

Liye Zhang

List of Publications by Year in descending order

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Version: 2024-02-01

25
papers

1,306
citations

687363

13
h-index

580821

25
g-index

28
all docs

28
docs citations

28
times ranked

2486
citing authors

#	ARTICLE	IF	CITATIONS
1	Targetable genetic features of primary testicular and primary central nervous system lymphomas. <i>Blood</i> , 2016, 127, 869-881.	1.4	429
2	A proximity-tagging system to identify membrane protein-protein interactions. <i>Nature Methods</i> , 2018, 15, 715-722.	19.0	148
3	A YAP/TAZ-Regulated Molecular Signature Is Associated with Oral Squamous Cell Carcinoma. <i>Molecular Cancer Research</i> , 2015, 13, 957-968.	3.4	107
4	Altered RNA editing in 3' UTR perturbs microRNA-mediated regulation of oncogenes and tumor-suppressors. <i>Scientific Reports</i> , 2016, 6, 23226.	3.3	77
5	Lexis and Grammar of Mitochondrial RNA Processing in Trypanosomes. <i>Trends in Parasitology</i> , 2020, 36, 337-355.	3.3	71
6	Glutamine-utilizing transaminases are a metabolic vulnerability of TAZ/YAP-activated cancer cells. <i>EMBO Reports</i> , 2018, 19, .	4.5	70
7	Diffuse large B-cell lymphoma patient-derived xenograft models capture the molecular and biological heterogeneity of the disease. <i>Blood</i> , 2016, 127, 2203-2213.	1.4	68
8	RNA Binding and Core Complexes Constitute the U-Insertion/Deletion Editosome. <i>Molecular and Cellular Biology</i> , 2014, 34, 4329-4342.	2.3	67
9	Thyroid Progenitors Are Robustly Derived from Embryonic Stem Cells through Transient, Developmental Stage-Specific Overexpression of Nkx2-1. <i>Stem Cell Reports</i> , 2017, 8, 216-225.	4.8	44
10	Antisense Transcripts Delimit Exonucleolytic Activity of the Mitochondrial 3' Processome to Generate Guide RNAs. <i>Molecular Cell</i> , 2016, 61, 364-378.	9.7	35
11	Oncogenic ALK regulates EMT in non-small cell lung carcinoma through repression of the epithelial splicing regulatory protein 1. <i>Oncotarget</i> , 2016, 7, 33316-33330.	1.8	35
12	<sc>PPR</sc> polyadenylation factor defines mitochondrial <sc>mRNA</sc> identity and stability in trypanosomes. <i>EMBO Journal</i> , 2017, 36, 2435-2454.	7.8	20
13	Transcription initiation defines kinetoplast RNA boundaries. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E10323-E10332.	7.1	19
14	Constructive edge of uridylation-induced RNA degradation. <i>RNA Biology</i> , 2016, 13, 1078-1083.	3.1	17
15	Protein tyrosine phosphatase receptor type R (PTPRR) antagonizes the Wnt signaling pathway in ovarian cancer by dephosphorylating and inactivating β -catenin. <i>Journal of Biological Chemistry</i> , 2019, 294, 18306-18323.	3.4	15
16	XAB2 depletion induces intron retention in POLR2A to impair global transcription and promote cellular senescence. <i>Nucleic Acids Research</i> , 2019, 47, 8239-8254.	14.5	15
17	Pentatricopeptide repeat poly(A) binding protein KPAF4 stabilizes mitochondrial mRNAs in <i>Trypanosoma brucei</i> . <i>Nature Communications</i> , 2019, 10, 146.	12.8	14
18	Induction of senescence-associated secretory phenotype underlies the therapeutic efficacy of PRC2 inhibition in cancer. <i>Cell Death and Disease</i> , 2022, 13, 155.	6.3	14

#	ARTICLE	IF	CITATIONS
19	Investigating RNA editing factors from trypanosome mitochondria. <i>Methods</i> , 2016, 107, 23-33.	3.8	10
20	Inhibition of the FACT Complex Targets Aberrant Hedgehog Signaling and Overcomes Resistance to Smoothed Antagonists. <i>Cancer Research</i> , 2021, 81, 3105-3120.	0.9	9
21	Poly(A) binding KPAF4/5 complex stabilizes kinetoplast mRNAs in <i>Trypanosoma brucei</i> . <i>Nucleic Acids Research</i> , 2020, 48, 8645-8662.	14.5	7
22	Diffuse Large B-Cell Lymphoma Patient-Derived Xenograft Models Capture Molecular and Biologic Heterogeneity and Inform Therapy. <i>Blood</i> , 2015, 126, 817-817.	1.4	5
23	Application of RNA subcellular fraction estimation method to explore RNA localization regulation. <i>G3: Genes, Genomes, Genetics</i> , 2022, 12, .	1.8	4
24	Actionable Genetic Features of Primary Testicular and Primary Central Nervous System Lymphomas. <i>Blood</i> , 2014, 124, 74-74.	1.4	2
25	CTS tag-based methods for investigating mitochondrial RNA modification factors in <i>Trypanosoma brucei</i> . <i>Methods in Enzymology</i> , 2021, 658, 83-109.	1.0	1