

Xing-Jin He

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8169203/publications.pdf>

Version: 2024-02-01

134
papers

4,654
citations

185998

28
h-index

110170

64
g-index

137
all docs

137
docs citations

137
times ranked

4867
citing authors

#	ARTICLE	IF	CITATIONS
1	RASP (Reconstruct Ancestral State in Phylogenies): A tool for historical biogeography. <i>Molecular Phylogenetics and Evolution</i> , 2015, 87, 46-49.	1.2	1,049
2	S-DIVA (Statistical Dispersal-Vicariance Analysis): A tool for inferring biogeographic histories. <i>Molecular Phylogenetics and Evolution</i> , 2010, 56, 848-850.	1.2	667
3	RASP 4: Ancestral State Reconstruction Tool for Multiple Genes and Characters. <i>Molecular Biology and Evolution</i> , 2020, 37, 604-606.	3.5	338
4	Purification, composition analysis and antioxidant activity of the polysaccharides from <i>Dendrobium nobile</i> Lindl.. <i>Carbohydrate Polymers</i> , 2010, 79, 1014-1019.	5.1	228
5	Phylogeny and biogeography of <i>Allium</i> (Amaryllidaceae: Allieae) based on nuclear ribosomal internal transcribed spacer and chloroplast rps16 sequences, focusing on the inclusion of species endemic to China. <i>Annals of Botany</i> , 2010, 106, 709-733.	1.4	170
6	Composition analysis and antioxidant activity of polysaccharide from <i>Dendrobium denneanum</i> . <i>International Journal of Biological Macromolecules</i> , 2009, 45, 169-173.	3.6	152
7	In vitro antioxidant activities of a water-soluble polysaccharide derived from <i>Dendrobium nobile</i> Lindl. extracts. <i>International Journal of Biological Macromolecules</i> , 2009, 45, 359-363.	3.6	119
8	A tool for the analysis of chromosomes: KaryoType. <i>Taxon</i> , 2016, 65, 586-592.	0.4	110
9	Evolutionary events in <i>Lilium</i> (including <i>Nomocharis</i> , Liliaceae) are temporally correlated with orogenies of the Qâ€™T plateau and the Hengduan Mountains. <i>Molecular Phylogenetics and Evolution</i> , 2013, 68, 443-460.	1.2	97
10	Comparative Analysis of the Chloroplast Genomes of the Chinese Endemic Genus <i>Urophysa</i> and Their Contribution to Chloroplast Phylogeny and Adaptive Evolution. <i>International Journal of Molecular Sciences</i> , 2018, 19, 1847.	1.8	92
11	A large-scale phylogeny of the lycophyte genus <i>Selaginella</i> (Selaginellaceae: Lycopodiopsida) based on plastid and nuclear loci. <i>Cladistics</i> , 2016, 32, 360-389.	1.5	80
12	Warming and nitrogen deposition are interactive in shaping surface soil microbial communities near the alpine timberline zone on the eastern Qinghai-Tibet Plateau, southwestern China. <i>Applied Soil Ecology</i> , 2016, 101, 72-83.	2.1	78
13	Phylogeny of Chinese <i>Allium</i> Species in Section <i>Daghestanica</i> and Adaptive Evolution of <i>Allium</i> (Amaryllidaceae, Alliioideae) Species Revealed by the Chloroplast Complete Genome. <i>Frontiers in Plant Science</i> , 2019, 10, 460.	1.7	64
14	Phylogeographic analysis of a temperate-deciduous forest restricted plant (<i>Bupleurum longiradiatum</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf <i>Molecular Phylogenetics and Evolution</i> , 2013, 68, 628-643.	1.2	57
15	In Vitro and In Vivo Antioxidant Activity of a Water-Soluble Polysaccharide from <i>Dendrobium denneanum</i> . <i>Molecules</i> , 2011, 16, 1579-1592.	1.7	53
16	Molecular systematics of <i>Angelica</i> and allied genera (Apiaceae) from the Hengduan Mountains of China based on nrDNA ITS sequences: phylogenetic affinities and biogeographic implications. <i>Journal of Plant Research</i> , 2009, 122, 403-414.	1.2	51
17	Insights into phylogeny, age and evolution of <i>Allium</i> (Amaryllidaceae) based on the whole plastome sequences. <i>Annals of Botany</i> , 2020, 125, 1039-1055.	1.4	49
18	Backbone phylogeny and evolution of Apioideae (Apiaceae): New insights from phylogenomic analyses of plastome data. <i>Molecular Phylogenetics and Evolution</i> , 2021, 161, 107183.	1.2	47

