

Zhike Lu

List of Publications by Citations

Source: <https://exaly.com/author-pdf/11802055/zhike-lu-publications-by-citations.pdf>

Version: 2024-04-20

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.

41
papers

15,792
citations

35
h-index

48
g-index

48
ext. papers

21,203
ext. citations

21.4
avg, IF

6.29
L-index

#	Paper	IF	Citations
41	N6-methyladenosine-dependent regulation of messenger RNA stability. <i>Nature</i> , 2014 , 505, 117-20	50.4	1949
40	ALKBH5 is a mammalian RNA demethylase that impacts RNA metabolism and mouse fertility. <i>Molecular Cell</i> , 2013 , 49, 18-29	17.6	1627
39	N(6)-methyladenosine Modulates Messenger RNA Translation Efficiency. <i>Cell</i> , 2015 , 161, 1388-99	56.2	1493
38	A METTL3-METTL14 complex mediates mammalian nuclear RNA N6-adenosine methylation. <i>Nature Chemical Biology</i> , 2014 , 10, 93-5	11.7	1458
37	Identification of 67 histone marks and histone lysine crotonylation as a new type of histone modification. <i>Cell</i> , 2011 , 146, 1016-28	56.2	1150
36	mA Demethylase ALKBH5 Maintains Tumorigenicity of Glioblastoma Stem-like Cells by Sustaining FOXM1 Expression and Cell Proliferation Program. <i>Cancer Cell</i> , 2017 , 31, 591-606.e6	24.3	734
35	YTHDF3 facilitates translation and decay of N-methyladenosine-modified RNA. <i>Cell Research</i> , 2017 , 27, 315-328	24.7	696
34	mA RNA Methylation Regulates the Self-Renewal and Tumorigenesis of Glioblastoma Stem Cells. <i>Cell Reports</i> , 2017 , 18, 2622-2634	10.6	656
33	METTL14 Inhibits Hematopoietic Stem/Progenitor Differentiation and Promotes Leukemogenesis via mRNA mA Modification. <i>Cell Stem Cell</i> , 2018 , 22, 191-205.e9	18	476
32	RNA mA methylation regulates the ultraviolet-induced DNA damage response. <i>Nature</i> , 2017 , 543, 573-576.e4	56.4	449
31	Ythdc2 is an N-methyladenosine binding protein that regulates mammalian spermatogenesis. <i>Cell Research</i> , 2017 , 27, 1115-1127	24.7	404
30	Structural basis for selective binding of m6A RNA by the YTHDC1 YTH domain. <i>Nature Chemical Biology</i> , 2014 , 10, 927-9	11.7	383
29	mA mRNA methylation regulates AKT activity to promote the proliferation and tumorigenicity of endometrial cancer. <i>Nature Cell Biology</i> , 2018 , 20, 1074-1083	23.4	358
28	mA-dependent maternal mRNA clearance facilitates zebrafish maternal-to-zygotic transition. <i>Nature</i> , 2017 , 542, 475-478	50.4	293
27	Differential mA, m6A, and m6A Demethylation Mediated by FTO in the Cell Nucleus and Cytoplasm. <i>Molecular Cell</i> , 2018 , 71, 973-985.e5	17.6	289
26	N-methyladenosine (mA) recruits and repels proteins to regulate mRNA homeostasis. <i>Nature Structural and Molecular Biology</i> , 2017 , 24, 870-878	17.6	261
25	Lysine 2-hydroxyisobutyrylation is a widely distributed active histone mark. <i>Nature Chemical Biology</i> , 2014 , 10, 365-70	11.7	259

