16.3) affecting the non-coding mRNA AK127244 segregate with diverse psychopathological phenotypes in a family. <i>European Journal of Medical Genetics</i> , 2015 , 58, 650-3	2.6	10
19	Lack of association between the regulator of G-protein signaling 4 (RGS4) rs951436 polymorphism and schizophrenia. <i>Psychiatric Genetics</i> , 2012 , 22, 263-4	2.9	9
18	Associations between psychosis endophenotypes across brain functional, structural, and cognitive domains. <i>Psychological Medicine</i> , 2018 , 48, 1325-1340	6.9	9

17	Linkage and whole genome sequencing identify a locus on 6q25-26 for formal thought disorder and implicate MEF2A regulation. <i>Schizophrenia Research</i> , 2015 , 169, 441-446	3.6	8
16	Evaluation of shared genetic susceptibility loci between autoimmune diseases and schizophrenia based on genome-wide association studies. <i>Nordic Journal of Psychiatry</i> , 2017 , 71, 20-25	2.3	7
15	Predictive power of the ADHD GWAS 2019 polygenic risk scores in independent samples of bipolar patients with childhood ADHD. <i>Journal of Affective Disorders</i> , 2020 , 265, 651-659	6.6	7
14	Genetic copy number variants, cognition and psychosis: a meta-analysis and a family study. <i>Molecular Psychiatry</i> , 2021 , 26, 5307-5319	15.1	5
13	The Influence of CYP2D6 and CYP2C19 Genetic Variation on Diabetes Mellitus Risk in People Taking Antidepressants and Antipsychotics. <i>Genes</i> , 2021 , 12,	4.2	3
12	The effect of CYP2D6 variation on antipsychotic-induced hyperprolactinaemia: a systematic review and meta-analysis. <i>Pharmacogenomics Journal</i> , 2020 , 20, 629-637	3.5	2
11	Transcriptome-wide association study reveals two genes that influence mismatch negativity. <i>Cell Reports</i> , 2021 , 34, 108868	10.6	2
10	Understanding COVID-19 trajectories from a nationwide linked electronic health record cohort of 56 million people: phenotypes, severity, waves & vaccination		1
9	Using de-identified electronic health records to research mental health supported housing services: A feasibility study. <i>PLoS ONE</i> , 2020 , 15, e0237664	3.7	1
8	Evaluation of antithrombotic use and COVID-19 outcomes in a nationwide atrial fibrillation cohort		1
7	Evaluation of antithrombotic use and COVID-19 outcomes in a nationwide atrial fibrillation cohort.. <i>Heart</i> , 2022 ,	5.1	1
6	The influence of regression models on genome-wide association studies of alcohol dependence: a comparison of binary and quantitative analyses. <i>Psychiatric Genetics</i> , 2021 , 31, 13-20	2.9	0
5	Antipsychotic polypharmacy and adverse drug reactions among adults in a London mental health service, 2008-2018.. <i>Psychological Medicine</i> , 2022 , 1-8	6.9	0
4	Using de-identified electronic health records to research mental health supported housing services: A feasibility study 2020 , 15, e0237664		
3	Using de-identified electronic health records to research mental health supported housing services: A feasibility study 2020 , 15, e0237664		
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