#	ARTICLE	IF	CITATIONS
19	Morphological and ecological divergence of <i>Lilium</i> and <i>Nomocharis</i> within the Hengduan Mountains and Qinghai-Tibetan Plateau may result from habitat specialization and hybridization. <i>BMC Evolutionary Biology</i> , 2015, 15, 147.	3.2	42
20	Plastomes of eight <i>Ligusticum</i> species: characterization, genome evolution, and phylogenetic relationships. <i>BMC Plant Biology</i> , 2020, 20, 519.	1.6	42
21	Phylogeny and biogeography of Chinese <i>Heracleum</i> (Apiaceae tribe Tordylieae) with comments on their fruit morphology. <i>Plant Systematics and Evolution</i> , 2011, 296, 179-203.	0.3	41
22	Chromosome diversity and evolution in tribe Lilieae (Liliaceae) with emphasis on Chinese species. <i>Journal of Plant Research</i> , 2012, 125, 55-69.	1.2	40
23	A new species in the genus <i>Nomocharis</i> Franchet (Liliaceae): evidence that brings the genus <i>Nomocharis</i> into <i>Lilium</i> . <i>Plant Systematics and Evolution</i> , 2012, 298, 69-85.	0.3	39
24	New Insights into the Phylogeny of <i>Angelica</i> and its Allies (Apiaceae) with Emphasis on East Asian Species, Inferred from nrDNA, cpDNA, and Morphological Evidence. <i>Systematic Botany</i> , 2013, 38, 266-281.	0.2	37
25	A transcriptome-based study on the phylogeny and evolution of the taxonomically controversial subfamily Apioideae (Apiaceae). <i>Annals of Botany</i> , 2020, 125, 937-953.	1.4	35
26	Molecular phylogeny, divergence time estimates and historical biogeography within one of the world's largest monocot genera. <i>AoB PLANTS</i> , 2016, 8, plw041.	1.2	33
27	Phylogenetic Analysis of the Sonneratiaceae and its Relationship to Lythraceae Based on ITS Sequences of nrDNA. <i>Journal of Plant Research</i> , 2000, 113, 253-258.	1.2	31
28	Comparative Chloroplast Genomics of <i>Fritillaria</i> (Liliaceae), Inferences for Phylogenetic Relationships between <i>Fritillaria</i> and <i>Lilium</i> and Plastome Evolution. <i>Plants</i> , 2020, 9, 133.	1.6	31
29	The effect of Hengduan Mountains Region (HMR) uplift to environmental changes in the HMR and its eastern adjacent area: Tracing the evolutionary history of <i>Allium</i> section <i>Sikkimensia</i> (Amaryllidaceae). <i>Molecular Phylogenetics and Evolution</i> , 2019, 130, 380-396.	1.2	28
30	Revisiting the evolutionary events in <i>Allium</i> subgenus <i>Cyathophora</i> (Amaryllidaceae): Insights into the effect of the Hengduan Mountains Region (HMR) uplift and Quaternary climatic fluctuations to the environmental changes in the Qinghai-Tibet Plateau. <i>Molecular Phylogenetics and Evolution</i> , 2016, 94, 802-813.	1.2	27
31	Comparative Analysis of the Complete Plastid Genome of Five <i>Bupleurum</i> Species and New Insights into DNA Barcoding and Phylogenetic Relationship. <i>Plants</i> , 2020, 9, 543.	1.6	26
32	A taxonomic reassessment in the Chinese <i>Bupleurum</i> (Apiaceae): Insights from morphology, nuclear ribosomal internal transcribed spacer, and chloroplast (<i>trnH-psbA</i> , <i>matK</i>) sequences. <i>Journal of Systematics and Evolution</i> , 2011, 49, 558-589.	1.6	25
33	Molecular phylogeny, biogeography and ecological niche modelling of <i>Cardiocrinum</i> (Liliaceae): insights into the evolutionary history of endemic genera distributed across the Sino-Japanese floristic region. <i>Annals of Botany</i> , 2017, 119, 59-72.	1.4	25
34	The complete plastomes of seven <i>Peucedanum</i> plants: comparative and phylogenetic analyses for the <i>Peucedanum</i> genus. <i>BMC Plant Biology</i> , 2022, 22, 101.	1.6	25
35	Molecular phylogenetics and historical biogeography of the tribe Lilieae (Liliaceae): bi-directional dispersal between biodiversity hotspots in Eurasia. <i>Annals of Botany</i> , 2018, 122, 1245-1262.	1.4	23
36	Intraspecific differentiation of <i>Allium wallichii</i> (Amaryllidaceae) inferred from chloroplast DNA and internal transcribed spacer fragments. <i>Journal of Systematics and Evolution</i> , 2014, 52, 341-354.	1.6	22

