Andrew T Mckenzie

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1178857/publications.pdf

Version: 2024-02-01

759233 16 1,692 12 citations papers

14 h-index g-index 20 20 20 3636 docs citations times ranked citing authors all docs

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#	Article	IF	CITATIONS
1	The innate immunity protein IFITM3 modulates γ-secretase in Alzheimer's disease. Nature, 2020, 586, 735-740.	27.8	219
2	Stress resilience is promoted by a Zfp189-driven transcriptional network in prefrontal cortex. Nature Neuroscience, 2019, 22, 1413-1423.	14.8	78
3	Perfusion fixation in brain banking: a systematic review. Acta Neuropathologica Communications, 2019, 7, 146.	5.2	36
4	The landscape of multiscale transcriptomic networks and key regulators in Parkinson's disease. Nature Communications, 2019, 10, 5234.	12.8	82
5	Brain Cell Type Specific Gene Expression and Co-expression Network Architectures. Scientific Reports, 2018, 8, 8868.	3.3	335
6	A common haplotype lowers PU.1 expression in myeloid cells and delays onset of Alzheimer's disease. Nature Neuroscience, 2017, 20, 1052-1061.	14.8	330
7	Multiscale network modeling of oligodendrocytes reveals molecular components of myelin dysregulation in Alzheimer's disease. Molecular Neurodegeneration, 2017, 12, 82.	10.8	100
8	DGCA: A comprehensive R package for Differential Gene Correlation Analysis. BMC Systems Biology, 2016, 10, 106.	3.0	171
9	S4â€02â€03: Accelerating Medicines Partnership: Coâ€Expression Networks. Alzheimer's and Dementia, 2016, 12, P322.	0.8	O
10	O2â€10â€06: A Common Allele in <i>SPI1</i> Lowers Risk and Delays Age at Onset for Alzheimer's Disease. Alzheimer's and Dementia, 2016, 12, P253.	0.8	0
11	The human-specific <i>CASP4</i> gene product contributes to Alzheimer-related synaptic and behavioural deficits. Human Molecular Genetics, 2016, 25, 4315-4327.	2.9	21
12	F2â€01â€01: Oligodendrocyteâ€Enriched Gene Networks Reveal Novel Pathways and Key Targets in the Pathogenesis of Alzheimer's Disease. Alzheimer's and Dementia, 2016, 12, P214.	0.8	0
13	Integrative network analysis of nineteen brain regions identifies molecular signatures and networks underlying selective regional vulnerability to Alzheimer's disease. Genome Medicine, 2016, 8, 104.	8.2	224
14	Transcriptome analysis identifies Bacillus anthracis genes that respond to CO2through an AtxA-dependent mechanism. BMC Genomics, 2014, 15, 229.	2.8	20
15	Transcriptional analysis of the three Nlrp1 paralogs in mice. BMC Genomics, 2013, 14, 188.	2.8	62
16	Tumor therapy with a urokinase plasminogen activator-activated anthrax lethal toxin alone and in combination with paclitaxel. Investigational New Drugs, 2013, 31, 206-212.	2.6	8