

Jeremy C Davis-Turak

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4532031/publications.pdf>

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14
papers

1,431
citations

686830

13
h-index

1058022

14
g-index

16
all docs

16
docs citations

16
times ranked

3386
citing authors

#	ARTICLE	IF	CITATIONS
1	A Systems-Level Analysis of the Peripheral Nerve Intrinsic Axonal Growth Program. <i>Neuron</i> , 2016, 89, 956-970.	3.8	314
2	Human-Specific Transcriptional Networks in the Brain. <i>Neuron</i> , 2012, 75, 601-617.	3.8	256
3	Transcriptional Architecture of the Primate Neocortex. <i>Neuron</i> , 2012, 73, 1083-1099.	3.8	234
4	Control of RelB during dendritic cell activation integrates canonical and noncanonical NF- κ B pathways. <i>Nature Immunology</i> , 2012, 13, 1162-1170.	7.0	166
5	Tauopathy with paired helical filaments in an aged chimpanzee. <i>Journal of Comparative Neurology</i> , 2008, 509, 259-270.	0.9	129
6	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, 266-271.	1.3	61
7	Genomics pipelines and data integration: challenges and opportunities in the research setting. <i>Expert Review of Molecular Diagnostics</i> , 2017, 17, 225-237.	1.5	54
8	μ Is 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, 3121-3132.	0.4	51
9	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, 2098-2111.	2.9	43
10	B-cell survival and development controlled by the coordination of NF- κ B family members RelB and cRel. <i>Blood</i> , 2016, 127, 1276-1286.	0.6	38
11	A Multiancestral Genome-Wide Exome Array Study of Alzheimer Disease, Frontotemporal Dementia, and Progressive Supranuclear Palsy. <i>JAMA Neurology</i> , 2015, 72, 414.	4.5	37
12	A multi-scale approach reveals that NF κ B el enforces a B cell decision to divide. <i>Molecular Systems Biology</i> , 2015, 11, 783.	3.2	25
13	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.	6.5	15
14	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.	6.5	8