#	ARTICLE	IF	CITATIONS
37	Molecular Authentication of the Traditional Chinese Medicinal Plant <i>Euphorbia pekinensis</i> . <i>Planta Medica</i> , 2007, 73, 91-93.	0.7	20
38	Historical biogeography of the <i>Angelica</i> group (Apiaceae tribe Selineae) inferred from analyses of nrDNA and cpDNA sequences. <i>Journal of Systematics and Evolution</i> , 2012, 50, 206-217.	1.6	20
39	Molecular phylogenetics of <i>Pimpinella</i> and allied genera (Apiaceae), with emphasis on Chinese native species, inferred from nrDNA ITS and cpDNA intron sequence data. <i>Nordic Journal of Botany</i> , 2014, 32, 642-657.	0.2	20
40	Phylogenetic reappraisal of <i>Allium</i> subgenus <i>Cyathophora</i> (Amaryllidaceae) and related taxa, with a proposal of two new sections. <i>Journal of Plant Research</i> , 2014, 127, 275-286.	1.2	19
41	Complete Plastid Genome Sequencing of Eight Species from <i>Hansenia</i> , <i>Haplosphaera</i> and <i>Sinodielsia</i> (Apiaceae): Comparative Analyses and Phylogenetic Implications. <i>Plants</i> , 2020, 9, 1523.	1.6	18
42	Sequencing and analyses on chloroplast genomes of <i>Tetrataenium candicans</i> and two allies give new insights on structural variants, DNA barcoding and phylogeny in Apiaceae subfamily Apioideae. <i>PeerJ</i> , 2019, 7, e8063.	0.9	18
43	Comparative Analysis of the Complete Chloroplast Genomes in <i>Allium</i> Subgenus <i>Cyathophora</i> (Amaryllidaceae): Phylogenetic Relationship and Adaptive Evolution. <i>BioMed Research International</i> , 2020, 2020, 1-17.	0.9	16
44	Phylogeny and Comparative Analysis of Chinese <i>Chamaesium</i> Species Revealed by the Complete Plastid Genome. <i>Plants</i> , 2020, 9, 965.	1.6	15
45	Comparative morphology of the leaf epidermis in <i>Fritillaria</i> (Liliaceae) from China. <i>Botanical Journal of the Linnean Society</i> , 2009, 160, 93-109.	0.8	14
46	Phylogenetic analyses and chromosome counts reveal multiple cryptic species in <i>Bupleurum commelynoideum</i> (Apiaceae). <i>Journal of Systematics and Evolution</i> , 2015, 53, 104-116.	1.6	14
47	Bright side? The impacts of Three Gorges Reservoir on local ecological service of soil conservation in southwestern China. <i>Environmental Earth Sciences</i> , 2017, 76, 1.	1.3	14
48	Phylogeography and genetic effects of habitat fragmentation on endemic <i>Urophysa</i> (Ranunculaceae) in Yungui Plateau and adjacent regions. <i>PLoS ONE</i> , 2017, 12, e0186378.	1.1	12
49	Molecular authentication of the traditional Chinese medicinal plant <i>Angelica sinensis</i> based on internal transcribed spacer of nrDNA. <i>Electronic Journal of Biotechnology</i> , 2010, 13, .	1.2	10
50	Karyotype studies in thirty-two species of <i>Lilium</i> (Liliaceae) from China. <i>Nordic Journal of Botany</i> , 2011, 29, 746-761.	0.2	10
51	Phylogeny and highland adaptation of Chinese species in <i>Allium</i> section <i>Daghestanica</i> (Amaryllidaceae) revealed by transcriptome sequencing. <i>Molecular Phylogenetics and Evolution</i> , 2020, 146, 106737.	1.2	10
52	Phylogeny, Age, and Evolution of Tribe Lilieae (Liliaceae) Based on Whole Plastid Genomes. <i>Frontiers in Plant Science</i> , 2021, 12, 699226.	1.7	10
53	Phylogenetic position of <i>Ligusticopsis</i> (Apiaceae, Apioideae): evidence from molecular data and carpological characters. <i>AoB PLANTS</i> , 2022, 14, plac008.	1.2	10
54	Karyotypes of 16 populations of eight species in the genus <i>Polygonatum</i> (Asparagaceae) from China. <i>Botanical Journal of the Linnean Society</i> , 2009, 159, 245-254.	0.8	9

