## Dan Dominissini

## List of Publications by Citations

Source: https://exaly.com/author-pdf/7717806/dan-dominissini-publications-by-citations.pdf

Version: 2024-04-28

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.

16 27 5,932 30 g-index h-index citations papers 5.61 19.6 30 7,795 avg, IF L-index ext. citations ext. papers

#	Paper	IF	Citations
27	Topology of the human and mouse m6A RNA methylomes revealed by m6A-seq. <i>Nature</i> , <b>2012</b> , 485, 201	- <b>6</b> 0.4	2387
26	Gene expression regulation mediated through reversible mA RNA methylation. <i>Nature Reviews Genetics</i> , <b>2014</b> , 15, 293-306	30.1	905
25	Stem cells. m6A mRNA methylation facilitates resolution of naMe pluripotency toward differentiation. <i>Science</i> , <b>2015</b> , 347, 1002-6	33.3	904
24	The dynamic N(1)-methyladenosine methylome in eukaryotic messenger RNA. <i>Nature</i> , <b>2016</b> , 530, 441-6	50.4	523
23	Transcriptome-wide mapping of N(6)-methyladenosine by m(6)A-seq based on immunocapturing and massively parallel sequencing. <i>Nature Protocols</i> , <b>2013</b> , 8, 176-89	18.8	358
22	Nm-seq maps 2VO-methylation sites in human mRNA with base precision. <i>Nature Methods</i> , <b>2017</b> , 14, 695-698	21.6	146
21	N(6)-methyladenosine in mRNA disrupts tRNA selection and translation-elongation dynamics. <i>Nature Structural and Molecular Biology</i> , <b>2016</b> , 23, 110-5	17.6	139
20	N6-methyladenosine modification and the YTHDF2 reader protein play cell type specific roles in lytic viral gene expression during Kaposi's sarcoma-associated herpesvirus infection. <i>PLoS Pathogens</i> , <b>2018</b> , 14, e1006995	7.6	102
19	The mA epitranscriptome: transcriptome plasticity in brain development and function. <i>Nature Reviews Neuroscience</i> , <b>2020</b> , 21, 36-51	13.5	86
18	Adenosine-to-inosine RNA editing meets cancer. <i>Carcinogenesis</i> , <b>2011</b> , 32, 1569-77	4.6	77
17	Epitranscriptomics: regulation of mRNA metabolism through modifications. <i>Current Opinion in Chemical Biology</i> , <b>2017</b> , 41, 93-98	9.7	73
16	2VO-methylation in mRNA disrupts tRNA decoding during translation elongation. <i>Nature Structural and Molecular Biology</i> , <b>2018</b> , 25, 208-216	17.6	61
15	High-Resolution N6-Methyladenosine (m6A) Map Using Photo-Crosslinking-Assisted m6A Sequencing. <i>Angewandte Chemie</i> , <b>2015</b> , 127, 1607-1610	3.6	26
14	The Epitranscriptome in Translation Regulation. Cold Spring Harbor Perspectives in Biology, 2019, 11,	10.2	23
13	Genomics and Proteomics. Roadmap to the epitranscriptome. <i>Science</i> , <b>2014</b> , 346, 1192	33.3	21
12	Transcriptome-Wide Mapping of NEMethyladenosine by mA-Seq. <i>Methods in Enzymology</i> , <b>2015</b> , 560, 131-47	1.7	16
11	5-methylcytosine mediates nuclear export of mRNA. <i>Cell Research</i> , <b>2017</b> , 27, 717-719	24.7	14

## LIST OF PUBLICATIONS

10	Loud and Clear Epitranscriptomic mA Signals: Now in Single-Base Resolution. <i>Molecular Cell</i> , <b>2017</b> , 68, 825-826	17.6	12	
9	N-acetylation of Cytidine in mRNA by NAT10 Regulates Stability and Translation. <i>Cell</i> , <b>2018</b> , 175, 1725	-1 <i>7</i> ,2672	12	
8	Single base resolution mapping of 2VO-methylation sites in human mRNA and in 3Vterminal ends of small RNAs. <i>Methods</i> , <b>2019</b> , 156, 85-90	4.6	10	
7	Deep and accurate detection of m6A RNA modifications using miCLIP2 and m6Aboost machine learning. <i>Nucleic Acids Research</i> , <b>2021</b> , 49, e92	20.1	8	
6	N-methyladenosinelmethylation in tRNA drives liver tumourigenesis by regulating cholesterol metabolism. <i>Nature Communications</i> , <b>2021</b> , 12, 6314	17.4	6	
5	Micro-editing mistake translates into a devastating brain tumor. <i>Journal of Clinical Investigation</i> , <b>2012</b> , 122, 3842-5	15.9	5	
4	Identification of a homozygous VRK1 mutation in two patients with adult-onset distal hereditary motor neuropathy. <i>Muscle and Nerve</i> , <b>2020</b> , 61, 395-400	3.4	4	
3	The epitranscriptome toolbox <i>Cell</i> , <b>2022</b> , 185, 764-776	56.2	4	
2	A single center experience with publicly funded clinical exome sequencing for neurodevelopmental disorders or multiple congenital anomalies. <i>Scientific Reports</i> , <b>2021</b> , 11, 19099	4.9	2	
1	Early and late manifestations of neuropathy due to HSPB1 mutation in the Jewish Iranian population. <i>Annals of Clinical and Translational Neurology</i> , <b>2021</b> , 8, 1260-1268	5.3	O	