

Qiu-ping Zhang

List of Publications by Year in descending order

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Version: 2024-02-01

11
papers

150
citations

1478505

6
h-index

1281871

11
g-index

12
all docs

12
docs citations

12
times ranked

127
citing authors

#	ARTICLE	IF	CITATIONS
1	Construction of a High-Density Genetic Map and Identification of Quantitative Trait Loci Linked to Fruit Quality Traits in Apricots Using Specific-Locus Amplified Fragment Sequencing. <i>Frontiers in Plant Science</i> , 2022, 13, 798700.	3.6	5
2	Genetic diversity analysis of Chinese plum (<i>Prunus salicina</i> L.) based on whole-genome resequencing. <i>Tree Genetics and Genomes</i> , 2021, 17, 1.	1.6	12
3	Phylogeography of <i>Prunus armeniaca</i> L. revealed by chloroplast DNA and nuclear ribosomal sequences. <i>Scientific Reports</i> , 2021, 11, 13623.	3.3	3
4	Frequent germplasm exchanges drive the high genetic diversity of Chinese-cultivated common apricot germplasm. <i>Horticulture Research</i> , 2021, 8, 215.	6.3	16
5	Genetic diversity, population structure, and relationships of apricot (<i>Prunus</i>) based on restriction site-associated DNA sequencing. <i>Horticulture Research</i> , 2020, 7, 69.	6.3	26
6	Construction of an SNP-based high-density genetic map for Japanese plum in a Chinese population using specific length fragment sequencing. <i>Tree Genetics and Genomes</i> , 2020, 16, 1.	1.6	7
7	Differential expression of genes encoding phenylpropanoid enzymes in an apricot cultivar (<i>Prunus</i>) Tj ETQq1 1 0.784314 rgBT ₉ /Overlook	1.9	
8	The genetic relationship and structure of some natural interspecific hybrids in <i>Prunus</i> subgenus <i>Prunophora</i> , based on nuclear and chloroplast simple sequence repeats. <i>Genetic Resources and Crop Evolution</i> , 2018, 65, 625-636.	1.6	8
9	Comparative transcriptome profiling and morphology provide insights into endocarp cleaving of apricot cultivar (<i>Prunus armeniaca</i> L.). <i>BMC Plant Biology</i> , 2017, 17, 72.	3.6	40
10	Allelic variation of simple sequence repeats markers linked to PPV resistance in Chinese apricot. <i>Zahradnictvi (Prague, Czech Republic: 1992)</i> , 2017, 44, 6-13.	0.9	1
11	Genetic diversity and relationships of common apricot (<i>Prunus armeniaca</i> L.) in China based on simple sequence repeat (SSR) markers. <i>Genetic Resources and Crop Evolution</i> , 2014, 61, 357-368.	1.6	23