#	ARTICLE	IF	CITATIONS
55	Phylogeography of an alpine plant (<i>Bupleurum smithii</i> , Apiaceae) endemic to the Qinghai-Tibetan Plateau and adjacent regions inferred from chloroplast DNA sequence variation. <i>Journal of Systematics and Evolution</i> , 2013, 51, 382-395.	1.6	9
56	<i>Lilium yapingense</i> (Liliaceae), a New Species from Yunnan, China, and its Systematic Significance Relative to <i>Nomocharis</i> . <i>Annales Botanici Fennici</i> , 2013, 50, 187-194.	0.0	9
57	Adaptation Evolution and Phylogenetic Analyses of Species in Chinese <i>Allium</i> Section <i>Pallasia</i> and Related Species Based on Complete Chloroplast Genome Sequences. <i>BioMed Research International</i> , 2020, 2020, 1-13.	0.9	9
58	Effects of Mountain Uplift and Climatic Oscillations on Phylogeography and Species Divergence of <i>Chamaesium</i> (Apiaceae). <i>Frontiers in Plant Science</i> , 2021, 12, 673200.	1.7	9
59	<i>In Vivo</i> Immunomodulatory Activities of Neutral Polysaccharide (DDP1-1) from <i>Dendrobium denneanum</i> . <i>Ying Yong Yu Huan Jing Sheng Wu Xue Bao = Chinese Journal of Applied and Environmental Biology</i> , 2010, 16, 376-379.	0.1	9
60	Molecular Authentication of the Traditional Dai Medicinal Plant <i>Croton caudatus</i> . <i>Planta Medica</i> , 2007, 73, 611-613.	0.7	8
61	Evolution of the platycodonoid group with particular references to biogeography and character evolution. <i>Journal of Integrative Plant Biology</i> , 2014, 56, 995-1008.	4.1	8
62	Characterization of the complete chloroplast genome of <i>Allium prattii</i> . <i>Mitochondrial DNA Part B: Resources</i> , 2018, 3, 153-154.	0.2	8
63	A phylogenetic study of Chinese <i>Polygonatum</i> (Polygonateae, Asparagaceae). <i>Nordic Journal of Botany</i> , 2019, 37, .	0.2	8
64	Chloroplast genomic comparison of two sister species <i>Allium macranthum</i> and <i>A. fasciculatum</i> provides valuable insights into adaptive evolution. <i>Genes and Genomics</i> , 2020, 42, 507-517.	0.5	8
65	Driving forces for recovery of forest vegetation after harvesting a subalpine oak forest in eastern Tibetan Plateau. <i>Environmental Science and Pollution Research</i> , 2021, 28, 67748-67763.	2.7	8
66	Morphological cladistic analysis of <i>Ligusticum</i> (Umbelliferae) in China. <i>Nordic Journal of Botany</i> , 2008, 26, 118-128.	0.2	7
67	Karyotype and cytogeography of the genus <i>Heracleum</i> (Apiaceae) in the Hengduan Mountains. <i>Journal of Systematics and Evolution</i> , 2009, 47, 273-285.	1.6	7
68	Phylogeography of <i>Angelica nitida</i> (Apiaceae) endemic to the Qinghai-Tibet Plateau based on chloroplast DNA sequences. <i>Journal of Systematics and Evolution</i> , 2013, 51, 564-577.	1.6	7
69	Cytotaxonomy of <i>Allium</i> (Amaryllidaceae) subgenera <i>Cyathophora</i> and <i>Amerallium</i> sect. <i>Bromatorrhiza</i> . <i>Phytotaxa</i> , 2017, 331, 185.	0.1	7
70	<i>Hansenia pinnatiinvolucellata</i> is conspecific with <i>H. weberbaueriana</i> (Apiaceae) based on morphology and molecular data. <i>Phytotaxa</i> , 2019, 418, 203-210.	0.1	7
71	A Phytogeographic Divide Along the 500 mm Isohyet in the Qinghai-Tibet Plateau: Insights From the Phylogeographic Evidence of Chinese <i>Alliums</i> (Amaryllidaceae). <i>Frontiers in Plant Science</i> , 2019, 10, 149.	1.7	7
72	Out of the Qinghai-Tibetan Plateau and rapid radiation across Eurasia for <i>Allium</i> section <i>Daghestanica</i> (Amaryllidaceae). <i>AoB PLANTS</i> , 2021, 13, plab017.	1.2	7

