

Namjin Koo

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11021564/publications.pdf>

Version: 2024-02-01

13
papers

573
citations

1163117

8
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1281871

11
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14
all docs

14
docs citations

14
times ranked

914
citing authors

#	ARTICLE	IF	CITATIONS
1	New reference genome sequences of hot pepper reveal the massive evolution of plant disease-resistance genes by retroduplication. <i>Genome Biology</i> , 2017, 18, 210.	8.8	255
2	Genome Sequence of <i>Striga asiatica</i> Provides Insight into the Evolution of Plant Parasitism. <i>Current Biology</i> , 2019, 29, 3041-3052.e4.	3.9	109
3	Transcriptome profiling of abiotic responses to heat, cold, salt, and osmotic stress of <i>Capsicum annum</i> L. <i>Scientific Data</i> , 2020, 7, 17.	5.3	76
4	Genome analysis of <i>Hibiscus syriacus</i> provides insights of polyploidization and indeterminate flowering in woody plants. <i>DNA Research</i> , 2017, 24, dsw049.	3.4	38
5	Transcriptome analysis of the oriental melon (<i>Cucumis melo</i> L. var. <i>makuwa</i>) during fruit development. <i>PeerJ</i> , 2017, 5, e2834.	2.0	31
6	TGFamã€Finder : a novel solution for targetã€gene family annotation in plants. <i>New Phytologist</i> , 2020, 227, 1568-1581.	7.3	23
7	Draft genome sequences of two oriental melons, <i>Cucumis melo</i> L. var. <i>makuwa</i> . <i>Scientific Data</i> , 2019, 6, 220.	5.3	13
8	Identification of transcriptome-wide, nut weight-associated SNPs in <i>Castanea crenata</i> . <i>Scientific Reports</i> , 2019, 9, 13161.	3.3	10
9	Universal gene co-expression network reveals receptor-like protein genes involved in broad-spectrum resistance in pepper (<i>Capsicum annum</i> L.). <i>Horticulture Research</i> , 2022, , .	6.3	10
10	Comprehensive analysis of Translationally Controlled Tumor Protein (TCTP) provides insights for lineage-specific evolution and functional divergence. <i>PLoS ONE</i> , 2020, 15, e0232029.	2.5	3
11	Admixture of divergent genomes facilitates hybridization across species in the family Brassicaceae. <i>New Phytologist</i> , 2022, 235, 743-758.	7.3	3
12	Prometheus, an omics portal for interkingdom comparative genomic analyses. <i>PLoS ONE</i> , 2020, 15, e0240191.	2.5	1
13	Inter-kingdom Comparative Analysis of Translationally Controlled Tumor Protein (TCTP) Provides Clues for Their Lineage-specific Evolution. , 2018, , .		0