

Brian T Do

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/9571177/brian-t-do-publications-by-year.pdf>
Version: 2024-04-09

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.
The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

13 papers	1,456 citations	11 h-index	15 g-index
15 ext. papers	2,162 ext. citations	15.2 avg, IF	4.03 L-index

#	Paper	IF	Citations
13	Regulation of chromatin accessibility by the histone chaperone CAF-1 sustains lineage fidelity.. <i>Nature Communications</i> , 2022 , 13, 2350	17.4	0
12	Cell-programmed nutrient partitioning in the tumour microenvironment. <i>Nature</i> , 2021 , 593, 282-288	50.4	111
11	Hepcidin sequesters iron to sustain nucleotide metabolism and mitochondrial function in colorectal cancer epithelial cells. <i>Nature Metabolism</i> , 2021 , 3, 969-982	14.6	12
10	Increased demand for NAD relative to ATP drives aerobic glycolysis. <i>Molecular Cell</i> , 2021 , 81, 691-707.e6	17.6	58
9	Loss of FLT3 Sensitizes Myeloid Cells to Differentiation Via DHODH Inhibition. <i>Blood</i> , 2019 , 134, 2712-2712	12	2
8	Opportunities and obstacles for deep learning in biology and medicine. <i>Journal of the Royal Society Interface</i> , 2018 , 15,	4.1	780
7	7SK-BAF axis controls pervasive transcription at enhancers. <i>Nature Structural and Molecular Biology</i> , 2016 , 23, 231-8	17.6	60
6	Medulloblastoma-associated DDX3 variant selectively alters the translational response to stress. <i>Oncotarget</i> , 2016 , 7, 28169-82	3.3	44
5	Stress from Nucleotide Depletion Activates the Transcriptional Regulator HEXIM1 to Suppress Melanoma. <i>Molecular Cell</i> , 2016 , 62, 34-46	17.6	52
4	irCLIP platform for efficient characterization of protein-RNA interactions. <i>Nature Methods</i> , 2016 , 13, 489-92	21.6	151
3	Dissecting noncoding and pathogen RNA-protein interactomes. <i>Rna</i> , 2015 , 21, 135-43	5.8	61
2	Dicer-microRNA-Myc circuit promotes transcription of hundreds of long noncoding RNAs. <i>Nature Structural and Molecular Biology</i> , 2014 , 21, 585-90	17.6	80
1	Opportunities and obstacles for deep learning in biology and medicine		45