#	ARTICLE	IF	CITATIONS
73	Phylogeny and Comparative Analysis for the Plastid Genomes of Five Tulipa (Liliaceae). BioMed Research International, 2021, 2021, 1-10.	0.9	7
74	Molecular evolution and phylogenetic relationships of Ligusticum (Apiaceae) inferred from the whole plastome sequences. BMC Ecology and Evolution, 2022, 22, 55.	0.7	7
75	Epidermal Morphology of <i>Ligusticum</i> (Apiaceae) from China. Annales Botanici Fennici, 2010, 47, 261-279.	0.0	6
76	A cytotaxonomic analysis of Chinese <i>Polygonatum</i> (Asparagaceae) species. Nordic Journal of Botany, 2014, 32, 441-451.	0.2	6
77	<i>Angelica onosepala</i> and <i>Heracleum yunnanense</i> are synonyms and refer to a species of <i>Tetrataenium</i> (Apiaceae). Nordic Journal of Botany, 2018, 36, njb-01563.	0.2	6
78	Phylogeny and taxonomy of <i>Meeboldia</i> , <i>Sinodielsia</i> and their relatives (Apiaceae: Apioideae) inferred from nrDNA ITS, plastid DNA intron (<i>rpl16</i> and <i>rps16</i>) sequences and morphological characters. Phytotaxa, 2021, 482, 121-142.	0.1	6
79	New insights into the phylogeny of <i>Sinocarum</i> (Apiaceae, Apioideae) based on morphological and molecular data. PhytoKeys, 2021, 175, 13-32.	0.4	6
80	Karyotypes of Four Genera in Liliaceae (<i>s.str.</i>) from Hengduan Mountains of Southwestern China. Acta Botanica Yunnanica, 2010, 31, 399-405.	0.1	6
81	<i>Bupleurum baimaense</i> (Apiaceae), a New Species from Hengduan Mountains, China. Annales Botanici Fennici, 2013, 50, 379-385.	0.0	5
82	<i>Polygonatum gongshanense</i> (Asparagaceae), a New Species from Gongshan, Yunnan, China. Annales Botanici Fennici, 2014, 51, 333-336.	0.0	5
83	<i>Pimpinella rhomboidea</i> var. <i>tenuiloba</i> is a synonym of <i>Melanosciadium bipinnatum</i> (Apiaceae). Nordic Journal of Botany, 2015, 33, 659-661.	0.2	5
84	Intraspecific differentiation of <i>Pleurospermum hookeri</i> (Apiaceae), and its interspecific relationships with two close relatives in the genus <i>Pleurospermum</i> . Journal of Systematics and Evolution, 2015, 53, 308-320.	1.6	5
85	<i>Semenovia torilifolia</i> is conspecific with <i>S. malcolmii</i> (Apiaceae) based on morphology and molecular data. Phytotaxa, 2017, 321, 225.	0.1	5
86	The complete chloroplast genome of <i>Nomocharis pardanthina</i> . Mitochondrial DNA Part B: Resources, 2018, 3, 103-104.	0.2	5
87	<i>Allium xinlongense</i> (Amaryllidaceae, Allioideae), a new species from western Sichuan. Phytotaxa, 2020, 432, 274-282.	0.1	5
88	Complete Chloroplast Genome of <i>Cnidium monnieri</i> (Apiaceae) and Comparisons with Other Tribe Selineae Species. Diversity, 2022, 14, 323.	0.7	5
89	<i>Bupleurum candollei</i> var. <i>paucefulcrans</i> comb. nov. (Apiaceae) from Guizhou, China: comparison of allied species based on morphology, anatomy and molecular data. Nordic Journal of Botany, 2011, 29, 424-430.	0.2	4
90	Fragmented habitat drives significant genetic divergence in the Chinese endemic plant, <i>Urophysa henryi</i> (Ranunculaceae). Biochemical Systematics and Ecology, 2016, 69, 76-82.	0.6	4

