Chengguo Yao

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/2023618/publications.pdf

Version: 2024-02-01

14	770	1040056	1125743
papers	citations	9	13
		h-index	g-index

15	15	15	1037
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Specific Regulation of m6A by SRSF7 Promotes the Progression of Glioblastoma. Genomics, Proteomics and Bioinformatics, 2023, 21, 707-728.	6.9	16
2	CFIm25 regulates human stem cell function independently of its role in mRNA alternative polyadenylation. RNA Biology, 2022, 19, 686-702.	3.1	0
3	U1 snRNP telescripting: molecular mechanisms and beyond. RNA Biology, 2021, 18, 1512-1523.	3.1	11
4	A potential mechanism underlying U1 snRNP inhibition of the cleavage step of mRNA 3â€™ processing. Biochemical and Biophysical Research Communications, 2020, 530, 196-202.	2.1	10
5	Suboptimal RNAâ€“RNA interaction limits U1 snRNP inhibition of canonical mRNA 3â€™ processing. RNA Biology, 2019, 16, 1448-1460.	3.1	11
6	snoRNAs associate with mRNA 3â€² processing complex: New wine in old bottles. RNA Biology, 2018, 15, 194-197.	3.1	18
7	A snoRNA modulates mRNA 3â€² end processing and regulates the expression of a subset of mRNAs. Nucleic Acids Research, 2017, 45, 8647-8660.	14.5	73
8	Coupling between alternative polyadenylation and alternative splicing is limited to terminal introns. RNA Biology, 2016, 13, 646-655.	3.1	34
9	CPSF30 and Wdr33 directly bind to AAUAAA in mammalian mRNA 3â€² processing. Genes and Development, 2014, 28, 2370-2380.	5.9	193
10	Fip1 regulates mRNA alternative polyadenylation to promote stem cell self-renewal. EMBO Journal, 2014, 33, 878-889.	7.8	136
11	Global and Quantitative Profiling of Polyadenylated RNAs Using PAS-seq. Methods in Molecular Biology, 2014, 1125, 179-185.	0.9	8
12	Global Proteinâ€“RNA Interaction Mapping at Single Nucleotide Resolution by iCLIP-Seq. Methods in Molecular Biology, 2014, 1126, 399-410.	0.9	7
13	Overlapping and distinct functions of CstF64 and CstF64Ï„ in mammalian mRNA 3â€² processing. Rna, 2013, 19, 1781-1790.	3.5	59
14	Transcriptome-wide analyses of CstF64â€“RNA interactions in global regulation of mRNA alternative polyadenylation. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 18773-18778.	7.1	194