

Jinyang Zhang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4867832/publications.pdf>

Version: 2024-02-01

11
papers

1,280
citations

1162889

8
h-index

1281743

11
g-index

12
all docs

12
docs citations

12
times ranked

1437
citing authors

#	ARTICLE	IF	CITATIONS
1	Circular RNA identification based on multiple seed matching. <i>Briefings in Bioinformatics</i> , 2018, 19, 803-810.	3.2	498
2	Comprehensive identification of internal structure and alternative splicing events in circular RNAs. <i>Nature Communications</i> , 2016, 7, 12060.	5.8	249
3	Expanded Expression Landscape and Prioritization of Circular RNAs in Mammals. <i>Cell Reports</i> , 2019, 26, 3444-3460.e5.	2.9	179
4	Accurate quantification of circular RNAs identifies extensive circular isoform switching events. <i>Nature Communications</i> , 2020, 11, 90.	5.8	140
5	Comprehensive profiling of circular RNAs with nanopore sequencing and CIRI-long. <i>Nature Biotechnology</i> , 2021, 39, 836-845.	9.4	108
6	circVAMP3 Drives CAPRIN1 Phase Separation and Inhibits Hepatocellular Carcinoma by Suppressing c-Myc Translation. <i>Advanced Science</i> , 2022, 9, e2103817.	5.6	38
7	Exploring the cellular landscape of circular RNAs using full-length single-cell RNA sequencing. <i>Nature Communications</i> , 2022, 13, .	5.8	25
8	The oral microbiome of pregnant women facilitates gestational diabetes discrimination. <i>Journal of Genetics and Genomics</i> , 2021, 48, 32-39.	1.7	16
9	Reconstruction of circular RNAs using Illumina and Nanopore RNA-seq datasets. <i>Methods</i> , 2021, 196, 17-22.	1.9	9
10	Characterizing Circular RNAs Using Nanopore Sequencing. <i>Trends in Biochemical Sciences</i> , 2021, 46, 785-786.	3.7	8
11	Screening Linear and Circular RNA Transcripts from Stress Granules. <i>Genomics, Proteomics and Bioinformatics</i> , 2023, 21, 886-893.	3.0	8