Yao Li

List of Publications by Citations

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	Phylogeography of the temperate tree species Quercus acutissima in China: Inferences from chloroplast DNA variations. *Biochemical Systematics and Ecology*, **2015**, 63, 190-197	1.4	21
24	Atmospheric deposition of heavy metals in Wuxi, China: estimation based on native moss analysis. *Environmental Monitoring and Assessment*, **2016**, 188, 360	3.1	17
23	Landscape Features and Climatic Forces Shape the Genetic Structure and Evolutionary History of an Oak Species () in East China. *Frontiers in Plant Science*, **2019**, 10, 1060	6.2	14
22	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	2.1	14
21	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, 81-8	6.7	12
20	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, 3578-85	5.1	11
19	Complete Chloroplast Genome Sequence of Chinese Lacquer Tree (, Anacardiaceae) and Its Phylogenetic Significance. *BioMed Research International*, **2020**, 2020, 9014873	3	9
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. *Plant Journal*, **2020**, 103, 357-378	6.9	8
17	Alternative splicing and its regulatory role in woody plants. *Tree Physiology*, **2020**, 40, 1475-1486	4.2	5
16	Community characteristics of a subtropical evergreen broad-leaved forest in Huangshan, Anhui Province, East China. *Biodiversity Science*, **2016**, 24, 875-887	1.3	5
15	The defense system for Bidens pilosa root exudate treatments in Pteris multifida gametophyte. *Ecotoxicology and Environmental Safety*, **2019**, 173, 203-213	7	5
14	Molecular characterization of sawtooth oak (Quercus acutissima) germplasm based on randomly amplified polymorphic DNA. *Plant Systematics and Evolution*, **2013**, 299, 1829-1837	1.3	4
13	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	5.3	4
12	The complete chloroplast genome sequence of (Hamamelidaceae). *Mitochondrial DNA Part B: Resources*, **2020**, 5, 701-702	0.5	2
11	Environmental Heterogeneity Affecting Community Assembly Patterns and Phylogenetic Diversity of Three Forest Communities at Mt. Huangshan, China. *Forests*, **2022**, 13, 133	2.8	2
10	The complete chloroplast genome of Chun ex Walker, a rare and endangered plant. *Mitochondrial DNA Part B: Resources*, **2020**, 5, 430-431	0.5	1
9	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	6.2	1

List of Publications

8	Influence of Pliocene and Pleistocene climates on hybridization patterns between two closely related oak species in China. *Annals of Botany*, **2021**,	4.1	1
7	A Meta-Analysis Indicates Positive Correlation between Genetic Diversity and Species Diversity. *Biology*, **2021**, 10,	4.9	1
6	The complete chloroplast genome sequence of (Fagaceae). *Mitochondrial DNA Part B: Resources*, **2019**, 4, 3139-3140	0.5	0
5	The complete chloroplast genome sequence of (Cleomaceae). *Mitochondrial DNA Part B: Resources*, **2021**, 6, 1909-1910	0.5	0
4	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	2.8	0
3	Complete chloroplast genome sequence and phylogenetic analysis of Champ. ex Benth. *Mitochondrial DNA Part B: Resources*, **2020**, 5, 914-915	0.5	
2	The complete plastid genome sequence of Maxim. *Mitochondrial DNA Part B: Resources*, **2020**, 5, 916-917	0.5	
1	The complete chloroplast genome sequence of Blume (Fagaceae).. *Mitochondrial DNA Part B: Resources*, **2022**, 7, 182-184	0.5	