#	ARTICLE	IF	CITATIONS
91	The complete chloroplast genome of <i>Semenovia gyirongensis</i> (Tribe Tordylieae, Apiaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1863-1864.	0.2	4
92	The complete chloroplast genome of <i>Chamaesium paradoxum</i> . Mitochondrial DNA Part B: Resources, 2019, 4, 2069-2070.	0.2	4
93	Sequence and phylogenetic analysis of complete plastid genome of a medicinal plant <i>Heracleum moellendorffii</i> . Mitochondrial DNA Part B: Resources, 2019, 4, 1251-1252.	0.2	4
94	New insights into the phylogeny and taxonomy of Chinese <i>Physospermopsis</i> (Apiaceae). <i>PhytoKeys</i> , 2021, 175, 67-88.	0.4	4
95	Accommodating <i>Haplosphaera</i> and <i>Hansenia</i> (Apiaceae) based on morphological and molecular evidence. <i>Phytotaxa</i> , 2020, 464, 207-216.	0.1	4
96	Notes on two species of <i>Brotherella</i> (Bryopsida: Sematophyllaceae) from Asia. <i>Journal of Bryology</i> , 2006, 28, 268-271.	0.4	3
97	The complete chloroplast genome of <i>Angelica nitida</i> . Mitochondrial DNA Part B: Resources, 2017, 2, 694-695.	0.2	3
98	The complete chloroplast genome of a wild onion species <i>Allium monanthum</i> (Alliaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 854-855.	0.2	3
99	Characterization of the complete chloroplast genome of <i>Allium kingdonii</i> . Mitochondrial DNA Part B: Resources, 2019, 4, 868-869.	0.2	3
100	Chromosomal Study on Chinese <i>Bupleurum</i> L. (Apiaceae). <i>Zhi Wu Ke Xue Xue Bao</i> , 2013, 31, 11.	0.1	3
101	<i>Lysimachia xuyongensis</i> (Primulaceae), a new species from Sichuan, China. <i>Phytotaxa</i> , 2021, 525, 59-64.	0.1	3
102	<i>Angelica dabashanensis</i> (Apiaceae), a New Species from Shaanxi, China. <i>Annales Botanici Fennici</i> , 2012, 49, 125-133.	0.0	2
103	<i>Pternopetalum monophyllum</i> (Apiaceae), a New Species from Sichuan, China. <i>Annales Botanici Fennici</i> , 2014, 51, 414-418.	0.0	2
104	On the identity of <i>Pternopetalum botrychioides</i> (Apiaceae), introducing <i>P. latipinnulatum</i> comb. & stat. nov.. <i>Phytotaxa</i> , 2015, 226, 233.	0.1	2
105	<i>Bupleurum shanianum</i> sp. nov. (Apiaceae) from China. <i>Nordic Journal of Botany</i> , 2015, 33, 71-73.	0.2	2
106	Phylogeography of two closely related species of <i>Allium</i> endemic to East Asia: Population evolution in response to climate oscillations. <i>Ecology and Evolution</i> , 2018, 8, 7986-7999.	0.8	2
107	Comparative Complete Chloroplast Genome Analyses and Contribution to the Understanding of Chloroplast Phylogeny and Adaptive Evolution in Subgenus <i>Anguinum</i> . <i>Russian Journal of Genetics</i> , 2019, 55, 872-884.	0.2	2
108	The complete chloroplast genome of <i>Lilium Lankongense</i> Franchet (Liliaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1824-1825.	0.2	2

