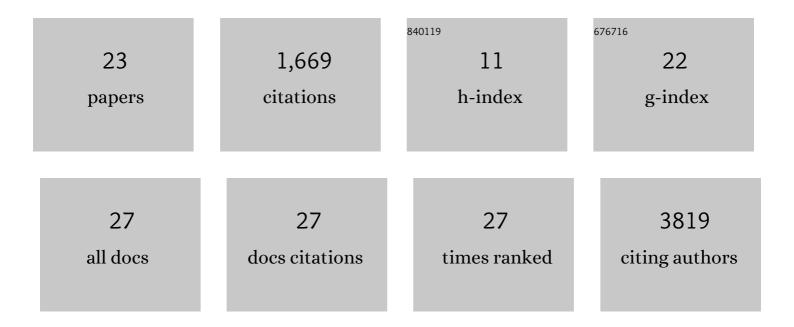
Nuria P Torres-Aguila

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6942480/publications.pdf

Version: 2024-02-01



#	Article	IF	CITATIONS
1	Evolutionary diversification of the canonical Wnt signaling effector TCF/LEF in chordates. Development Growth and Differentiation, 2022, , .	0.6	4
2	Multi-ancestry GWAS reveals excitotoxicity associated with outcome after ischaemic stroke. Brain, 2022, 145, 2394-2406.	3.7	15
3	DNA Methylation and Ischemic Stroke Risk: An Epigenome-Wide Association Study. Thrombosis and Haemostasis, 2022, 122, 1767-1778.	1.8	12
4	Early Neurological Change After Ischemic Stroke Is Associated With 90-Day Outcome. Stroke, 2021, 52, 132-141.	1.0	36
5	Single nucleotide variations in <i>ZBTB46</i> are associated with post-thrombolytic parenchymal haematoma. Brain, 2021, 144, 2416-2426.	3.7	10
6	Genome-wide transcriptome study in skin biopsies reveals an association of E2F4 with cadasil and cognitive impairment. Scientific Reports, 2021, 11, 6846.	1.6	5
7	Genetic basis of lacunar stroke: a pooled analysis of individual patient data and genome-wide association studies. Lancet Neurology, The, 2021, 20, 351-361.	4.9	95
8	A parsimonious score with a free web tool for predicting disability after an ischemic stroke: the Parsifal Score. Journal of Neurology, 2020, 267, 2871-2880.	1.8	0
9	Genome-Wide Association Study of VKORC1 and CYP2C9 on acenocoumarol dose, stroke recurrence and intracranial haemorrhage in Spain. Scientific Reports, 2020, 10, 2806.	1.6	7
10	DNA methylation of MMPs and TIMPs in atherothrombosis process in carotid plaques and blood tissues. Oncotarget, 2020, 11, 905-912.	0.8	4
11	Validation of a clinical-genetics score to predict hemorrhagic transformations after rtPA. Neurology, 2019, 93, e851-e863.	1.5	10
12	Genome-Wide Association Study of White Blood Cell Counts in Patients With Ischemic Stroke. Stroke, 2019, 50, 3618-3621.	1.0	13
13	Serum magnesium and calcium levels in relation to ischemic stroke. Neurology, 2019, 92, e944-e950.	1.5	38
14	<i>PATJ</i> Low Frequency Variants Are Associated With Worse Ischemic Stroke Functional Outcome. Circulation Research, 2019, 124, 114-120.	2.0	49
15	Clinical Variables and Genetic Risk Factors Associated with the Acute Outcome of Ischemic Stroke: A Systematic Review. Journal of Stroke, 2019, 21, 276-289.	1.4	27
16	DNA Methylation in Stroke. Update of Latest Advances. Computational and Structural Biotechnology Journal, 2018, 16, 1-5.	1.9	29
17	Pharmacogenetic studies with oral anticoagulants. Genome-wide association studies in vitamin K antagonist and direct oral anticoagulants. Oncotarget, 2018, 9, 29238-29258.	0.8	26
18	Diatom bloom-derived biotoxins cause aberrant development and gene expression in the appendicularian chordate Oikopleura dioica. Communications Biology, 2018, 1, 121.	2.0	12

#	Article	IF	CITATIONS
19	Multiancestry genome-wide association study of 520,000 subjects identifies 32 loci associated with stroke and stroke subtypes. Nature Genetics, 2018, 50, 524-537.	9.4	1,124
20	Role of TRAF3 in neurological and cardiovascular diseases: an overview of recent studies. Biomolecular Concepts, 2017, 8, 197-202.	1.0	9
21	Systematic Review of Cysteine-Sparing NOTCH3 Missense Mutations in Patients with Clinical Suspicion of CADASIL. International Journal of Molecular Sciences, 2017, 18, 1964.	1.8	62
22	Coelimination and Survival in Gene Network Evolution: Dismantling the RA-Signaling in a Chordate. Molecular Biology and Evolution, 2016, 33, 2401-2416.	3.5	39
23	<i>Oikopleura dioica</i> culturing made easy: A Lowâ€Cost facility for an emerging animal model in <scp>E</scp> vo <scp>D</scp> evo. Genesis, 2015, 53, 183-193.	0.8	31