

# Zhaodong Hao

## 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.

33  
papers

403  
citations

10  
h-index

19  
g-index

38  
ext. papers

755  
ext. citations

4.7  
avg, IF

3.64  
L-index

#	Paper	IF	Citations
33	ICE-CBF-COR Signaling Cascade and Its Regulation in Plants Responding to Cold Stress.. <i>International Journal of Molecular Sciences</i> , <b>2022</b> , 23,	6.3	11
32	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> , <b>2022</b> , 22, 25	5.3	5
31	Exploring the (Lamb.) Hook Genome by BAC Sequencing.. <i>Frontiers in Bioengineering and Biotechnology</i> , <b>2022</b> , 10, 854130	5.8	1
30	-Mediated Genetic Transformation of Embryogenic Callus in a Hybrid (L).. <i>Frontiers in Plant Science</i> , <b>2022</b> , 13, 802128	6.2	1
29	PIN3 from <i>Liriodendron</i> May Function in Inflorescence Development and Root Elongation. <i>Forests</i> , <b>2022</b> , 13, 568	2.8	0
28	The complete chloroplast genome of (Thunb.) Sweet, a traditional Chinese medicinal plant. <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 851-852	0.5	0
27	The role of ̢-aminobutyric acid in aluminum stress tolerance in a woody plant, <i>Liriodendron chinense</i> Lulipifera. <i>Horticulture Research</i> , <b>2021</b> , 8, 80	7.7	11
26	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> , <b>2021</b> , 21, 230	5.3	2
25	Conserved, divergent and heterochronic gene expression during <i>Brachypodium</i> and <i>Arabidopsis</i> embryo development. <i>Plant Reproduction</i> , <b>2021</b> , 34, 207-224	3.9	5
24	The Full-Length Transcriptome Sequencing and Identification of Na/H Antiporter Genes in Halophyte Bobrov. <i>Genes</i> , <b>2021</b> , 12,	4.2	2
23	Gibberellin Oxidase Gene Family in : Genome-Wide Identification and Gene Expression Analysis. <i>International Journal of Molecular Sciences</i> , <b>2021</b> , 22,	6.3	5
22	Characterization of the Gene Family and Its Role in Abiotic Stress Response. <i>Frontiers in Plant Science</i> , <b>2021</b> , 12, 641280	6.2	7
21	Transcriptome and proteome analysis suggest enhanced photosynthesis in tetraploid <i>Liriodendron sino-americanum</i> . <i>Tree Physiology</i> , <b>2021</b> , 41, 1953-1971	4.2	1
20	Identification of miR397a and Its Functional Characterization in Callus Growth and Development by Regulating Its Target in <i>Liriodendron</i> . <i>Forests</i> , <b>2021</b> , 12, 912	2.8	1
19	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> , <b>2021</b> , 8, 174	7.7	4
18	Overexpression of From Halophyte Plant Enhances Tolerance to Salt Stress in. <i>Frontiers in Plant Science</i> , <b>2021</b> , 12, 716855	6.2	1
17	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> , <b>2020</b> , 20, 508	5.3	4

16	Genome Sequence and Comparative Analysis of Isolated from Leaves. <i>Phytopathology</i> , <b>2020</b> , 110, 1260-1269	2.9	2
15	Transcriptome analysis and metabolic profiling reveal the key role of carotenoids in the petal coloration of. <i>Horticulture Research</i> , <b>2020</b> , 7, 70	7.7	20
14	: drawing SVG graphics to visualize and map genome-wide data on the idiograms. <i>PeerJ Computer Science</i> , <b>2020</b> , 6, e251	2.7	73
13	The Liriodendron chinense MKK2 Gene Enhances Arabidopsis thaliana Salt Resistance. <i>Forests</i> , <b>2020</b> , 11, 1160	2.8	2
12	Molecular Cloning and Functional Characterization of the DELLA Gene Family in Liriodendron Hybrids. <i>Forests</i> , <b>2020</b> , 11, 1363	2.8	0
11	Genetic Diversity and Differentiation of Relict Plant Liriodendron Populations Based on 29 Novel EST-SSR Markers. <i>Forests</i> , <b>2019</b> , 10, 334	2.8	7
10	The Transcriptional Landscape of Polyploid Wheats and Their Diploid Ancestors during Embryogenesis and Grain Development. <i>Plant Cell</i> , <b>2019</b> , 31, 2888-2911	11.6	25
9	Peptide Hormone Genes Promote Primary Root Growth and Adventitious Root Formation. <i>Plants</i> , <b>2019</b> , 8,	4.5	7
8	Liriodendron genome sheds light on angiosperm phylogeny and species-pair differentiation. <i>Nature Plants</i> , <b>2019</b> , 5, 18-25	11.5	77
7	The chloroplast genome of (Maxim.) A.N. Vassiljeva. <i>Mitochondrial DNA Part B: Resources</i> , <b>2018</b> , 3, 222-224	4.5	1
6	Phylogenetic studies and comparative chloroplast genome analyses elucidate the basal position of halophyte Nitraria sibirica (Nitrariaceae) in the Sapindales. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , <b>2018</b> , 29, 745-755	1.3	9
5	Complete chloroplast genome sequence of a major economic species, Ziziphus jujuba (Rhamnaceae). <i>Current Genetics</i> , <b>2017</b> , 63, 117-129	2.9	20
4	Comparative Analysis of the Chloroplast Genomic Information of Cunninghamia lanceolata (Lamb.) Hook with Sibling Species from the Genera Cryptomeria D. Don, Taiwania Hayata, and Calocedrus Kurz. <i>International Journal of Molecular Sciences</i> , <b>2016</b> , 17,	6.3	15
3	The Complete Chloroplast Genome Sequence of a Relict Conifer Glyptostrobus pensilis: Comparative Analysis and Insights into Dynamics of Chloroplast Genome Rearrangement in Cupressophytes and Pinaceae. <i>PLoS ONE</i> , <b>2016</b> , 11, e0161809	3.7	18
2	The complete chloroplast genome sequence of the relict woody plant Metasequoia glyptostroboides Hu et Cheng. <i>Frontiers in Plant Science</i> , <b>2015</b> , 6, 447	6.2	61
1	Rldeogram: drawing SVG graphics to visualize and map genome-wide data on the idiograms		2