

Zhaohui Zhong

List of Publications by Year in descending order

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Version: 2024-02-01

22
papers

2,268
citations

471061

17
h-index

642321

23
g-index

25
all docs

25
docs citations

25
times ranked

2049
citing authors

#	ARTICLE	IF	CITATIONS
1	Protoplast Isolation and Transfection in Rice. <i>Methods in Molecular Biology</i> , 2022, 2464, 83-90.	0.4	2
2	Genome-wide analyses of PAM-relaxed Cas9 genome editors reveal substantial off-target effects by ABE8e in rice. <i>Plant Biotechnology Journal</i> , 2022, 20, 1670-1682.	4.1	23
3	Improved plant cytosine base editors with high editing activity, purity, and specificity. <i>Plant Biotechnology Journal</i> , 2021, 19, 2052-2068.	4.1	55
4	PAM-less plant genome editing using a CRISPR-SpRY toolbox. <i>Nature Plants</i> , 2021, 7, 25-33.	4.7	140
5	The Improvement of CRISPR-Cas9 System With Ubiquitin-Associated Domain Fusion for Efficient Plant Genome Editing. <i>Frontiers in Plant Science</i> , 2020, 11, 621.	1.7	12
6	CRISPR-Cas12b enables efficient plant genome engineering. <i>Nature Plants</i> , 2020, 6, 202-208.	4.7	116
7	Knockout of the OsNAC006 Transcription Factor Causes Drought and Heat Sensitivity in Rice. <i>International Journal of Molecular Sciences</i> , 2020, 21, 2288.	1.8	69
8	Intron-Based Single Transcript Unit CRISPR Systems for Plant Genome Editing. <i>Rice</i> , 2020, 13, 8.	1.7	22
9	Bidirectional Promoter-Based CRISPR-Cas9 Systems for Plant Genome Editing. <i>Frontiers in Plant Science</i> , 2019, 10, 1173.	1.7	39
10	Improving Plant Genome Editing with High-Fidelity xCas9 and Non-canonical PAM-Targeting Cas9-NG. <i>Molecular Plant</i> , 2019, 12, 1027-1036.	3.9	159
11	Multiplex QTL editing of grain-related genes improves yield in elite rice varieties. <i>Plant Cell Reports</i> , 2019, 38, 475-485.	2.8	136
12	Single transcript unit <sc>CRISPR</sc> 2.0 systems for robust Cas9 and Cas12a mediated plant genome editing. <i>Plant Biotechnology Journal</i> , 2019, 17, 1431-1445.	4.1	120
13	A Single Transcript CRISPR-Cas9 System for Multiplex Genome Editing in Plants. <i>Methods in Molecular Biology</i> , 2019, 1917, 75-82.	0.4	3
14	Knocking Out MicroRNA Genes in Rice with CRISPR-Cas9. <i>Methods in Molecular Biology</i> , 2019, 1917, 109-119.	0.4	8
15	Robust Transcriptional Activation in Plants Using Multiplexed CRISPR-Act2.0 and mTALE-Act Systems. <i>Molecular Plant</i> , 2018, 11, 245-256.	3.9	179
16	Plant Genome Editing Using FnCpf1 and LbCpf1 Nucleases at Redefined and Altered PAM Sites. <i>Molecular Plant</i> , 2018, 11, 999-1002.	3.9	136
17	A large-scale whole-genome sequencing analysis reveals highly specific genome editing by both Cas9 and Cpf1 (Cas12a) nucleases in rice. <i>Genome Biology</i> , 2018, 19, 84.	3.8	230
18	CRISPRMatch: An Automatic Calculation and Visualization Tool for High-throughput CRISPR Genome-editing Data Analysis. <i>International Journal of Biological Sciences</i> , 2018, 14, 858-862.	2.6	53

#	ARTICLE	IF	CITATIONS
19	A CRISPR-Cpf1 system for efficient genome editing and transcriptional repression in plants. <i>Nature Plants</i> , 2017, 3, 17018.	4.7	425
20	CRISPR-Cas9 Based Genome Editing Reveals New Insights into MicroRNA Function and Regulation in Rice. <i>Frontiers in Plant Science</i> , 2017, 8, 1598.	1.7	150
21	Construction of a Single Transcriptional Unit for Expression of Cas9 and Single-guide RNAs for Genome Editing in Plants. <i>Bio-protocol</i> , 2017, 7, e2546.	0.2	2
22	Effective screen of CRISPR/Cas9-induced mutants in rice by single-strand conformation polymorphism. <i>Plant Cell Reports</i> , 2016, 35, 1545-1554.	2.8	74