

Zhong Zhang

List of Publications by Year in descending order

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Version: 2024-02-01

10
papers

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citations

1039406

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1372195

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all docs

10
docs citations

10
times ranked

391
citing authors

#	ARTICLE	IF	CITATIONS
1	The Jujube Genome Provides Insights into Genome Evolution and the Domestication of Sweetness/Acidity Taste in Fruit Trees. <i>PLoS Genetics</i> , 2016, 12, e1006433.	1.5	136
2	Genome-wide identification of AP2/ERF superfamily genes and their expression during fruit ripening of Chinese jujube. <i>Scientific Reports</i> , 2018, 8, 15612.	1.6	67
3	Comparative Analysis of Pigments, Phenolics, and Antioxidant Activity of Chinese Jujube (Ziziphus Tj ETQq1 1 0.784314 rgBT /Overlo	1.7	55
4	Construction of a high-density genetic map of Ziziphus jujuba Mill. using genotyping by sequencing technology. <i>Tree Genetics and Genomes</i> , 2016, 12, 1.	0.6	42
5	Transcript analyses of ethylene pathway genes during ripening of Chinese jujube fruit. <i>Journal of Plant Physiology</i> , 2018, 224-225, 1-10.	1.6	26
6	Jujube metabolome selection determined the edible properties acquired during domestication. <i>Plant Journal</i> , 2022, 109, 1116-1133.	2.8	25
7	Influences of Environmental Factors on Leaf Morphology of Chinese Jujubes. <i>PLoS ONE</i> , 2015, 10, e0127825.	1.1	16
8	High-Density Genetic Map Construction and QTL Mapping of Leaf and Needling Traits in Ziziphus jujuba Mill. <i>Frontiers in Plant Science</i> , 2019, 10, 1424.	1.7	13
9	Transcript analyses reveal a comprehensive role of abscisic acid in modulating fruit ripening in Chinese jujube. <i>BMC Plant Biology</i> , 2019, 19, 189.	1.6	10
10	Metabolite and Gene Expression Analysis Underlying Temporal and Spatial Accumulation of Pentacyclic Triterpenoids in Jujube. <i>Genes</i> , 2022, 13, 823.	1.0	8