

Zhaodong Hao

List of Publications by Citations

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Version: 2024-04-26

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papers

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755
ext. citations

4.7
avg, IF

3.64
L-index

#	Paper	IF	Citations
33	Liriodendron genome sheds light on angiosperm phylogeny and species-pair differentiation. <i>Nature Plants</i> , 2019 , 5, 18-25	11.5	77
32	: drawing SVG graphics to visualize and map genome-wide data on the idiograms. <i>PeerJ Computer Science</i> , 2020 , 6, e251	2.7	73
31	The complete chloroplast genome sequence of the relict woody plant <i>Metasequoia glyptostroboides</i> Hu et Cheng. <i>Frontiers in Plant Science</i> , 2015 , 6, 447	6.2	61
30	The Transcriptional Landscape of Polyploid Wheats and Their Diploid Ancestors during Embryogenesis and Grain Development. <i>Plant Cell</i> , 2019 , 31, 2888-2911	11.6	25
29	Complete chloroplast genome sequence of a major economic species, <i>Ziziphus jujuba</i> (Rhamnaceae). <i>Current Genetics</i> , 2017 , 63, 117-129	2.9	20
28	Transcriptome analysis and metabolic profiling reveal the key role of carotenoids in the petal coloration of. <i>Horticulture Research</i> , 2020 , 7, 70	7.7	20
27	The Complete Chloroplast Genome Sequence of a Relict Conifer <i>Glyptostrobus pensilis</i> : Comparative Analysis and Insights into Dynamics of Chloroplast Genome Rearrangement in Cupressophytes and Pinaceae. <i>PLoS ONE</i> , 2016 , 11, e0161809	3.7	18
26	Comparative Analysis of the Chloroplast Genomic Information of <i>Cunninghamia lanceolata</i> (Lamb.) Hook with Sibling Species from the Genera <i>Cryptomeria</i> D. Don, <i>Taiwania</i> Hayata, and <i>Calocedrus Kurz</i> . <i>International Journal of Molecular Sciences</i> , 2016 , 17,	6.3	15
25	ICE-CBF-COR Signaling Cascade and Its Regulation in Plants Responding to Cold Stress.. <i>International Journal of Molecular Sciences</i> , 2022 , 23,	6.3	11
24	The role of β -aminobutyric acid in aluminum stress tolerance in a woody plant, <i>Liriodendron chinense</i> Tulipifera. <i>Horticulture Research</i> , 2021 , 8, 80	7.7	11
23	Phylogenetic studies and comparative chloroplast genome analyses elucidate the basal position of halophyte <i>Nitraria sibirica</i> (Nitrariaceae) in the Sapindales. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2018 , 29, 745-755	1.3	9
22	Genetic Diversity and Differentiation of Relict Plant <i>Liriodendron</i> Populations Based on 29 Novel EST-SSR Markers. <i>Forests</i> , 2019 , 10, 334	2.8	7
21	Characterization of the Gene Family and Its Role in Abiotic Stress Response. <i>Frontiers in Plant Science</i> , 2021 , 12, 641280	6.2	7
20	Peptide Hormone Genes Promote Primary Root Growth and Adventitious Root Formation. <i>Plants</i> , 2019 , 8,	4.5	7
19	Genome-wide identification of the <i>Liriodendron chinense</i> WRKY gene family and its diverse roles in response to multiple abiotic stress.. <i>BMC Plant Biology</i> , 2022 , 22, 25	5.3	5
18	Conserved, divergent and heterochronic gene expression during <i>Brachypodium</i> and <i>Arabidopsis</i> embryo development. <i>Plant Reproduction</i> , 2021 , 34, 207-224	3.9	5
17	Gibberellin Oxidase Gene Family in : Genome-Wide Identification and Gene Expression Analysis. <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	5

16	The Transcriptome of <i>Cunninghamia lanceolata</i> male/female cone reveal the association between MIKC MADS-box genes and reproductive organs development. <i>BMC Plant Biology</i> , 2020 , 20, 508	5.3	4
15	Morphological, phenological, and transcriptional analyses provide insight into the diverse flowering traits of a mutant of the relic woody plant <i>Liriodendron chinense</i> . <i>Horticulture Research</i> , 2021 , 8, 174	7.7	4
14	Genome Sequence and Comparative Analysis of Isolated from Leaves. <i>Phytopathology</i> , 2020 , 110, 1260-1269	3.8	2
13	RIdeogram: drawing SVG graphics to visualize and map genome-wide data on the idiograms		2
12	The <i>Liriodendron chinense</i> MKK2 Gene Enhances <i>Arabidopsis thaliana</i> Salt Resistance. <i>Forests</i> , 2020 , 11, 1160	2.8	2
11	Integrative analysis of transcriptome and proteome revealed nectary and nectar traits in the plant-pollinator interaction of <i>Nitraria tangutorum</i> Bobrov. <i>BMC Plant Biology</i> , 2021 , 21, 230	5.3	2
10	The Full-Length Transcriptome Sequencing and Identification of Na/H Antiporter Genes in Halophyte Bobrov. <i>Genes</i> , 2021 , 12,	4.2	2
9	The chloroplast genome of (<i>Maxim.</i>) A.N. Vassiljeva. <i>Mitochondrial DNA Part B: Resources</i> , 2018 , 3, 222-224	5	1
8	Transcriptome and proteome analysis suggest enhanced photosynthesis in tetraploid <i>Liriodendron sino-americanum</i> . <i>Tree Physiology</i> , 2021 , 41, 1953-1971	4.2	1
7	Identification of miR397a and Its Functional Characterization in Callus Growth and Development by Regulating Its Target in <i>Liriodendron</i> . <i>Forests</i> , 2021 , 12, 912	2.8	1
6	Overexpression of From Halophyte Plant Enhances Tolerance to Salt Stress in. <i>Frontiers in Plant Science</i> , 2021 , 12, 716855	6.2	1
5	Exploring the (<i>Lamb.</i>) Hook Genome by BAC Sequencing.. <i>Frontiers in Bioengineering and Biotechnology</i> , 2022 , 10, 854130	5.8	1
4	-Mediated Genetic Transformation of Embryogenic Callus in a Hybrid (). <i>Frontiers in Plant Science</i> , 2022 , 13, 802128	6.2	1
3	Molecular Cloning and Functional Characterization of the DELLA Gene Family in <i>Liriodendron</i> Hybrids. <i>Forests</i> , 2020 , 11, 1363	2.8	0
2	The complete chloroplast genome of (<i>Thunb.</i>) Sweet, a traditional Chinese medicinal plant. <i>Mitochondrial DNA Part B: Resources</i> , 2021 , 6, 851-852	0.5	0
1	PIN3 from <i>Liriodendron</i> May Function in Inflorescence Development and Root Elongation. <i>Forests</i> , 2022 , 13, 568	2.8	0