

Zhaohe Yuan

List of Publications by Citations

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Version: 2024-04-20

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

37
papers

409
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46
ext. papers

561
ext. citations

2.9
avg, IF

3.71
L-index

#	Paper	IF	Citations
37	The pomegranate (<i>Punica granatum</i> L.) genome provides insights into fruit quality and ovule developmental biology. <i>Plant Biotechnology Journal</i> , 2018 , 16, 1363-1374	11.6	71
36	Population genetic diversity in Chinese pomegranate (<i>Punica granatum</i> L.) cultivars revealed by fluorescent-AFLP markers. <i>Journal of Genetics and Genomics</i> , 2007 , 34, 1061-71	4	61
35	Characterization and evaluation of major anthocyanins in pomegranate (<i>Punica granatum</i> L.) peel of different cultivars and their development phases. <i>European Food Research and Technology</i> , 2013 , 236, 109-117	3.4	42
34	Cloning and expression of anthocyanin biosynthetic genes in red and white pomegranate. <i>Journal of Plant Research</i> , 2015 , 128, 687-96	2.6	33
33	Genetic structure of <i>Malus sieversii</i> population from Xinjiang, China, revealed by SSR markers. <i>Journal of Genetics and Genomics</i> , 2007 , 34, 947-55	4	27
32	The Complete Chloroplast Genomes of and a Comparison with Other Species in Lythraceae. <i>International Journal of Molecular Sciences</i> , 2019 , 20,	6.3	25
31	Effects of Salt Stress on Growth, Photosynthesis, and Mineral Nutrients of 18 Pomegranate (<i>Punica granatum</i>) Cultivars. <i>Agronomy</i> , 2020 , 10, 27	3.6	20
30	Population genetic structure in apricot (<i>Prunus armeniaca</i> L.) cultivars revealed by fluorescent-AFLP markers in southern Xinjiang, China. <i>Journal of Genetics and Genomics</i> , 2007 , 34, 1037-47	4	18
29	Proteome comparison following self- and across-pollination in self-incompatible apricot (<i>Prunus armeniaca</i> L.). <i>Protein Journal</i> , 2006 , 25, 328-35	3.9	16
28	Land-plant Phylogenomic and Pomegranate Transcriptomic Analyses Reveal an Evolutionary Scenario of CYP75 Genes Subsequent to Whole Genome Duplications 2019 , 62, 48-60		9
27	Genome-wide identification and expression of YABBY genes family during flower development in <i>Punica granatum</i> L. <i>Gene</i> , 2020 , 752, 144784	3.8	9
26	PATTERNS OF PIGMENT CHANGES IN POMEGRANATE (<i>PUNICA GRANATUM</i> L.) PEEL DURING FRUIT RIPENING. <i>Acta Horticulturae</i> , 2015 , 83-89	0.3	8
25	Genome-Wide Identification of the NHX Gene Family in <i>Punica granatum</i> L. and Their Expressional Patterns under Salt Stress. <i>Agronomy</i> , 2021 , 11, 264	3.6	6
24	Genome-wide identification and expression analysis of the CLC gene family in pomegranate (<i>Punica granatum</i>) reveals its roles in salt resistance. <i>BMC Plant Biology</i> , 2020 , 20, 560	5.3	5
23	Transcriptomic Profiling of Pomegranate Provides Insights into Salt Tolerance. <i>Agronomy</i> , 2020 , 10, 44	3.6	5
22	Characterization and comparative analysis of the complete chloroplast genome sequence from SummitV <i>PeerJ</i> , 2019 , 7, e8210	3.1	4
21	Fruit Breeding in Regard to Color and Seed Hardness: A Genomic View from Pomegranate. <i>Agronomy</i> , 2020 , 10, 991	3.6	4

20	Genome-wide identification, gene cloning, subcellular location and expression analysis of SPL gene family in <i>P. granatum</i> L. <i>BMC Plant Biology</i> , 2021 , 21, 400	5.3	4
19	Genome-Wide Identification and Expression Analysis of MIKC-Type MADS-Box Gene Family in <i>Punica granatum</i> L.. <i>Agronomy</i> , 2020 , 10, 1197	3.6	3
18	Advances in Mechanisms and Omics Pertaining to Fruit Cracking in Horticultural Plants. <i>Agronomy</i> , 2021 , 11, 1045	3.6	3
17	Anthocyanins from Pomegranate (<i>Punica granatum</i> L.) and Their Role in Antioxidant Capacities in Vitro. <i>Chemistry and Biodiversity</i> , 2021 , 18, e2100399	2.5	3
16	The complete chloroplast genome sequence of <i>G. Don</i> (Rosaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 3671-3672	0.5	2
15	Genome-wide Identification and Expression Analysis of TALE Gene Family in Pomegranate (<i>Punica granatum</i> L.). <i>Agronomy</i> , 2020 , 10, 829	3.6	2
14	The complete chloroplast genome sequence of cv.. <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 3236-3237	0.5	2
13	POLLEN MORPHOLOGY OF POMEGRANATE (<i>PUNICA GRANATUM</i> L.) FROM DIFFERENT ECO-GEOGRAPHICAL POPULATIONS IN CHINA. <i>Acta Horticulturae</i> , 2015 , 269-277	0.3	2
12	Genome-Wide Identification and Evolutionary Analysis of AOMT Gene Family in Pomegranate (<i>Punica granatum</i>). <i>Agronomy</i> , 2021 , 11, 318	3.6	2
11	Characterization of complete chloroplast genome of <i>L.</i> <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 2357-2358	0.5	1
10	The complete chloroplast genome sequence of <i>Ehrh. Ψissardii</i> (Rosaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 3744-3745	0.5	1
9	Flavonoid profiles in peels and arils of pomegranate cultivars. <i>Journal of Food Measurement and Characterization</i> , 2022 , 16, 880	2.8	1
8	Genome-wide identification and characterization of bZIP gene family and cloning of candidate genes for anthocyanin biosynthesis in pomegranate (<i>Punica granatum</i>).. <i>BMC Plant Biology</i> , 2022 , 22, 170	5.3	1
7	The complete chloroplast genome sequence of (<i>L.</i>) DC. Ψ leniflora(Rosaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 3723-3724	0.5	0
6	Identification and Function Analysis Provide Insights Into Flower Development of <i>L.</i> <i>Frontiers in Plant Science</i> , 2022 , 13, 833747	6.2	0
5	The complete chloroplast genome of apple rootstock Ψ 9V <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 2187-2188	0.5	0
4	Characterization of the complete chloroplast genome of var. xiaojinensis. <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 2487-2488	0.5	0
3	BIBLIOMETRIC ANALYSIS ON THE SITUATION AND TENDENCY OF POMEGRANATE RESEARCH IN THE WORLD. <i>Acta Horticulturae</i> , 2015 , 43-51	0.3	0

2	Genome-wide analysis of the family of light-harvesting chlorophyll a/b-binding proteins in pomegranate (<i>Punica granatum</i> L.). <i>Acta Horticulturae</i> , 2020 , 647-652	0.3
1	Genome-Wide Identification and Expression Analysis of MAPK and MAPKK Gene Family in Pomegranate (<i>Punica Granatum</i> L.). <i>Agronomy</i> , 2020 , 10, 1015	3.6