

Fei Qi

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9373144/publications.pdf>

Version: 2024-02-01

8
papers

241
citations

1306789

7
h-index

1588620

8
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8
all docs

8
docs citations

8
times ranked

321
citing authors

#	ARTICLE	IF	CITATIONS
1	Methanol-Independent Protein Expression by AOX1 Promoter with trans-Acting Elements Engineering and Glucose-Glycerol-Shift Induction in <i>Pichia pastoris</i> . <i>Scientific Reports</i> , 2017, 7, 41850.	1.6	88
2	A CRISPR/Cas13-based approach demonstrates biological relevance of vlinc class of long non-coding RNAs in anticancer drug response. <i>Scientific Reports</i> , 2020, 10, 1794.	1.6	49
3	Evolutionary analysis of polyproline motifs in <i>Escherichia coli</i> reveals their regulatory role in translation. <i>PLoS Computational Biology</i> , 2018, 14, e1005987.	1.5	31
4	Hovlinc is a recently evolved class of ribozyme found in human lincRNA. <i>Nature Chemical Biology</i> , 2021, 17, 601-607.	3.9	26
5	Transcriptome analysis of <i>mig1</i> mutant reveals their roles in methanol catabolism, peroxisome biogenesis and autophagy in methylotrophic yeast <i>Pichia pastoris</i> . <i>Genes and Genomics</i> , 2018, 40, 399-412.	0.5	15
6	Proline codon pair selection determines ribosome pausing strength and translation efficiency in bacteria. <i>Communications Biology</i> , 2021, 4, 589.	2.0	13
7	Melting temperature highlights functionally important RNA structure and sequence elements in yeast mRNA coding regions. <i>Nucleic Acids Research</i> , 2017, 45, 6109-6118.	6.5	12
8	Complex Age- and Cancer-Related Changes in Human Blood Transcriptome—Implications for Pan-Cancer Diagnostics. <i>Frontiers in Genetics</i> , 2021, 12, 746879.	1.1	7