

Boxuan Simen Zhao

List of Publications by Year in Descending Order

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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.

33
papers

7,271
citations

25
h-index

38
g-index

38
ext. papers

9,942
ext. citations

20.2
avg, IF

6.17
L-index

#	Paper	IF	Citations
33	DNA 5-Methylcytosine-Specific Amplification and Sequencing. <i>Journal of the American Chemical Society</i> , 2020 , 142, 4539-4543	16.4	8
32	N-methyladenosine modification enables viral RNA to escape recognition by RNA sensor RIG-I. <i>Nature Microbiology</i> , 2020 , 5, 584-598	26.6	91
31	RNA-protein interaction mapping via MS2- or Cas13-based APEX targeting. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 22068-22079	11.5	42
30	Histone H3 trimethylation at lysine 36 guides m ⁶ A RNA modification co-transcriptionally. <i>Nature</i> , 2019 , 567, 414-419	50.4	232
29	Viral N-methyladenosine upregulates replication and pathogenesis of human respiratory syncytial virus. <i>Nature Communications</i> , 2019 , 10, 4595	17.4	35
28	Recognition of RNA N-methyladenosine by IGF2BP proteins enhances mRNA stability and translation. <i>Nature Cell Biology</i> , 2018 , 20, 285-295	23.4	795
27	METTL14 Inhibits Hematopoietic Stem/Progenitor Differentiation and Promotes Leukemogenesis via mRNA m ⁶ A Modification. <i>Cell Stem Cell</i> , 2018 , 22, 191-205.e9	18	476
26	Our views of dynamic m ⁶ A-methyladenosine RNA methylation. <i>Rna</i> , 2018 , 24, 268-272	5.8	35
25	Long genes linked to autism spectrum disorders harbor broad enhancer-like chromatin domains. <i>Genome Research</i> , 2018 , 28, 933-942	9.7	17
24	YTHDF3 facilitates translation and decay of N-methyladenosine-modified RNA. <i>Cell Research</i> , 2017 , 27, 315-328	24.7	696
23	m ⁶ A-dependent maternal mRNA clearance facilitates zebrafish maternal-to-zygotic transition. <i>Nature</i> , 2017 , 542, 475-478	50.4	293
22	m ⁶ A 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
21	Evolution of transcript modification by m ⁶ A-methyladenosine in primates. <i>Genome Research</i> , 2017 , 27, 385-397	9.7	34
20	"Gamete On" for m ⁶ A: YTHDF2 Exerts Essential Functions in Female Fertility. <i>Molecular Cell</i> , 2017 , 67, 903-905	17.6	17
19	Post-transcriptional gene regulation by mRNA modifications. <i>Nature Reviews Molecular Cell Biology</i> , 2017 , 18, 31-42	48.7	909
18	Quantifying mammalian genomic DNA hydroxymethylcytosine content using solid-state nanopores. <i>Scientific Reports</i> , 2016 , 6, 29565	4.9	26
17	Dynamics of Human and Viral RNA Methylation during Zika Virus Infection. <i>Cell Host and Microbe</i> , 2016 , 20, 666-673	23.4	221

16	Nucleic Acid Modifications in Regulation of Gene Expression. <i>Cell Chemical Biology</i> , 2016 , 23, 74-85	8.2	155
15	The N6-Adenine Methyltransferase METTL14 Plays an Oncogenic Role in Acute Myeloid Leukemia. <i>Blood</i> , 2016 , 128, 1536-1536	2.2	1
14	N(6)-methyladenosine of HIV-1 RNA regulates viral infection and HIV-1 Gag protein expression. <i>ELife</i> , 2016 , 5,	8.9	167
13	Fate by RNA methylation: m6A steers stem cell pluripotency. <i>Genome Biology</i> , 2015 , 16, 43	18.3	64
12	Pseudouridine in a new era of RNA modifications. <i>Cell Research</i> , 2015 , 25, 153-4	24.7	41
11	N(6)-methyladenosine Modulates Messenger RNA Translation Efficiency. <i>Cell</i> , 2015 , 161, 1388-99	56.2	1493
10	Base-resolution maps of 5-formylcytosine and 5-carboxylcytosine reveal genome-wide DNA demethylation dynamics. <i>Cell Research</i> , 2015 , 25, 386-9	24.7	64
9	TET family proteins: oxidation activity, interacting molecules, and functions in diseases. <i>Chemical Reviews</i> , 2015 , 115, 2225-39	68.1	75
8	The multiple antibiotic resistance regulator MarR is a copper sensor in Escherichia coli. <i>Nature Chemical Biology</i> , 2014 , 10, 21-8	11.7	95
7	5mC oxidation by Tet2 modulates enhancer activity and timing of transcriptome reprogramming during differentiation. <i>Molecular Cell</i> , 2014 , 56, 286-297	17.6	226
6	A highly sensitive and genetically encoded fluorescent reporter for ratiometric monitoring of quinones in living cells. <i>Chemical Communications</i> , 2013 , 49, 8027-9	5.8	2
5	Probing subcellular organic hydroperoxide formation via a genetically encoded ratiometric and reversible fluorescent indicator. <i>Integrative Biology (United Kingdom)</i> , 2013 , 5, 1485-9	3.7	5
4	A Selective Fluorescent Probe for Carbon Monoxide Imaging in Living Cells. <i>Angewandte Chemie</i> , 2012 , 124, 9790-9794	3.6	19
3	A selective fluorescent probe for carbon monoxide imaging in living cells. <i>Angewandte Chemie - International Edition</i> , 2012 , 51, 9652-6	16.4	117
2	A highly selective fluorescent probe for visualization of organic hydroperoxides in living cells. <i>Journal of the American Chemical Society</i> , 2010 , 132, 17065-7	16.4	51
1	RNA-protein interaction mapping via MS2 or Cas13-based APEX targeting		2