Qin Qiao

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8179501/publications.pdf

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11	389	1040056	1281871
papers	citations	h-index	g-index
11	11	11	561
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Genome of <i>Crucihimalaya himalaica</i> , a close relative of <i>Arabidopsis</i> , shows ecological adaptation to high altitude. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 7137-7146.	7.1	108
2	Transcriptome sequencing of Crucihimalaya himalaica (Brassicaceae) reveals how Arabidopsis close relative adapt to the Qinghai-Tibet Plateau. Scientific Reports, 2016, 6, 21729.	3.3	47
3	Identification of Glutathione S-Transferase (GST) Genes from a Dark Septate Endophytic Fungus (Exophiala pisciphila) and Their Expression Patterns under Varied Metals Stress. PLoS ONE, 2015, 10, e0123418.	2.5	44
4	Reply to: Revisiting the origin of octoploid strawberry. Nature Genetics, 2020, 52, 5-7.	21.4	44
5	Evolutionary history and pan-genome dynamics of strawberry (<i>Fragaria </i> spp.). Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	43
6	Comparative Transcriptomics of Strawberries (Fragaria spp.) Provides Insights into Evolutionary Patterns. Frontiers in Plant Science, 2016, 7, 1839.	3.6	33
7	Genomic analysis of field pennycress (Thlaspi arvense) provides insights into mechanisms of adaptation to high elevation. BMC Biology, 2021, 19, 143.	3.8	23
8	Probing the transcriptome of Aconitum carmichaelii reveals the candidate genes associated with the biosynthesis of the toxic aconitine-type C19-diterpenoid alkaloids. Phytochemistry, 2018, 152, 113-124.	2.9	20
9	Enriched networks †nucleoside/nucleotide and ribonucleoside/ribonucleotide metabolic processes†and †response to stimulus†potentially conferred to drought adaptation of the epiphytic orchid Dendrobium wangliangii. Physiology and Molecular Biology of Plants, 2019, 25, 31-45.	3.1	13
10	Evaluation of genetic diversity and population structure of Fragaria nilgerrensis using EST-SSR markers. Gene, 2021, 796-797, 145791.	2.2	9
11	Dynamic Changes of DNA Methylation During Wild Strawberry (Fragaria nilgerrensis) Tissue Culture. Frontiers in Plant Science, 2021, 12, 765383.	3.6	5