

# Yuanchao Zhang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2233365/publications.pdf>

Version: 2024-02-01

9  
papers

521  
citations

1307594

7  
h-index

1474206

9  
g-index

12  
all docs

12  
docs citations

12  
times ranked

886  
citing authors

#	ARTICLE	IF	CITATIONS
1	Comprehensive Multi-omics Analysis Reveals Mitochondrial Stress as a Central Biological Hub for Spaceflight Impact. <i>Cell</i> , 2020, 183, 1185-1201.e20.	28.9	161
2	Massively Systematic Transcript End Readout, "MASTER" Transcription Start Site Selection, Transcriptional Slippage, and Transcript Yields. <i>Molecular Cell</i> , 2015, 60, 953-965.	9.7	72
3	CapZyme-Seq Comprehensively Defines Promoter-Sequence Determinants for RNA 5' Capping with NAD+. <i>Molecular Cell</i> , 2018, 70, 553-564.e9.	9.7	64
4	Multiplexed protein-DNA cross-linking: Scrunching in transcription start site selection. <i>Science</i> , 2016, 351, 1090-1093.	12.6	62
5	The Pediatric Cell Atlas: Defining the Growth Phase of Human Development at Single-Cell Resolution. <i>Developmental Cell</i> , 2019, 49, 10-29.	7.0	57
6	Interactions between RNA polymerase and the core recognition element are a determinant of transcription start site selection. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E2899-905.	7.1	36
7	XACT-Seq Comprehensively Defines the Promoter-Position and Promoter-Sequence Determinants for Initial-Transcription Pausing. <i>Molecular Cell</i> , 2020, 79, 797-811.e8.	9.7	20
8	Scedar: A scalable Python package for single-cell RNA-seq exploratory data analysis. <i>PLoS Computational Biology</i> , 2020, 16, e1007794.	3.2	9
9	Promoter-sequence determinants and structural basis of primer-dependent transcription initiation in <i>Escherichia coli</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	4