

Huan Zhong

List of Publications by Year in descending order

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Version: 2024-02-01

11
papers

264
citations

1040056

9
h-index

1281871

11
g-index

14
all docs

14
docs citations

14
times ranked

353
citing authors

#	ARTICLE	IF	CITATIONS
1	Arabidopsis PUB2 and PUB4 connect signaling components of pattern-triggered immunity. <i>New Phytologist</i> , 2022, 233, 2249-2265.	7.3	17
2	SPAAC-NAD-seq, a sensitive and accurate method to profile NAD ⁺ -capped transcripts. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	26
3	Use of NAD tagSeq II to identify growth phase-dependent alterations in <i>E. coli</i> RNA NAD ⁺ capping. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	17
4	AtHDA6 functions as an H3K18ac eraser to maintain pericentromeric CHG methylation in <i>Arabidopsis thaliana</i> . <i>Nucleic Acids Research</i> , 2021, 49, 9755-9767.	14.5	6
5	<i>Arabidopsis</i> DXO1 possesses deNADding and exonuclease activities and its mutation affects defense-related and photosynthetic gene expression. <i>Journal of Integrative Plant Biology</i> , 2020, 62, 967-983.	8.5	29
6	NAD tagSeq for transcriptome-wide identification and characterization of NAD ⁺ -capped RNAs. <i>Nature Protocols</i> , 2020, 15, 2813-2836.	12.0	13
7	Retrospective analysis of LNM risk factors and the effect of chemotherapy in early colorectal cancer: A Chinese multicenter study. <i>BMC Cancer</i> , 2020, 20, 1067.	2.6	2
8	Redox-sensitive bZIP68 plays a role in balancing stress tolerance with growth in <i>Arabidopsis</i> . <i>Plant Journal</i> , 2019, 100, 768-783.	5.7	21
9	NAD tagSeq reveals that NAD ⁺ -capped RNAs are mostly produced from a large number of protein-coding genes in <i>Arabidopsis</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 12072-12077.	7.1	61
10	Bisphenol S induced epigenetic and transcriptional changes in human breast cancer cell line MCF-7. <i>Environmental Pollution</i> , 2019, 246, 697-703.	7.5	42
11	Predicting gene expression using DNA methylation in three human populations. <i>PeerJ</i> , 2019, 7, e6757.	2.0	28