

Yao Li

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.

25
papers

145
citations

8
h-index

11
g-index

34
ext. papers

234
ext. citations

3
avg, IF

2.94
L-index

#	Paper	IF	Citations
25	The complete chloroplast genome sequence of Blume (Fagaceae).. <i>Mitochondrial DNA Part B: Resources</i> , 2022 , 7, 182-184	0.5	
24	Environmental Heterogeneity Affecting Community Assembly Patterns and Phylogenetic Diversity of Three Forest Communities at Mt. Huangshan, China. <i>Forests</i> , 2022 , 13, 133	2.8	2
23	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. <i>Forests</i> , 2022 , 13, 367	2.8	0
22	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.. <i>Frontiers in Plant Science</i> , 2021 , 12, 706006	6.2	1
21	Influence of Pliocene and Pleistocene climates on hybridization patterns between two closely related oak species in China. <i>Annals of Botany</i> , 2021 ,	4.1	1
20	A Meta-Analysis Indicates Positive Correlation between Genetic Diversity and Species Diversity. <i>Biology</i> , 2021 , 10,	4.9	1
19	The complete chloroplast genome sequence of (Cleomaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2021 , 6, 1909-1910	0.5	0
18	Phylogenetic comparison of 5asplice site determination in central spliceosomal proteins of the U1-70K gene family, in response to developmental cues and stress conditions. <i>Plant Journal</i> , 2020 , 103, 357-378	6.9	8
17	Alternative splicing and its regulatory role in woody plants. <i>Tree Physiology</i> , 2020 , 40, 1475-1486	4.2	5
16	The complete chloroplast genome of Chun ex Walker, a rare and endangered plant. <i>Mitochondrial DNA Part B: Resources</i> , 2020 , 5, 430-431	0.5	1
15	The complete chloroplast genome sequence of (Hamamelidaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2020 , 5, 701-702	0.5	2
14	Complete chloroplast genome sequence and phylogenetic analysis of Champ. ex Benth. <i>Mitochondrial DNA Part B: Resources</i> , 2020 , 5, 914-915	0.5	
13	The complete plastid genome sequence of Maxim. <i>Mitochondrial DNA Part B: Resources</i> , 2020 , 5, 916-917	0.5	
12	Complete Chloroplast Genome Sequence of Chinese Lacquer Tree (, Anacardiaceae) and Its Phylogenetic Significance. <i>BioMed Research International</i> , 2020 , 2020, 9014873	3	9
11	Systematic characterization of the branch point binding protein, splicing factor 1, gene family in plant development and stress responses. <i>BMC Plant Biology</i> , 2020 , 20, 379	5.3	4
10	The complete chloroplast genome sequence of (Fagaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 3139-3140	0.5	0
9	Landscape Features and Climatic Forces Shape the Genetic Structure and Evolutionary History of an Oak Species () in East China. <i>Frontiers in Plant Science</i> , 2019 , 10, 1060	6.2	14

8	The defense system for <i>Bidens pilosa</i> root exudate treatments in <i>Pteris multifida</i> gametophyte. <i>Ecotoxicology and Environmental Safety</i> , 2019 , 173, 203-213	7	5
7	Ancient east-west divergence, recent admixture, and multiple marginal refugia shape genetic structure of a widespread oak species (<i>Quercus acutissima</i>) in China. <i>Tree Genetics and Genomes</i> , 2018 , 14, 1	2.1	14
6	Changes in gametophyte physiology of <i>Pteris multifida</i> induced by the leaf leachate treatment of the invasive <i>Bidens pilosa</i> . <i>Environmental Science and Pollution Research</i> , 2016 , 23, 3578-85	5.1	11
5	Photosynthetic electron-transfer reactions in the gametophyte of <i>Pteris multifida</i> reveal the presence of allelopathic interference from the invasive plant species <i>Bidens pilosa</i> . <i>Journal of Photochemistry and Photobiology B: Biology</i> , 2016 , 158, 81-8	6.7	12
4	Community characteristics of a subtropical evergreen broad-leaved forest in Huangshan, Anhui Province, East China. <i>Biodiversity Science</i> , 2016 , 24, 875-887	1.3	5
3	Atmospheric deposition of heavy metals in Wuxi, China: estimation based on native moss analysis. <i>Environmental Monitoring and Assessment</i> , 2016 , 188, 360	3.1	17
2	Phylogeography of the temperate tree species <i>Quercus acutissima</i> in China: Inferences from chloroplast DNA variations. <i>Biochemical Systematics and Ecology</i> , 2015 , 63, 190-197	1.4	21
1	Molecular characterization of sawtooth oak (<i>Quercus acutissima</i>) germplasm based on randomly amplified polymorphic DNA. <i>Plant Systematics and Evolution</i> , 2013 , 299, 1829-1837	1.3	4