Yao Li

List of Publications by Year
in descending order

31	316	932766 10	940134 16
papers	citations	h-index	g-index

34	34	34	260
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Genome-wide analyses of introgression between two sympatric Asian oak species. Nature Ecology and Evolution, 2022, 6, 924-935.	3.4	32
2	Alternative splicing and its regulatory role in woody plants. Tree Physiology, 2020, 40, 1475-1486.	1.4	31
3	Phylogeography of the temperate tree species Quercus acutissima in China: Inferences from chloroplast DNA variations. Biochemical Systematics and Ecology, 2015, 63, 190-197.	0.6	30
4	Phylogenetic comparison of 5â€² splice site determination in central spliceosomal proteins of the U1â€70K gene family, in response to developmental cues and stress conditions. Plant Journal, 2020, 103, 357-378.	2.8	30
5	Landscape Features and Climatic Forces Shape the Genetic Structure and Evolutionary History of an Oak Species (Quercus chenii) in East China. Frontiers in Plant Science, 2019, 10, 1060.	1.7	26
6	Photosynthetic electron-transfer reactions in the gametophyte of Pteris multifida reveal the presence of allelopathic interference from the invasive plant species Bidens pilosa. Journal of Photochemistry and Photobiology B: Biology, 2016, 158, 81-88.	1.7	23
7	Atmospheric deposition of heavy metals in Wuxi, China: estimation based on native moss analysis. Environmental Monitoring and Assessment, 2016, 188, 360.	1.3	20
8	Ancient east-west divergence, recent admixture, and multiple marginal refugia shape genetic structure of a widespread oak species (Quercus acutissima) in China. Tree Genetics and Genomes, 2018, 14, 1.	0.6	19
9	Changes in gametophyte physiology of Pteris multifida induced by the leaf leachate treatment of the invasive Bidens pilosa. Environmental Science and Pollution Research, 2016, 23, 3578-3585.	2.7	16
10	Complete Chloroplast Genome Sequence of Chinese Lacquer Tree (Toxicodendron vernicifluum,) Tj ETQq0 0 0 rgBT /Overlock 10	0.9	15
11	The defense system for Bidens pilosa root exudate treatments in Pteris multifida gametophyte. Ecotoxicology and Environmental Safety, 2019, 173, 203-213.	2.9	13
12	Community characteristics of a subtropical evergreen broad-leaved forest in Huangshan, Anhui Province, East China. Biodiversity Science, 2016, 24, 875-887.	0.2	8
13	Influence of Pliocene and Pleistocene climates on hybridization patterns between two closely related oak species in China. Annals of Botany, 2022, 129, 231-245.	1.4	7
14	The complete chloroplast genome sequence of Toxicodendron succedaneum (Anacardiaceae). Mitochondrial DNA Part B: Resources, 2020, 5, 1956-1957.	0.2	6
15	Systematic characterization of the branch point binding protein, splicing factor 1, gene family in plant development and stress responses. BMC Plant Biology, 2020, 20, 379.	1.6	5
16	Molecular characterization of sawtooth oak (Quercus acutissima) germplasm based on randomly amplified polymorphic DNA. Plant Systematics and Evolution, 2013, 299, 1829-1837.	0.3	4
17	A Meta-Analysis Indicates Positive Correlation between Genetic Diversity and Species Diversity. Biology, 2021, 10, 1089.	1.3	4
18	Mismatch Between Specific and Genetic Diversity in an Evergreen Broadleaf Forest in Southeast China: A Study Case of 10.24 ha Forest Dynamics Plot of Huangshan. Frontiers in Plant Science, 2021, 12, 706006.	1.7	4

#	Article	IF	Citations
19	The complete plastid genome sequence of Quercus ciliaris (Fagaceae). Mitochondrial DNA Part B: Resources, 2020, 5, 1954-1955.	0.2	3
20	The complete chloroplast genome sequence of Toxicodendron sylvestre (Anacardiaceae). Mitochondrial DNA Part B: Resources, 2020, 5, 2008-2009.	0.2	3
21	Stepped Geomorphology Shaped the Phylogeographic Structure of a Widespread Tree Species (Toxicodendron vernicifluum, Anacardiaceae) in East Asia. Frontiers in Plant Science, 2022, 13, .	1.7	3
22	The complete chloroplast genome sequence of Quercus myrsinifolia (Fagaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 3139-3140.	0.2	2
23	The complete chloroplast genome sequence of Quercus stewardiana (Fagaceae). Mitochondrial DNA Part B: Resources, 2020, 5, 1958-1959.	0.2	2
24	The complete chloroplast genome sequence of Distylium macrophyllum (Hamamelidaceae). Mitochondrial DNA Part B: Resources, 2020, 5, 701-702.	0.2	2
25	The complete chloroplast genome sequence of Quercus sessilifolia Blume (Fagaceae). Mitochondrial DNA Part B: Resources, 2022, 7, 182-184.	0.2	2
26	Environmental Heterogeneity Affecting Community Assembly Patterns and Phylogenetic Diversity of Three Forest Communities at Mt. Huangshan, China. Forests, 2022, 13, 133.	0.9	2
27	Combining the Effects of Global Warming, Land Use Change and Dispersal Limitations to Predict the Future Distributions of East Asian Cerris Oaks (Quercus Section Cerris, Fagaceae) in China. Forests, 2022, 13, 367.	0.9	2
28	The complete chloroplast genome of Distylium tsiangii Chun ex Walker, a rare and endangered plant. Mitochondrial DNA Part B: Resources, 2020, 5, 430-431.	0.2	1
29	The complete chloroplast genome sequence of Gynandropsis gynandra (Cleomaceae). Mitochondrial DNA Part B: Resources, 2021, 6, 1909-1910.	0.2	1
30	Complete chloroplast genome sequence and phylogenetic analysis of Ilex viridis Champ. ex Benth. Mitochondrial DNA Part B: Resources, 2020, 5, 914-915.	0.2	0
31	The complete plastid genome sequence of Ilex micrococca Maxim. Mitochondrial DNA Part B: Resources, 2020, 5, 916-917.	0.2	0