Zhaohe Yuan

List of Publications by Year in Descending Order

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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
citations

9
h-index

9
g-index

46
ext. papers

2.9
avg, IF

L-index

#	Paper	IF	Citations
37	Flavonoid profiles in peels and arils of pomegranate cultivars. <i>Journal of Food Measurement and Characterization</i> , 2022 , 16, 880	2.8	1
36	Identification and Function Analysis Provide Insights Into Flower Development of L <i>Frontiers in Plant Science</i> , 2022 , 13, 833747	6.2	O
35	Genome-wide identification and characterization of bZIP gene family and cloning of candidate genes for anthocyanin biosynthesis in pomegranate (Punica granatum) <i>BMC Plant Biology</i> , 2022 , 22, 170	5.3	1
34	Advances in Mechanisms and Omics Pertaining to Fruit Cracking in Horticultural Plants. <i>Agronomy</i> , 2021 , 11, 1045	3.6	3
33	Genome-Wide Identification of the NHX Gene Family in Punica granatum L. and Their Expressional Patterns under Salt Stress. <i>Agronomy</i> , 2021 , 11, 264	3.6	6
32	Genome-Wide Identification and Evolutionary Analysis of AOMT Gene Family in Pomegranate (Punica granatum). <i>Agronomy</i> , 2021 , 11, 318	3.6	2
31	Genome-wide identification, gene cloning, subcellular location and expression analysis of SPL gene family in P. granatum L. <i>BMC Plant Biology</i> , 2021 , 21, 400	5.3	4
30	Anthocyanins from Pomegranate (Punica granatum L.) and Their Role in Antioxidant Capacities in Vitro. <i>Chemistry and Biodiversity</i> , 2021 , 18, e2100399	2.5	3
29	Genome-wide identification and expression analysis of the CLC gene family in pomegranate (Punica granatum) reveals its roles in salt resistance. <i>BMC Plant Biology</i> , 2020 , 20, 560	5.3	5
28	Genome-wide Identification and Expression Analysis of TALE Gene Family in Pomegranate (Punica granatum L.). <i>Agronomy</i> , 2020 , 10, 829	3.6	2
27	Transcriptomic Profiling of Pomegranate Provides Insights into Salt Tolerance. <i>Agronomy</i> , 2020 , 10, 44	3.6	5
26	Effects of Salt Stress on Growth, Photosynthesis, and Mineral Nutrients of 18 Pomegranate (Punica granatum) Cultivars. <i>Agronomy</i> , 2020 , 10, 27	3.6	20
25	Genome-wide analysis of the family of light-harvesting chlorophyll a/b-binding proteins in pomegranate (Punica granatum L.). <i>Acta Horticulturae</i> , 2020 , 647-652	0.3	
24	Genome-wide identification and expression of YABBY genes family during flower development in Punica granatum L. <i>Gene</i> , 2020 , 752, 144784	3.8	9
23	Fruit Breeding in Regard to Color and Seed Hardness: A Genomic View from Pomegranate. <i>Agronomy</i> , 2020 , 10, 991	3.6	4
22	Genome-Wide Identification and Expression Analysis of MAPK and MAPKK Gene Family in Pomegranate (Punica Granatum L.). <i>Agronomy</i> , 2020 , 10, 1015	3.6	
21	Genome-Wide Identification and Expression Analysis of MIKC-Type MADS-Box Gene Family in Punica granatum L <i>Agronomy</i> , 2020 , 10, 1197	3.6	3

(2007-2019)

20	The complete chloroplast genome sequence of G. Don (Rosaceae). <i>Mitochondrial DNA Part B:</i> Resources, 2019 , 4, 3671-3672	0.5	2
19	The complete chloroplast genome of apple rootstock W9V Mitochondrial DNA Part B: Resources, 2019, 4, 2187-2188	0.5	
18	Characterization of complete chloroplast genome of L. <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 2357-2358	0.5	1
17	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
16	Land-plant Phylogenomic and Pomegranate Transcriptomic Analyses Reveal an Evolutionary Scenario of CYP75 Genes Subsequent to Whole Genome Duplications 2019 , 62, 48-60		9
15	Characterization of the complete chloroplast genome of var. xiaojinensis. <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 2487-2488	0.5	
14	The complete chloroplast genome sequence of Ehrh. PissardiiV(Rosaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 3744-3745	0.5	1
13	The complete chloroplast genome sequence of cv Mitochondrial DNA Part B: Resources, 2019, 4, 3236-3	32357	2
12	Characterization and comparative analysis of the complete chloroplast genome sequence from <code>\u00e4ummitVPeerJ</code> , 2019, 7, e8210	3.1	4
11	The complete chloroplast genome sequence of (L.) DC. \(\psi\)leniflora\(\psi\)Rosaceae). \(Mitochondrial DNA \) Part B: Resources, \(2019 \), 4, 3723-3724	0.5	O
10	The pomegranate (Punica granatum L.) genome provides insights into fruit quality and ovule developmental biology. <i>Plant Biotechnology Journal</i> , 2018 , 16, 1363-1374	11.6	71
9	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
8	BIBLIOMETRIC ANALYSIS ON THE SITUATION AND TENDENCY OF POMEGRANATE RESEARCH IN THE WORLD. <i>Acta Horticulturae</i> , 2015 , 43-51	0.3	
7	PATTERNS OF PIGMENT CHANGES IN POMEGRANATE (PUNICA GRANATUM L.) PEEL DURING FRUIT RIPENING. <i>Acta Horticulturae</i> , 2015 , 83-89	0.3	8
6	POLLEN MORPHOLOGY OF POMEGRANATE (PUNICA GRANATUM L.) FROM DIFFERENT ECO-GEOGRAPHICAL POPULATIONS IN CHINA. <i>Acta Horticulturae</i> , 2015 , 269-277	0.3	2
5	Characterization and evaluation of major anthocyanins in pomegranate (Punica granatum L.) peel of different cultivars and their development phases. <i>European Food Research and Technology</i> , 2013 , 236, 109-117	3.4	42
4	Genetic structure of Malus sieversii population from Xinjiang, China, revealed by SSR markers. Journal of Genetics and Genomics, 2007 , 34, 947-55	4	27
3	Population genetic structure in apricot (Prunus armeniaca L.) cultivars revealed by fluorescent-AFLP markers in southern Xinjiang, China. <i>Journal of Genetics and Genomics</i> , 2007 , 34, 1037-	-47	18

Population genetic diversity in Chinese pomegranate (Punica granatum L.) cultivars revealed by fluorescent-AFLP markers. *Journal of Genetics and Genomics*, **2007**, 34, 1061-71

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Proteome comparison following self- and across-pollination in self-incompatible apricot (Prunus armeniaca L.). *Protein Journal*, **2006**, 25, 328-35

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