

Wouter J Peyrot

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

3,983
citations

21
h-index

36
g-index

36
ext. papers

5,558
ext. citations

15
avg, IF

4.39
L-index

| # | Paper | IF | Citations |
|----|---|------|-----------|
| 34 | Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. <i>Nature Genetics</i> , 2018 , 50, 668-681 | 36.3 | 1301 |
| 33 | Genome-wide association study identifies 74 loci associated with educational attainment. <i>Nature</i> , 2016 , 533, 539-42 | 50.4 | 850 |
| 32 | GWAS of 126,559 individuals identifies genetic variants associated with educational attainment. <i>Science</i> , 2013 , 340, 1467-71 | 33.3 | 563 |
| 31 | Genetic evidence of assortative mating in humans. <i>Nature Human Behaviour</i> , 2017 , 1, | 12.8 | 137 |
| 30 | Effect of polygenic risk scores on depression in childhood trauma. <i>British Journal of Psychiatry</i> , 2014 , 205, 113-9 | 5.4 | 133 |
| 29 | Genetic Association of Major Depression With Atypical Features and Obesity-Related Immunometabolic Dysregulations. <i>JAMA Psychiatry</i> , 2017 , 74, 1214-1225 | 14.5 | 109 |
| 28 | Genome-wide physical activity interactions in adiposity - A meta-analysis of 200,452 adults. <i>PLoS Genetics</i> , 2017 , 13, e1006528 | 6 | 103 |
| 27 | Genetic correlations of polygenic disease traits: from theory to practice. <i>Nature Reviews Genetics</i> , 2019 , 20, 567-581 | 30.1 | 98 |
| 26 | Genetic variants linked to education predict longevity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 13366-13371 | 11.5 | 90 |
| 25 | Risk of psychiatric illness from advanced paternal age is not predominantly from de novo mutations. <i>Nature Genetics</i> , 2016 , 48, 718-24 | 36.3 | 74 |
| 24 | The Genetics of the Mood Disorder Spectrum: Genome-wide Association Analyses of More Than 185,000 Cases and 439,000 Controls. <i>Biological Psychiatry</i> , 2020 , 88, 169-184 | 7.9 | 57 |
| 23 | Does Childhood Trauma Moderate Polygenic Risk for Depression? A Meta-analysis of 5765 Subjects From the Psychiatric Genomics Consortium. <i>Biological Psychiatry</i> , 2018 , 84, 138-147 | 7.9 | 48 |
| 22 | Genome-wide gene-environment analyses of major depressive disorder and reported lifetime traumatic experiences in UK Biobank. <i>Molecular Psychiatry</i> , 2020 , 25, 1430-1446 | 15.1 | 47 |
| 21 | Genome-wide gene-environment interaction in depression: A systematic evaluation of candidate genes: The childhood trauma working-group of PGC-MDD. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018 , 177, 40-49 | 3.5 | 43 |
| 20 | Collaborative stepped care v. care as usual for common mental disorders: 8-month, cluster randomised controlled trial. <i>British Journal of Psychiatry</i> , 2013 , 203, 132-9 | 5.4 | 39 |
| 19 | Exploring Boundaries for the Genetic Consequences of Assortative Mating for Psychiatric Traits. <i>JAMA Psychiatry</i> , 2016 , 73, 1189-1195 | 14.5 | 36 |
| 18 | Disease and Polygenic Architecture: Avoid Trio Design and Appropriately Account for Unscreened Control Subjects for Common Disease. <i>American Journal of Human Genetics</i> , 2016 , 98, 382-91 | 11 | 27 |

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| 17 | Using Clinical Characteristics to Identify Which Patients With Major Depressive Disorder Have a Higher Genetic Load for Three Psychiatric Disorders. <i>Biological Psychiatry</i> , 2017 , 81, 316-324 | 7.9 | 26 |
| 16 | Glucocorticoid and mineralocorticoid receptor polymorphisms and recurrence of major depressive disorder. <i>Psychoneuroendocrinology</i> , 2015 , 55, 154-63 | 5 | 23 |
| 15 | Strong effects of environmental factors on prevalence and course of major depressive disorder are not moderated by 5-HTTLPR polymorphisms in a large Dutch sample. <i>Journal of Affective Disorders</i> , 2013 , 146, 91-9 | 6.6 | 23 |
| 14 | Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depressive disorder | | 21 |
| 13 | A role for vitamin D and omega-3 fatty acids in major depression? An exploration using genomics. <i>Translational Psychiatry</i> , 2019 , 9, 219 | 8.6 | 19 |
| 12 | Association of polygenic score for major depression with response to lithium in patients with bipolar disorder. <i>Molecular Psychiatry</i> , 2021 , 26, 2457-2470 | 15.1 | 17 |
| 11 | Quantifying between-cohort and between-sex genetic heterogeneity in major depressive disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019 , 180, 439-447 | 3.5 | 16 |
| 10 | Identifying loci with different allele frequencies among cases of eight psychiatric disorders using CC-GWAS. <i>Nature Genetics</i> , 2021 , 53, 445-454 | 36.3 | 16 |
| 9 | Association of Whole-Genome and NETRIN1 Signaling Pathway-Derived Polygenic Risk Scores for Major Depressive Disorder and White Matter Microstructure in the UK Biobank. <i>Biological Psychiatry: Cognitive Neuroscience and Neuroimaging</i> , 2019 , 4, 91-100 | 3.4 | 12 |
| 8 | Genome-wide gene-environment analyses of depression and reported lifetime traumatic experiences in UK Biobank | | 9 |
| 7 | Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. <i>Biological Psychiatry</i> , 2020 , 87, 419-430 | 7.9 | 9 |
| 6 | The Genetic Architecture of Depression in Individuals of East Asian Ancestry: A Genome-Wide Association Study. <i>JAMA Psychiatry</i> , 2021 , 78, 1258-1269 | 14.5 | 7 |
| 5 | Polygenic prediction of educational attainment within and between families from genome-wide association analyses in 3 million individuals.. <i>Nature Genetics</i> , 2022 , | 36.3 | 7 |
| 4 | Genomic aberrations relate early and advanced stage ovarian cancer. <i>Cellular Oncology (Dordrecht)</i> , 2012 , 35, 181-8 | 7.2 | 6 |
| 3 | Identifying the Common Genetic Basis of Antidepressant Response. <i>Biological Psychiatry Global Open Science</i> , 2021 , | | 4 |
| 2 | Leveraging fine-mapping and multipopulation training data to improve cross-population polygenic risk scores.. <i>Nature Genetics</i> , 2022 , 54, 450-458 | 36.3 | 3 |
| 1 | Identifying loci with different allele frequencies among cases of eight psychiatric disorders using CC-GWAS | | 1 |