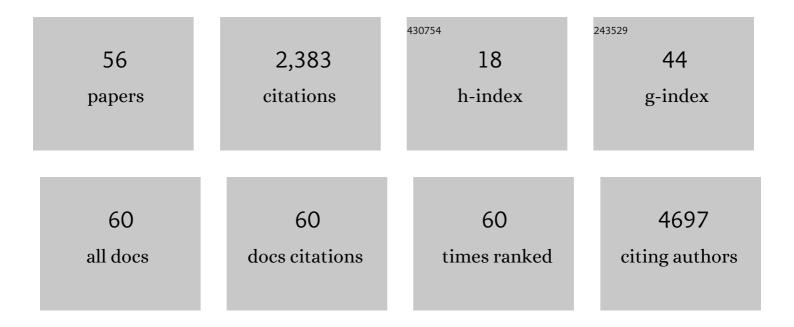
## Natalia Vilor-Tejedor

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9543925/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Polygenic risk for ADHD and ASD and their relation with cognitive measures in school children. Psychological Medicine, 2022, 52, 1356-1364.	2.7	14
2	Genome-wide Association Meta-analysis of Childhood and Adolescent Internalizing Symptoms. Journal of the American Academy of Child and Adolescent Psychiatry, 2022, 61, 934-945.	0.3	26
3	The protective gene dose effect of the <i>APOE ε2</i> allele on gray matter volume in cognitively unimpaired individuals. Alzheimer's and Dementia, 2022, 18, 1383-1395.	0.4	13
4	Study of the Combined Effect of Maternal Tobacco Smoking and Polygenic Risk Scores on Birth Weight and Body Mass Index in Childhood. Frontiers in Genetics, 2022, 13, .	1.1	1
5	Strategies to reduce sample sizes in Alzheimer's disease primary and secondary prevention trials using longitudinal amyloid PET imaging. Alzheimer's Research and Therapy, 2021, 13, 82.	3.0	14
6	Genetic Influences on Hippocampal Subfields. Neurology: Genetics, 2021, 7, e591.	0.9	8
7	Genetic Predisposition to Alzheimer's Disease Is Associated with Enlargement of Perivascular Spaces in Centrum Semiovale Region. Genes, 2021, 12, 825.	1.0	7
8	Genetic association study of childhood aggression across raters, instruments, and age. Translational Psychiatry, 2021, 11, 413.	2.4	31
9	Amyloid-β positive individuals with subjective cognitive decline present increased CSF neurofilament light levels that relate to lower hippocampal volume. Neurobiology of Aging, 2021, 104, 24-31.	1.5	13
10	Perivascular spaces are associated with tau pathophysiology and synaptic dysfunction in early Alzheimer's continuum. Alzheimer's Research and Therapy, 2021, 13, 135.	3.0	30
11	Continuity of Genetic Risk for Aggressive Behavior Across the Life-Course. Behavior Genetics, 2021, 51, 592-606.	1.4	13
12	CSF Synaptic Biomarkers in the Preclinical Stage of Alzheimer Disease and Their Association With MRI and PET. Neurology, 2021, 97, e2065-e2078.	1.5	40
13	DNA methylation signature as a biomarker of major neuropsychiatric disorders. Journal of Psychiatric Research, 2021, 141, 34-49.	1.5	24
14	Associations between air pollution and biomarkers of Alzheimer's disease in cognitively unimpaired individuals. Environment International, 2021, 157, 106864.	4.8	40
15	Multivariate Analysis and Modelling of multiple Brain endOphenotypes: Let's MAMBO!. Computational and Structural Biotechnology Journal, 2021, 19, 5800-5810.	1.9	4
16	Differences Between Plasma and Cerebrospinal Fluid Glial Fibrillary Acidic Protein Levels Across the Alzheimer Disease Continuum. JAMA Neurology, 2021, 78, 1471.	4.5	204
17	Single-cell Transcriptional Changes in Neurodegenerative Diseases. Neuroscience, 2021, 479, 192-205.	1.1	11
18	Association between telomere length and cognitive function among cognitively unimpaired individuals at risk of Alzheimer's disease. Alzheimer's and Dementia, 2021, 17, .	0.4	0