#	ARTICLE	IF	CITATIONS
109	The complete chloroplast genome of <i>Allium macrostemon</i> . Mitochondrial DNA Part B: Resources, 2019, 4, 1938-1939.	0.2	2
110	The complete chloroplast genome of <i>Meeboldia yunnanensis</i> (Apiaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 4176-4177.	0.2	2
111	The complete chloroplast genome sequence of <i>Heracleum yungningense</i> . Mitochondrial DNA Part B: Resources, 2020, 5, 1783-1784.	0.2	2
112	A Combined Morphological and Molecular Evolutionary Analysis of Karst-Environment Adaptation for the Genus <i>Urophysa</i> (Ranunculaceae). Frontiers in Plant Science, 2021, 12, 667988.	1.7	2
113	<i>Semenovia gyirongensis</i> (Apiaceae), a new species from Xizang, China. PhytoKeys, 2017, 82, 57-72.	0.4	2
114	<p>New insights into the phylogenetic position of Hymenidium dentatum (Apiaceae) inferred from nrDNA and morphological evidence</p>. Phytotaxa, 2020, 452, 46-54.	0.1	2
115	<p>Tongoloa tagongensis (Apiaceae), a new species from the Hengduan Mountains, China</p>. Phytotaxa, 2020, 461, 12-20.	0.1	2
116	<p>Notholirion campanulatum is co-specific with N. bulbuliferum (Liliaceae) based on morphology and molecular data</p>. Phytotaxa, 2020, 471, 234-246.	0.1	2
117	Comparative Plastome Analysis of Three Amaryllidaceae Subfamilies: Insights into Variation of Genome Characteristics, Phylogeny, and Adaptive Evolution. BioMed Research International, 2022, 2022, 1-20.	0.9	2
118	<i>Chamaesium jiulongense</i> sp. nov. (Apiaceae) from Sichuan, China. Nordic Journal of Botany, 2017, 35, 676-680.	0.2	1
119	Morphological and micromorphological data support the independent specific status of <i>Chamaesium spatuliferum</i> (Apiaceae) from China. Phytotaxa, 2017, 314, 81.	0.1	1
120	The complete chloroplast genome of <i>Notholirion macrophyllum</i> . Mitochondrial DNA Part B: Resources, 2018, 3, 1102-1103.	0.2	1
121	The complete chloroplast genome of <i>Haplosphaera phaea</i> (Apiaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1969-1970.	0.2	1
122	Comparative analysis of complete plastid genomes from <i>Lilium lankongense</i> Franchet and its closely related species and screening of <i>Lilium</i>-specific primers. PeerJ, 2021, 9, e10964.	0.9	1
123	The complete chloroplast genome of <i>Semenovia thomsonii</i> (Tordylieae: Apiaceae), a new record from Xizang, China. Mitochondrial DNA Part B: Resources, 2021, 6, 1911-1913.	0.2	1
124	<i>Hymenidium pilosum</i> (Apiaceae) is a synonym of <i>H. apiolens</i> based on morphology and molecular data. Nordic Journal of Botany, 2020, 38, .	0.2	1
125	Yuzurimine from <i>Daphniphyllum macropodum</i> Miq.. Acta Crystallographica Section E: Structure Reports Online, 2010, 66, o3013-o3013.	0.2	0
126	<i>Spiraea fangii</i> (Rosaceae), a new species from Sichuan, China. Phytotaxa, 2016, 268, 155.	0.1	0

#	ARTICLE	IF	CITATIONS
127	<i>Spiraea longifolia</i> (Rosaceae), a New Species from Sichuan, China. <i>Annales Botanici Fennici</i> , 2016, 53, 238-242.	0.0	0
128	Characterization of the complete plastid genome sequence of <i>Allium Fasciculatum</i> . <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 1782-1783.	0.2	0
129	Complete plastome sequence of <i>Tetrataenium Candicans</i> (tribe Tordylieae, Apiaceae): a medicinal plant. <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 3429-3431.	0.2	0
130	Characterization of the complete chloroplast genome of <i>Taibaisanqi</i> (<i>Tongoloa silaifolia</i>). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 2912-2913.	0.2	0
131	<i>Meeboldia linearis</i> sp. nov. (Apiaceae) from Xizang, China. <i>Nordic Journal of Botany</i> , 2020, 38, .	0.2	0
132	Complete plastome sequence of <i>Tetrataenium yunnanense</i> (tribe Tordylieae, Apiaceae) with anti-tumor activity. <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 1525-1527.	0.2	0
133	Resurrection of the genus <i>Similisinocarum</i> (Apiaceae) based on evidence from morphology and ITS sequences. <i>Phytotaxa</i> , 2021, 497, 127-137.	0.1	0
134	<i>Sinocarum muliense</i> (Apiaceae), a new species from Sichuan, China. <i>Nordic Journal of Botany</i> , 2022, 2022, .	0.2	0