

# Johan Hilge Thygesen

## List of Publications by Year in Descending Order

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

| #  | Paper  | IF   | Citations |
|----|--|------|-----------|
| 34 | Evaluation of antithrombotic use and COVID-19 outcomes in a nationwide atrial fibrillation cohort.. <i>Heart</i> , <b>2022</b> ,   | 5.1  | 1         |
| 33 | Antipsychotic polypharmacy and adverse drug reactions among adults in a London mental health service, 2008-2018.. <i>Psychological Medicine</i> , <b>2022</b> , 1-8  | 6.9  | 0         |
| 32 | The Influence of CYP2D6 and CYP2C19 Genetic Variation on Diabetes Mellitus Risk in People Taking Antidepressants and Antipsychotics. <i>Genes</i> , <b>2021</b> , 12,  | 4.2  | 3         |
| 31 | The influence of regression models on genome-wide association studies of alcohol dependence: a comparison of binary and quantitative analyses. <i>Psychiatric Genetics</i> , <b>2021</b> , 31, 13-20                             | 2.9  | 0         |
| 30 | Genetic copy number variants, cognition and psychosis: a meta-analysis and a family study. <i>Molecular Psychiatry</i> , <b>2021</b> , 26, 5307-5319   | 15.1 | 5         |
| 29 | Transcriptome-wide association study reveals two genes that influence mismatch negativity. <i>Cell Reports</i> , <b>2021</b> , 34, 108868  | 10.6 | 2         |
| 28 | The effect of CYP2D6 variation on antipsychotic-induced hyperprolactinaemia: a systematic review and meta-analysis. <i>Pharmacogenomics Journal</i> , <b>2020</b> , 20, 629-637  | 3.5  | 2         |
| 27 | 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> , <b>2020</b> , 23, 809-818                             | 25.5 | 69        |
| 26 | 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> , <b>2020</b> , 265, 651-659                                  | 6.6  | 7         |
| 25 | Using de-identified electronic health records to research mental health supported housing services: A feasibility study. <i>PLoS ONE</i> , <b>2020</b> , 15, e0237664  | 3.7  | 1         |
| 24 | Using de-identified electronic health records to research mental health supported housing services: A feasibility study <b>2020</b> , 15, e0237664   |      |           |
| 23 | Using de-identified electronic health records to research mental health supported housing services: A feasibility study <b>2020</b> , 15, e0237664   |      |           |
| 22 | Using de-identified electronic health records to research mental health supported housing services: A feasibility study <b>2020</b> , 15, e0237664   |      |           |
| 21 | Using de-identified electronic health records to research mental health supported housing services: A feasibility study <b>2020</b> , 15, e0237664   |      |           |
| 20 | Neurodevelopmental risk copy number variants in adults with intellectual disabilities and comorbid psychiatric disorders. <i>British Journal of Psychiatry</i> , <b>2018</b> , 212, 287-294                                      | 5.4  | 21        |
| 19 | 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> , <b>2018</b> , 177, 21-34 | 3.5  | 39        |
| 18 | Use of schizophrenia and bipolar disorder polygenic risk scores to identify psychotic disorders. <i>British Journal of Psychiatry</i> , <b>2018</b> , 213, 535-541   | 5.4  | 21        |

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|----|---|------|------|
| 17 | Associations between psychosis endophenotypes across brain functional, structural, and cognitive domains. <i>Psychological Medicine</i> , <b>2018</b> , 48, 1325-1340   | 6.9  | 9    |
| 16 | Evaluation of shared genetic susceptibility loci between autoimmune diseases and schizophrenia based on genome-wide association studies. <i>Nordic Journal of Psychiatry</i> , <b>2017</b> , 71, 20-25                                | 2.3  | 7    |
| 15 | Chromosomal microarray testing in adults with intellectual disability presenting with comorbid psychiatric disorders. <i>European Journal of Human Genetics</i> , <b>2016</b> , 25, 66-72   | 5.3  | 24   |
| 14 | 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> , <b>2015</b> , 58, 650-3 | 2.6  | 10   |
| 13 | Linkage and whole genome sequencing identify a locus on 6q25-26 for formal thought disorder and implicate MEF2A regulation. <i>Schizophrenia Research</i> , <b>2015</b> , 169, 441-446  | 3.6  | 8    |
| 12 | Variability in working memory performance explained by epistasis vs polygenic scores in the ZNF804A pathway. <i>JAMA Psychiatry</i> , <b>2014</b> , 71, 778-785   | 14.5 | 24   |
| 11 | CNVs conferring risk of autism or schizophrenia affect cognition in controls. <i>Nature</i> , <b>2014</b> , 505, 361-6  | 50.4 | 444  |
| 10 | Sequence analysis of 17 NRXN1 deletions. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2014</b> , 165B, 52-61  | 3.5  | 10   |
| 9  | Genetic schizophrenia risk variants jointly modulate total brain and white matter volume. <i>Biological Psychiatry</i> , <b>2013</b> , 73, 525-31   | 7.9  | 87   |
| 8  | Schizophrenia genetic variants are not associated with intelligence. <i>Psychological Medicine</i> , <b>2013</b> , 43, 2563-70  | 6.9  | 34   |
| 7  | Association between a genetic variant in the serotonin transporter gene (SLC6A4) and suicidal behavior in patients with schizophrenia. <i>Behavioral and Brain Functions</i> , <b>2012</b> , 8, 24                                    | 4.1  | 11   |
| 6  | Runs of homozygosity implicate autozygosity as a schizophrenia risk factor. <i>PLoS Genetics</i> , <b>2012</b> , 8, e1002656  | 9.1  | 91   |
| 5  | Lack of association between the regulator of G-protein signaling 4 (RGS4) rs951436 polymorphism and schizophrenia. <i>Psychiatric Genetics</i> , <b>2012</b> , 22, 263-4  | 2.9  | 9    |
| 4  | At-risk variant in TCF7L2 for type II diabetes increases risk of schizophrenia. <i>Biological Psychiatry</i> , <b>2011</b> , 70, 59-63  | 7.9  | 101  |
| 3  | Genome-wide association study identifies five new schizophrenia loci. <i>Nature Genetics</i> , <b>2011</b> , 43, 969-76   | 36.3 | 1508 |
| 2  | Understanding COVID-19 trajectories from a nationwide linked electronic health record cohort of 56 million people: phenotypes, severity, waves & vaccination  |      | 1    |
| 1  | Evaluation of antithrombotic use and COVID-19 outcomes in a nationwide atrial fibrillation cohort   |      | 1    |