NATALIA VILOR-TEJEDOR

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19	Sex differences in genetic susceptibility of hippocampal subfields: A polygenic association study. Alzheimer's and Dementia, 2021, 17, .	0.4	0
20	Neuroimaging in neurodegeneration: A global diversity problem. Alzheimer's and Dementia, 2021, 17, .	0.4	0
21	Perivascular spaces are associated with tau pathophysiology and synaptic dysfunction in early Alzheimer's continuum. Alzheimer's and Dementia, 2021, 17, .	0.4	2
22	Crossâ€sectional associations between sleep quality reports and core Alzheimer's disease biomarkers in cognitively unimpaired adults from the European Prevention of Alzheimer's Dementia Longitudinal Cohort Study (EPAD LCS). Alzheimer's and Dementia, 2021, 17, .	0.4	0
23	Transposable elements in brain health and disease. Ageing Research Reviews, 2020, 64, 101153.	5.0	24
24	Novel loci for childhood body mass index and shared heritability with adult cardiometabolic traits. PLoS Genetics, 2020, 16, e1008718.	1.5	95
25	Aging-Dependent Genetic Effects Associated to ADHD Predict Longitudinal Changes of Ventricular Volumes in Adulthood. Frontiers in Psychiatry, 2020, 11, 574.	1.3	3
26	Effect of BDNF Val66Met on hippocampal subfields volumes and compensatory interaction with APOE-ε4 in middle-age cognitively unimpaired individuals from the ALFA study. Brain Structure and Function, 2020, 225, 2331-2345.	1.2	5
27	Polygenicâ€wide analysis to assess the impact of genetic risk profiles on brain morphometry in the ALFA study. Alzheimer's and Dementia, 2020, 16, e042952.	0.4	0
28	Amyloidâ€î², tau, synaptic dysfunction, neurodegeneration, glial and vascular biomarkers in the preclinical stage of the Alzheimer's continuum. Alzheimer's and Dementia, 2020, 16, e044444.	0.4	0
29	Genetically predicted telomere length and Alzheimer's disease endophenotypes: A Mendelian randomization study. Alzheimer's and Dementia, 2020, 16, e044720.	0.4	0
30	The effect of physical activity on CSF biomarkers of Alzheimer's disease differs between men and women. Alzheimer's and Dementia, 2020, 16, e044722.	0.4	0
31	Multiple biological pathways associate with cerebral amyloid load in the early Alzheimer's continuum. Alzheimer's and Dementia, 2020, 16, e044733.	0.4	0
32	Air pollution and biomarkers of Alzheimer's disease in cognitively unimpaired individuals. Alzheimer's and Dementia, 2020, 16, e044802.	0.4	3
33	APOE ―ε4 shapes temporoâ€parietal network properties in middleâ€aged, cognitively unimpaired individuals: A graph theory analysis. Alzheimer's and Dementia, 2020, 16, e045092.	0.4	0
34	Incidence of subjective cognitive decline is associated with amyloidâ€Ŷ pathology, whereas stability relates to neurodegeneration. Alzheimer's and Dementia, 2020, 16, e045293.	0.4	0
35	Amyloidâ€positive individuals with subjective cognitive decline present increased CSF neurofilament light levels that relate to hippocampal volume. Alzheimer's and Dementia, 2020, 16, e045715.	0.4	0
36	Amyloid beta, tau, synaptic, neurodegeneration, and glial biomarkers in the preclinical stage of the Alzheimer's <i>continuum</i> . Alzheimer's and Dementia, 2020, 16, 1358-1371.	0.4	120

NATALIA VILOR-TEJEDOR

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37	A trans-ancestral meta-analysis of genome-wide association studies reveals loci associated with childhood obesity. Human Molecular Genetics, 2019, 28, 3327-3338.	1.4	76
38	Effects of prenatal exposure to particulate matter air pollution on corpus callosum and behavioral problems in children. Environmental Research, 2019, 178, 108734.	3.7	55
39	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
40	Independent Multiple Factor Association Analysis for Multiblock Data in Imaging Genetics. Neuroinformatics, 2019, 17, 583-592.	1.5	2
41	Prenatal Omega-6:Omega-3 Ratio and Attention Deficit and Hyperactivity Disorder Symptoms. Journal of Pediatrics, 2019, 209, 204-211.e4.	0.9	28
42	Assessment of Susceptibility Risk Factors for ADHD in Imaging Genetic Studies. Journal of Attention Disorders, 2019, 23, 671-681.	1.5	17
43	Traffic-Related Air Pollution, <i>APOE</i> ε4 Status, and Neurodevelopmental Outcomes among School Children Enrolled in the BREATHE Project (Catalonia, Spain). Environmental Health Perspectives, 2018, 126, 087001.	2.8	53
44	Strategies for integrated analysis in imaging genetics studies. Neuroscience and Biobehavioral Reviews, 2018, 93, 57-70.	2.9	7
45	Sparse multiple factor analysis to integrate genetic data, neuroimaging features, and attentionâ€deficit/hyperactivity disorder domains. International Journal of Methods in Psychiatric Research, 2018, 27, e1738.	1.1	10
46	Interaction between airborne copper exposure and ATP7B polymorphisms on inattentiveness in scholar children. International Journal of Hygiene and Environmental Health, 2017, 220, 51-56.	2.1	14
47	Imaging genetics in attention-deficit/hyperactivity disorder and related neurodevelopmental domains: state of the art. Brain Imaging and Behavior, 2017, 11, 1922-1931.	1.1	10
48	A Genome-Wide Association Study of Attention Function in a Population-Based Sample of Children. PLoS ONE, 2016, 11, e0163048.	1.1	11
49	Genome-wide associations for birth weight and correlations with adult disease. Nature, 2016, 538, 248-252.	13.7	406
50	A Genome-Wide Association Meta-Analysis of Attention-Deficit/Hyperactivity Disorder Symptoms in Population-Based Pediatric Cohorts. Journal of the American Academy of Child and Adolescent Psychiatry, 2016, 55, 896-905.e6.	0.3	112
51	A genome-wide association meta-analysis of diarrhoeal disease in young children identifies <i>FUT2</i> locus and provides plausible biological pathways. Human Molecular Genetics, 2016, 25, 4127-4142.	1.4	35
52	Heritability and Genome-Wide Association Analyses of Sleep Duration in Children: The EAGLE Consortium. Sleep, 2016, 39, 1859-1869.	0.6	34
53	Efficient and Powerful Method for Combining P-Values in Genome-Wide Association Studies. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2016, 13, 1100-1106.	1.9	4
54	Genome-wide association analysis identifies three new susceptibility loci for childhood body mass index. Human Molecular Genetics, 2016, 25, 389-403.	1.4	275

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55	New suggestive genetic loci and biological pathways for attention function in adult attentionâ€deficit/hyperactivity disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 459-470.	1.1	78
56	Global adaptive rank truncated product method for geneâ€set analysis in association studies. Biometrical Journal, 2014, 56, 901-911.	0.6	3