

Andrew T N Tebbenkamp

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

2,718
citations

567281

15
h-index

888059

17
g-index

19
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19
docs citations

19
times ranked

5871
citing authors

#	ARTICLE	IF	CITATIONS
1	Transcriptomic taxonomy and neurogenic trajectories of adult human, macaque, and pig hippocampal and entorhinal cells. <i>Neuron</i> , 2022, 110, 452-469.e14.	8.1	142
2	Whole-Genome and RNA Sequencing Reveal Variation and Transcriptomic Coordination in the Developing Human Prefrontal Cortex. <i>Cell Reports</i> , 2020, 31, 107489.	6.4	91
3	Integrative functional genomic analysis of human brain development and neuropsychiatric risks. <i>Science</i> , 2018, 362, .	12.6	516
4	The 7q11.23 Protein DNAJC30 Interacts with ATP Synthase and Links Mitochondria to Brain Development. <i>Cell</i> , 2018, 175, 1088-1104.e23.	28.9	46
5	Molecular and cellular reorganization of neural circuits in the human lineage. <i>Science</i> , 2017, 358, 1027-1032.	12.6	192
6	Zika Virus Disrupts Phospho-TBK1 Localization and Mitosis in Human Neuroepithelial Stem Cells and Radial Glia. <i>Cell Reports</i> , 2016, 16, 2576-2592.	6.4	253
7	The autism-associated chromatin modifier CHD8 regulates other autism risk genes during human neurodevelopment. <i>Nature Communications</i> , 2015, 6, 6404.	12.8	316
8	The developmental transcriptome of the human brain. <i>Current Opinion in Neurology</i> , 2014, 27, 149-156.	3.6	100
9	Experimental Mutagenesis of Huntingtin to Map Cleavage Sites: Different Outcomes in Cell and Mouse Models. <i>Journal of Huntington's Disease</i> , 2014, 3, 73-86.	1.9	1
10	Coexpression Networks Implicate Human Midfetal Deep Cortical Projection Neurons in the Pathogenesis of Autism. <i>Cell</i> , 2013, 155, 997-1007.	28.9	825
11	Analysis of Proteolytic Processes and Enzymatic Activities in the Generation of Huntingtin N-Terminal Fragments in an HEK293 Cell Model. <i>PLoS ONE</i> , 2012, 7, e50750.	2.5	22
12	Transgenic mice expressing caspase-6-derived N-terminal fragments of mutant huntingtin develop neurologic abnormalities with predominant cytoplasmic inclusion pathology composed largely of a smaller proteolytic derivative. <i>Human Molecular Genetics</i> , 2011, 20, 2770-2782.	2.9	39
13	Premature death and neurologic abnormalities in transgenic mice expressing a mutant huntingtin exon-2 fragment. <i>Human Molecular Genetics</i> , 2011, 20, 1633-1642.	2.9	22
14	Partial Depletion of CREB-Binding Protein Reduces Life Expectancy in a Mouse Model of Huntington Disease. <i>Journal of Neuropathology and Experimental Neurology</i> , 2010, 69, 396-404.	1.7	24
15	Analysis of Chaperone mRNA Expression in the Adult Mouse Brain by Meta Analysis of the Allen Brain Atlas. <i>PLoS ONE</i> , 2010, 5, e13675.	2.5	32
16	Protein Aggregate Characterization in Models of Neurodegenerative Disease. <i>Methods in Molecular Biology</i> , 2009, 566, 85-91.	0.9	11
17	Characterization of Huntingtin Pathologic Fragments in Human Huntington Disease, Transgenic Mice, and Cell Models. <i>Journal of Neuropathology and Experimental Neurology</i> , 2007, 66, 313-320.	1.7	72