

Jeremy C Davis-Turak

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

14
papers

1,044
citations

12
h-index

16
g-index

16
ext. papers

1,305
ext. citations

11.1
avg, IF

3.26
L-index

#	Paper	IF	Citations
14	A Systems-Level Analysis of the Peripheral Nerve Intrinsic Axonal Growth Program. <i>Neuron</i> , 2016 , 89, 956-70	13.9	206
13	Human-specific transcriptional networks in the brain. <i>Neuron</i> , 2012 , 75, 601-17	13.9	180
12	Transcriptional architecture of the primate neocortex. <i>Neuron</i> , 2012 , 73, 1083-99	13.9	170
11	Control of RelB during dendritic cell activation integrates canonical and noncanonical NF- κ B pathways. <i>Nature Immunology</i> , 2012 , 13, 1162-70	19.1	133
10	Tauopathy with paired helical filaments in an aged chimpanzee. <i>Journal of Comparative Neurology</i> , 2008 , 509, 259-70	3.4	105
9	A multiplex RNA-seq strategy to profile poly(A+) RNA: application to analysis of transcription response and 3'end formation. <i>Genomics</i> , 2011 , 98, 266-71	4.3	53
8	IB κ s a key regulator of B cell expansion by providing negative feedback on cRel and RelA in a stimulus-specific manner. <i>Journal of Immunology</i> , 2014 , 192, 3121-32	5.3	42
7	Genomics pipelines and data integration: challenges and opportunities in the research setting. <i>Expert Review of Molecular Diagnostics</i> , 2017 , 17, 225-237	3.8	38
6	A multi-ancestral genome-wide exome array study of Alzheimer disease, frontotemporal dementia, and progressive supranuclear palsy. <i>JAMA Neurology</i> , 2015 , 72, 414-22	17.2	33
5	A pathway switch directs BAFF signaling to distinct NF κ B transcription factors in maturing and proliferating B cells. <i>Cell Reports</i> , 2014 , 9, 2098-111	10.6	27
4	B-cell survival and development controlled by the coordination of NF- κ B family members RelB and cRel. <i>Blood</i> , 2016 , 127, 1276-86	2.2	24
3	A multi-scale approach reveals that NF- κ B cRel enforces a B-cell decision to divide. <i>Molecular Systems Biology</i> , 2015 , 11, 783	12.2	16
2	Considering the kinetics of mRNA synthesis in the analysis of the genome and epigenome reveals determinants of co-transcriptional splicing. <i>Nucleic Acids Research</i> , 2015 , 43, 699-707	20.1	12
1	Mathematical modeling identifies potential gene structure determinants of co-transcriptional control of alternative pre-mRNA splicing. <i>Nucleic Acids Research</i> , 2018 , 46, 10598-10607	20.1	5