Namjin Koo

List of Publications by Year in descending order

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

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		1163117	1281871	
13	573	8	11	
papers	citations	h-index	g-index	
14 all docs	14 docs citations	14 times ranked	914 citing authors	

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