24	mA mRNA demethylase FTO regulates melanoma tumorigenicity and response to anti-PD-1 blockade. <i>Nature Communications</i> , 2019 , 10, 2782	17.4	254
23	High-resolution N(6) -methyladenosine (m(6) A) map using photo-crosslinking-assisted m(6) A sequencing. <i>Angewandte Chemie - International Edition</i> , 2015 , 54, 1587-90	16.4	249
22	Unique features of the m6A methylome in Arabidopsis thaliana. <i>Nature Communications</i> , 2014 , 5, 5630	17.4	239
21	Histone H3 trimethylation at lysine 36 guides mA RNA modification co-transcriptionally. <i>Nature</i> , 2019 , 567, 414-419	50.4	232
20	Dynamics of Human and Viral RNA Methylation during Zika Virus Infection. <i>Cell Host and Microbe</i> , 2016 , 20, 666-673	23.4	221
19	mA facilitates hippocampus-dependent learning and memory through YTHDF1. <i>Nature</i> , 2018 , 563, 249-253	33.4	208
18	Mettl3-/Mettl14-mediated mRNA N-methyladenosine modulates murine spermatogenesis. <i>Cell Research</i> , 2017 , 27, 1216-1230	24.7	171
17	N(6)-methyladenosine of HIV-1 RNA regulates viral infection and HIV-1 Gag protein expression. <i>ELife</i> , 2016 , 5,	8.9	167
16	NMethyladenosine methyltransferase ZCCHC4 mediates ribosomal RNA methylation. <i>Nature Chemical Biology</i> , 2019 , 15, 88-94	11.7	149
15	Ythdf2-mediated mA mRNA clearance modulates neural development in mice. <i>Genome Biology</i> , 2018 , 19, 69	18.3	129
14	ALKBH10B Is an RNA -Methyladenosine Demethylase Affecting Arabidopsis Floral Transition. <i>Plant Cell</i> , 2017 , 29, 2995-3011	11.6	124
13	The mA Reader ECT2 Controls Trichome Morphology by Affecting mRNA Stability in Arabidopsis. <i>Plant Cell</i> , 2018 , 30, 968-985	11.6	121
12	YTHDF2 reduction fuels inflammation and vascular abnormalization in hepatocellular carcinoma. <i>Molecular Cancer</i> , 2019 , 18, 163	42.1	113
11	Transfer RNA demethylase ALKBH3 promotes cancer progression via induction of tRNA-derived small RNAs. <i>Nucleic Acids Research</i> , 2019 , 47, 2533-2545	20.1	108
10	-methyladenosine RNA modification-mediated cellular metabolism rewiring inhibits viral replication. <i>Science</i> , 2019 , 365, 1171-1176	33.3	78
9	Mettl14 Is Essential for Epitranscriptomic Regulation of Striatal Function and Learning. <i>Neuron</i> , 2018 , 99, 283-292.e5	13.9	71
8	A metabolic labeling method detects mA transcriptome-wide at single base resolution. <i>Nature Chemical Biology</i> , 2020 , 16, 887-895	11.7	70
7	The RNA-binding protein FMRP facilitates the nuclear export of -methyladenosine-containing mRNAs. <i>Journal of Biological Chemistry</i> , 2019 , 294, 19889-19895	5.4	41

6	Viral N-methyladenosine upregulates replication and pathogenesis of human respiratory syncytial virus. <i>Nature Communications</i> , 2019 , 10, 4595	17.4	35
5	Steady-state hydrogen peroxide induces glycolysis in <i>Staphylococcus aureus</i> and <i>Pseudomonas aeruginosa</i> . <i>Journal of Bacteriology</i> , 2014 , 196, 2499-513	3.5	26
4	High-Resolution N6-Methyladenosine (m6A) Map Using Photo-Crosslinking-Assisted m6A Sequencing. <i>Angewandte Chemie</i> , 2015 , 127, 1607-1610	3.6	26
3	Direct-seq: a programmed gRNA scaffold for streamlined scRNA-seq in CRISPR screen. <i>Genome Biology</i> , 2020 , 21, 136	18.3	3
2	New Chromatin Run-On Reaction Enables Global Mapping of Active RNA Polymerase Locations in an Enrichment-free Manner.. <i>ACS Chemical Biology</i> , 2022 ,	4.9	1
1	m6A facilitates hippocampus-dependent learning and memory through Ythdf1. <i>FASEB Journal</i> , 2018 , 32, 787.6	0.9	0