

Yu-Sheng Chen

List of Publications by Year in descending order

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26
papers

8,921
citations

394421

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526287

27
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docs citations

29
times ranked

8266
citing authors

#	ARTICLE	IF	CITATIONS
1	Dynamic DNA 5-Hydroxymethylcytosine and RNA 5-Methylcytosine Reprogramming During Early Human Development. <i>Genomics, Proteomics and Bioinformatics</i> , 2023, 21, 805-822.	6.9	1
2	Comprehensive analysis of RNA-seq and whole genome sequencing data reveals no evidence for SARS-CoV-2 integrating into host genome. <i>Protein and Cell</i> , 2022, 13, 379-385.	11.0	3
3	RNA 5-methylcytosine regulates YBX2-dependent liquid-liquid phase separation. <i>Fundamental Research</i> , 2022, 2, 48-55.	3.3	8
4	Differential transcriptomic landscapes of multiple organs from SARS-CoV-2 early infected rhesus macaques. <i>Protein and Cell</i> , 2022, 13, 920-939.	11.0	9
5	Aberrant APOBEC3C expression induces characteristic genomic instability in pancreatic ductal adenocarcinoma. <i>Oncogenesis</i> , 2022, 11, .	4.9	7
6	Dynamic transcriptomic ⁵ C and its regulatory role in RNA processing. <i>Wiley Interdisciplinary Reviews RNA</i> , 2021, 12, e1639.	6.4	101
7	N6-methyladenosine regulates RNA abundance of SARS-CoV-2. <i>Cell Discovery</i> , 2021, 7, 7.	6.7	7
8	Reorganized 3D Genome Structures Support Transcriptional Regulation in Mouse Spermatogenesis. <i>IScience</i> , 2020, 23, 101034.	4.1	36
9	5-methylcytosine promotes pathogenesis of bladder cancer through stabilizing mRNAs. <i>Nature Cell Biology</i> , 2019, 21, 978-990.	10.3	410
10	RNA 5-Methylcytosine Facilitates the Maternal-to-Zygotic Transition by Preventing Maternal mRNA Decay. <i>Molecular Cell</i> , 2019, 75, 1188-1202.e11.	9.7	242
11	Single-cell RNA-seq highlights intra-tumoral heterogeneity and malignant progression in pancreatic ductal adenocarcinoma. <i>Cell Research</i> , 2019, 29, 725-738.	12.0	661
12	m6A promotes R-loop formation to facilitate transcription termination. <i>Cell Research</i> , 2019, 29, 1035-1038.	12.0	101
13	Dynamic methylome of internal mRNA N7-methylguanosine and its regulatory role in translation. <i>Cell Research</i> , 2019, 29, 927-941.	12.0	154
14	Insight into novel RNA-binding activities via large-scale analysis of lncRNA-bound proteome and IDH1-bound transcriptome. <i>Nucleic Acids Research</i> , 2019, 47, 2244-2262.	14.5	29
15	An alternative CTCF isoform antagonizes canonical CTCF occupancy and changes chromatin architecture to promote apoptosis. <i>Nature Communications</i> , 2019, 10, 1535.	12.8	39
16	5-Methylcytosine Analysis by RNA-BisSeq. <i>Methods in Molecular Biology</i> , 2019, 1870, 237-248.	0.9	10
17	A novel m6A reader Prrc2a controls oligodendroglial specification and myelination. <i>Cell Research</i> , 2019, 29, 23-41.	12.0	250
18	Endothelial-specific m6A modulates mouse hematopoietic stem and progenitor cell development via Notch signaling. <i>Cell Research</i> , 2018, 28, 249-252.	12.0	84

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19	Dynamic transcriptomic m6A decoration: writers, erasers, readers and functions in RNA metabolism. Cell Research, 2018, 28, 616-624.	12.0	1,045
20	Cytoplasmic m6A reader YTHDF3 promotes mRNA translation. Cell Research, 2017, 27, 444-447.	12.0	606
21	5-methylcytosine promotes mRNA export â€” NSUN2 as the methyltransferase and ALYREF as an m5C reader. Cell Research, 2017, 27, 606-625.	12.0	666
22	Mettl3-mediated m6A regulates spermatogonial differentiation and meiosis initiation. Cell Research, 2017, 27, 1100-1114.	12.0	306
23	Nuclear m 6 A Reader YTHDC1 Regulates mRNA Splicing. Molecular Cell, 2016, 61, 507-519.	9.7	1,432
24	Smg6/Est1 licenses embryonic stem cell differentiation via nonsenseâ€”mediated <scp>mRNA</scp> decay. EMBO Journal, 2015, 34, 1630-1647.	7.8	108
25	FTO-dependent demethylation of N6-methyladenosine regulates mRNA splicing and is required for adipogenesis. Cell Research, 2014, 24, 1403-1419.	12.0	869
26	Mammalian WTAP is a regulatory subunit of the RNA N6-methyladenosine methyltransferase. Cell Research, 2014, 24, 177-189.	12.0	1,719