

Joseph S Reddy

List of Publications by Year in descending order

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Version: 2024-02-01

24
papers

768
citations

687363

13
h-index

642732

23
g-index

34
all docs

34
docs citations

34
times ranked

1855
citing authors

#	ARTICLE	IF	CITATIONS
1	Systematic analysis of dark and camouflaged genes reveals disease-relevant genes hiding in plain sight. <i>Genome Biology</i> , 2019, 20, 97.	8.8	122
2	ABI3 and PLCG2 missense variants as risk factors for neurodegenerative diseases in Caucasians and African Americans. <i>Molecular Neurodegeneration</i> , 2018, 13, 53.	10.8	75
3	Large eQTL meta-analysis reveals differing patterns between cerebral cortical and cerebellar brain regions. <i>Scientific Data</i> , 2020, 7, 340.	5.3	75
4	TLR5 decoy receptor as a novel anti-amyloid therapeutic for Alzheimer's disease. <i>Journal of Experimental Medicine</i> , 2018, 215, 2247-2264.	8.5	50
5	Divergent brain gene expression patterns associate with distinct cell-specific tau neuropathology traits in progressive supranuclear palsy. <i>Acta Neuropathologica</i> , 2018, 136, 709-727.	7.7	47
6	Comparative Genomics and Transcriptional Analysis of <i>Flavobacterium columnare</i> Strain ATCC 49512. <i>Frontiers in Microbiology</i> , 2017, 8, 588.	3.5	46
7	Identifying drug targets for neurological and psychiatric disease via genetics and the brain transcriptome. <i>PLoS Genetics</i> , 2021, 17, e1009224.	3.5	43
8	Deciphering cellular transcriptional alterations in Alzheimer's disease brains. <i>Molecular Neurodegeneration</i> , 2020, 15, 38.	10.8	42
9	Extensive transcriptomic study emphasizes importance of vesicular transport in C9orf72 expansion carriers. <i>Acta Neuropathologica Communications</i> , 2019, 7, 150.	5.2	40
10	TMEM106B haplotypes have distinct gene expression patterns in aged brain. <i>Molecular Neurodegeneration</i> , 2018, 13, 35.	10.8	30
11	<i>ABCA7</i> loss-of-function variants, expression, and neurologic disease risk. <i>Neurology: Genetics</i> , 2017, 3, e126.	1.9	26
12	The Effect of Oxygen on Bile Resistance in <i>Listeria monocytogenes</i> . <i>Journal of Proteomics and Bioinformatics</i> , 2016, 04, 107-119.	0.4	25
13	<i>MAPT</i> haplotype-stratified GWAS reveals differential association for AD risk variants. <i>Alzheimer's and Dementia</i> , 2020, 16, 983-1002.	0.8	21
14	Alzheimer's disease and progressive supranuclear palsy share similar transcriptomic changes in distinct brain regions. <i>Journal of Clinical Investigation</i> , 2022, 132, .	8.2	13
15	Whole genome sequencing-based copy number variations reveal novel pathways and targets in Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2022, 18, 1846-1867.	0.8	13
16	Transcriptome profile of a bovine respiratory disease pathogen: <i>Mannheimia haemolytica</i> PHL213. <i>BMC Bioinformatics</i> , 2012, 13, S4.	2.6	11
17	Plasma Biomarkers of Alzheimer's Disease in African Americans. <i>Journal of Alzheimer's Disease</i> , 2021, 79, 323-334.	2.6	11
18	Comparative Proteomic Analysis of Cotton Fiber Development and Protein Extraction Method Comparison in Late Stage Fibers. <i>Proteomes</i> , 2016, 4, 7.	3.5	10

#	ARTICLE	IF	CITATIONS
19	Identification of missing variants by combining multiple analytic pipelines. BMC Bioinformatics, 2018, 19, 139.	2.6	10
20	Genome-wide analysis identifies a novel LINC-PINT splice variant associated with vascular amyloid pathology in Alzheimer's disease. Acta Neuropathologica Communications, 2021, 9, 93.	5.2	9
21	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
22	Latent trait modeling of tau neuropathology in progressive supranuclear palsy. Acta Neuropathologica, 2021, 141, 667-680.	7.7	5
23	Impact of variant-level batch effects on identification of genetic risk factors in large sequencing studies. PLoS ONE, 2021, 16, e0249305.	2.5	5
24	Transcript levels in plasma contribute substantial predictive value as potential Alzheimer's disease biomarkers in African Americans. EBioMedicine, 2022, , 103929.	6.1	2