

Wenbing Su

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

214
citations

1040056

9
h-index

1058476

14
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15
all docs

15
docs citations

15
times ranked

123
citing authors

#	ARTICLE	IF	CITATIONS
1	Integrated analysis of the metabolome, transcriptome and miRNome reveals crucial roles of auxin and heat shock proteins in the heat stress response of loquat fruit. <i>Scientia Horticulturae</i> , 2022, 294, 110764.	3.6	3
2	Integration of genomics, transcriptomics and metabolomics identifies candidate loci underlying fruit weight in loquat. <i>Horticulture Research</i> , 2022, .	6.3	12
3	Polyploidy underlies co-option and diversification of biosynthetic triterpene pathways in the apple tribe. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	37
4	EjBZR1 represses fruit enlargement by binding to the <i>EjCYP90</i> promoter in loquat. <i>Horticulture Research</i> , 2021, 8, 152.	6.3	18
5	EjFWLs are repressors of cell division during early fruit morphogenesis of loquat. <i>Scientia Horticulturae</i> , 2021, 287, 110261.	3.6	3
6	Gibberellin Induced Transcriptome Profiles Reveal Gene Regulation of Loquat Flowering. <i>Frontiers in Genetics</i> , 2021, 12, 703688.	2.3	4
7	Method for fast staining and obtaining high-magnification and high-resolution cell images of <i>Nicotiana benthamiana</i> . <i>Physiology and Molecular Biology of Plants</i> , 2021, 27, 181-188.	3.1	3
8	EjRAV1/2 Delay Flowering Through Transcriptional Repression of EjFTs and EjSOC1s in Loquat. <i>Frontiers in Plant Science</i> , 2021, 12, 816086.	3.6	5
9	EjTFL1 Genes Promote Growth but Inhibit Flower Bud Differentiation in Loquat. <i>Frontiers in Plant Science</i> , 2020, 11, 576.	3.6	18
10	The Role of EjSPL3, EjSPL4, EjSPL5, and EjSPL9 in Regulating Flowering in Loquat (<i>Eriobotrya japonica</i>) Tj ETQq0 0 0 rgBT /Overlock 10 T	4.4	22
11	The Role of EjSOC1s in Flower Initiation in <i>Eriobotrya japonica</i> . <i>Frontiers in Plant Science</i> , 2019, 10, 253.	3.6	26
12	Functional characterization of GI and CO homologs from <i>Eriobotrya deflexa</i> Nakai forma <i>koshunensis</i> . <i>Plant Cell Reports</i> , 2019, 38, 533-543.	5.6	16
13	Selection of the optimal reference genes for expression analyses in different materials of <i>Eriobotrya japonica</i> . <i>Plant Methods</i> , 2019, 15, 7.	4.3	18
14	Differential gene expression between the vigorous and dwarf litchi cultivars based on RNA-Seq transcriptome analysis. <i>PLoS ONE</i> , 2018, 13, e0208771.	2.5	12
15	The cellular physiology of loquat (<i>Eriobotrya japonica</i> Lindl.) fruit with a focus on how cell division and cell expansion processes contribute to pome morphogenesis. <i>Scientia Horticulturae</i> , 2017, 224, 142-149.	3.6	17