Joseph S Reddy

List of Publications by Year
in descending order

24	768	687363 13	642732 23
papers	citations	h-index	g-index

34	34	34	1855
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Alzheimerâ€™s disease and progressive supranuclear palsy share similar transcriptomic changes in distinct brain regions. Journal of Clinical Investigation, 2022, 132, .	8.2	13
2	Transcript levels in plasma contribute substantial predictive value as potential Alzheimer's disease biomarkers in African Americans. EBioMedicine, 2022, , 103929.	6.1	2
3	Whole genome sequencingâ€“based copy number variations reveal novel pathways and targets in Alzheimer's disease. Alzheimer's and Dementia, 2022, 18, 1846-1867.	0.8	13
4	Plasma Biomarkers of Alzheimerâ€™s Disease in African Americans. Journal of Alzheimer's Disease, 2021, 79, 323-334.	2.6	11
5	Identifying drug targets for neurological and psychiatric disease via genetics and the brain transcriptome. PLoS Genetics, 2021, 17, e1009224.	3.5	43
6	Latent trait modeling of tau neuropathology in progressive supranuclear palsy. Acta Neuropathologica, 2021, 141, 667-680.	7.7	5
7	Impact of variant-level batch effects on identification of genetic risk factors in large sequencing studies. PLoS ONE, 2021, 16, e0249305.	2.5	5
8	Genome-wide analysis identifies a novel LINC-PINT splice variant associated with vascular amyloid pathology in Alzheimerâ€™s disease. Acta Neuropathologica Communications, 2021, 9, 93.	5.2	9
9	Large eQTL meta-analysis reveals differing patterns between cerebral cortical and cerebellar brain regions. Scientific Data, 2020, 7, 340.	5.3	75
10	Deciphering cellular transcriptional alterations in Alzheimerâ€™s disease brains. Molecular Neurodegeneration, 2020, 15, 38.	10.8	42
11	Association of ABI3 and PLCG2 missense variants with disease risk and neuropathology in Lewy body disease and progressive supranuclear palsy. Acta Neuropathologica Communications, 2020, 8, 172.	5.2	8
12	MAPT haplotypeâ€“stratified GWAS reveals differential association for AD risk variants. Alzheimer's and Dementia, 2020, 16, 983-1002.	0.8	21
13	Extensive transcriptomic study emphasizes importance of vesicular transport in C9orf72 expansion carriers. Acta Neuropathologica Communications, 2019, 7, 150.	5.2	40
14	Systematic analysis of dark and camouflaged genes reveals disease-relevant genes hiding in plain sight. Genome Biology, 2019, 20, 97.	8.8	122
15	ABI3 and PLCG2 missense variants as risk factors for neurodegenerative diseases in Caucasians and African Americans. Molecular Neurodegeneration, 2018, 13, 53.	10.8	75
16	TLR5 decoy receptor as a novel anti-amyloid therapeutic for Alzheimerâ€™s disease. Journal of Experimental Medicine, 2018, 215, 2247-2264.	8.5	50
17	TMEM106B haplotypes have distinct gene expression patterns in aged brain. Molecular Neurodegeneration, 2018, 13, 35.	10.8	30
18	Identification of missing variants by combining multiple analytic pipelines. BMC Bioinformatics, 2018, 19, 139.	2.6	10

#	Article	IF	Citations
19	Divergent brain gene expression patterns associate with distinct cell-specific tau neuropathology traits in progressive supranuclear palsy. Acta Neuropathologica, 2018, 136, 709-727.	7.7	47
20	ABCA7 loss-of-function variants, expression, and neurologic disease risk. Neurology: Genetics, 2017, 3, e126.	1.9	26
21	Comparative Genomics and Transcriptional Analysis of Flavobacterium columnare Strain ATCC 49512. Frontiers in Microbiology, 2017, 8, 588.	3.5	46
22	The Effect of Oxygen on Bile Resistance in Listeria monocytogenes. Journal of Proteomics and Bioinformatics, 2016, 04, 107-119.	0.4	25
23	Comparative Proteomic Analysis of Cotton Fiber Development and Protein Extraction Method Comparison in Late Stage Fibers. Proteomes, 2016, 4, 7.	3.5	10
24	Transcriptome profile of a bovine respiratory disease pathogen: Mannheimia haemolytica PHL213. BMC Bioinformatics, 2012, 13, S4.	2.6	11