Qiao-Ping Xiang

List of Publications by Year
in descending order

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29	647	623734	580821
papers	citations	14 h-index	25 g-index

32	32	32	639
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Phylogeography of the Sino-Himalayan Fern Lepisorus clathratus on â€œThe Roof of the Worldâ€• PLoS ONE, 2011, 6, e25896.	2.5	72
2	Phylogenetic relationships, possible ancient hybridization, and biogeographic history of Abies (Pinaceae) based on data from nuclear, plastid, and mitochondrial genomes. Molecular Phylogenetics and Evolution, 2015, 82, 1-14.	2.7	72
3	Plastid Phylogenomics Resolve Deep Relationships among Eupolypod II Ferns with Rapid Radiation and Rate Heterogeneity. Genome Biology and Evolution, 2017, 9, 1646-1657.	2.5	67
4	Phylogeny of the paleotropical fern genus Lepisorus (Polypodiaceae, Polypodiopsida) inferred from four chloroplast DNA regions. Molecular Phylogenetics and Evolution, 2010, 54, 211-225.	2.7	59
5	Phylogeny of Abies (Pinaceae) inferred from nrITS sequence data. Taxon, 2009, 58, 141-152.	0.7	46
6	A molecular phylogeny and a revised classification of tribe Lepisoreae (Polypodiaceae) based on an analysis of four plastid DNA regions. Botanical Journal of the Linnean Society, 2010, 162, 28-38.	1.6	35
7	Phylogenetic relationships in Abies (Pinaceae): evidence from PCR-RFLP of the nuclear ribosomal DNA internal transcribed spacer region. Botanical Journal of the Linnean Society, 2004, 145, 425-435.	1.6	31
8	The Unique Evolutionary Trajectory and Dynamic Conformations of DR and IR/DR-Coexisting Plastomes of the Early Vascular Plant Selaginellaceae (Lycophyte). Genome Biology and Evolution, 2019, 11, 1258-1274.	2.5	26
9	Distinctive evolutionary pattern of organelle genomes linked to the nuclear genome in Selaginellaceae. Plant Journal, 2020, 104, 1657-1672.	5.7	26
10	Fire-prone Rhamnaceae with South African affinities in Cretaceous Myanmar amber. Nature Plants, 2022, 8, 125-135.	9.3	24
11	Phylogeny and Biogeography ofThujaL. (Cupressaceae), an Eastern Asian and North American Disjunct Genus. Journal of Integrative Plant Biology, 2005, 47, 651-659.	8.5	23
12	Species delimitation and phylogeography of theAbies chensiensiscomplex inferred from morphological and molecular data. Botanical Journal of the Linnean Society, 2015, 177, 175-188.	1.6	22
13	Plastome-based phylogenomics resolves the placement of the sanguinolenta group in the spikemoss of lycophyte (Selaginellaceae). Molecular Phylogenetics and Evolution, 2020, 147, 106788.	2.7	21
14	Elevation Shift in Abies Mill. (Pinaceae) of Subtropical and Temperate China and Vietnamâ€”Corroborative Evidence from Cytoplasmic DNA and Ecological Niche Modeling. Frontiers in Plant Science, 2017, 8, 578.	3.6	15
15	Backbone phylogeny of Lepisorus (Polypodiaceae) and a novel infrageneric classification based on the total evidence from plastid and morphological data. Cladistics, 2020, 36, 235-258.	3.3	15
16	A new species of Keteleeria (Pinaceae) in the Shanwang Miocene flora of China and its phytogeographic connection with North America. Taxon, 2006, 55, 165-171.	0.7	12
17	Directed Repeats Co-occur with Few Short-Dispersed Repeats in Plastid Genome of a Spikemoss, Selaginella vardei (Selaginellaceae, Lycopodiopsida). BMC Genomics, 2019, 20, 484.	2.8	12
18	New infrageneric classification of Abies in light of molecular phylogeny and high diversity in western North America. Journal of Systematics and Evolution, 2018, 56, 562-572.	3.1	11

#	Article	IF	Citations
19	Integrative taxonomy of the Selaginella helvetica group based on morphological, molecular and ecological data. Taxon, 2021, 70, 1163-1187.	0.7	10
20	Species delimitation and phylogeography of Abies delavayi complex: Inferred from morphological, molecular, and climatic data. Journal of Systematics and Evolution, 2020, 58, 234-246.	3.1	9
21	The evolution of extremely diverged plastomes in Selaginellaceae (lycophyte) is driven by repeat patterns and the underlying <scp>DNA</scp> maintenance machinery. Plant Journal, 2022, 111, 768-784.	5.7	8
22	Molecular Phylogeny of the Cliff Ferns (Woodsiaceae: Polypodiales) with a Proposed Infrageneric Classification. PLoS ONE, 2015, 10, e0136318.	2.5	7
23	Plastid phylogenomic analyses of the Selaginella sanguinolenta group (Selaginellaceae) reveal conflict signatures resulting from sequence types, outlier genes, and pervasive RNA editing. Molecular Phylogenetics and Evolution, 2022, 173, 107507.	2.7	7
24	Recognizing the species of Thuja (Cupressaceae) based on their cone and foliage morphology. Phytotaxa, 2015, 219, 101.	0.3	4
25	The origin of allotetraploid Lepisorus inaequibasis (Polypodiaceae) and paternal bias in its morphology and abiotic niche. Taxon, 2020, 69, 43-55.	0.7	4
26	Ellipinema andÂÃ—Ellipisorus? Just Lepisorus (Polypodiaceae)!. Molecular Phylogenetics and Evolution, 2021, 161, 107176.	2.7	4
27	Isolation and characterization of microsatellite markers in the Lepisorus clathratus complex (Polypodiaceae). Applications in Plant Sciences, 2016, 4, 1600069.	2.1	2
28	Correctability of the epithet minchegense to mingcheense in Lycopodium (Lycopodiaceae) and lectotypification of Lycopodium mingcheense. Taxon, 2003, 52, 857-858.	0.7	1
29	The complete chloroplast genome of the Abies yuanbaoshanensis, an endangered Pinaceae species in Southern China. Mitochondrial DNA Part B: Resources, 2019, 4, 3386-3387.	0.4	0