

Xing-Jin He

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	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
2	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
3	Comparative Plastome Analysis of Three Amaryllidaceae Subfamilies: Insights into Variation of Genome Characteristics, Phylogeny, and Adaptive Evolution. <i>BioMed Research International</i> , 2022, 2022, 1-20.	0.9	2
4	<i>Sinocarum muliense</i> (Apiaceae), a new species from Sichuan, China. <i>Nordic Journal of Botany</i> , 2022, 2022, .	0.2	0
5	Molecular evolution and phylogenetic relationships of <i>Ligusticum</i> (Apiaceae) inferred from the whole plastome sequences. <i>Bmc Ecology and Evolution</i> , 2022, 22, 55.	0.7	7
6	Complete Chloroplast Genome of <i>Cnidium monnieri</i> (Apiaceae) and Comparisons with Other Tribe Selineae Species. <i>Diversity</i> , 2022, 14, 323.	0.7	5
7	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. <i>Phytotaxa</i> , 2021, 482, 121-142.	0.1	6
8	New insights into the phylogeny and taxonomy of Chinese <i>Physospermopsis</i> (Apiaceae). <i>PhytoKeys</i> , 2021, 175, 67-88.	0.4	4
9	New insights into the phylogeny of <i>Sinocarum</i> (Apiaceae, Apioideae) based on morphological and molecular data. <i>PhytoKeys</i> , 2021, 175, 13-32.	0.4	6
10	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. <i>PeerJ</i> , 2021, 9, e10964.	0.9	1
11	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
12	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
13	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
14	A Combined Morphological and Molecular Evolutionary Analysis of Karst-Environment Adaptation for the Genus <i>Urophysa</i> (Ranunculaceae). <i>Frontiers in Plant Science</i> , 2021, 12, 667988.	1.7	2
15	The complete chloroplast genome of <i>Semenovia thomsonii</i> (Tordylieae: Apiaceae), a new record from Xizang, China. <i>Mitochondrial DNA Part B: Resources</i> , 2021, 6, 1911-1913.	0.2	1
16	Phylogeny and Comparative Analysis for the Plastid Genomes of Five <i>Tulipa</i> (Liliaceae). <i>BioMed Research International</i> , 2021, 2021, 1-10.	0.9	7
17	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
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

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19	<i>Lysimachia xuyongensis</i> (Primulaceae), a new species from Sichuan, China. <i>Phytotaxa</i> , 2021, 525, 59-64.	0.1	3
20	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
21	RASP 4: Ancestral State Reconstruction Tool for Multiple Genes and Characters. <i>Molecular Biology and Evolution</i> , 2020, 37, 604-606.	3.5	338
22	Plastomes of eight <i>Ligusticum</i> species: characterization, genome evolution, and phylogenetic relationships. <i>BMC Plant Biology</i> , 2020, 20, 519.	1.6	42
23	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
24	<i>Meeboldia linearis</i> sp. nov. (Apiaceae) from Xizang, China. <i>Nordic Journal of Botany</i> , 2020, 38, .	0.2	0
25	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
26	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
27	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
28	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
29	The complete chloroplast genome sequence of <i>Heracleum yungningense</i> . <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 1783-1784.	0.2	2
30	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
31	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
32	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
33	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
34	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
35	<i>Allium xinlongense</i> (Amaryllidaceae, Alliioideae), a new species from western Sichuan. <i>Phytotaxa</i> , 2020, 432, 274-282.	0.1	5
36	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

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37	Hymenidium pilosum (Apiaceae) is a synonym of H. apiolens based on morphology and molecular data. Nordic Journal of Botany, 2020, 38, .	0.2	1
38	New insights into the phylogenetic position of Hymenidium dentatum (Apiaceae) inferred from nrDNA and morphological evidence. Phytotaxa, 2020, 452, 46-54.	0.1	2
39	Tongoloa tagongensis (Apiaceae), a new species from the Hengduan Mountains, China. Phytotaxa, 2020, 461, 12-20.	0.1	2
40	Accommodating Haplosphaera Hansenia (Apiaceae) based on morphological and molecular evidence. Phytotaxa, 2020, 464, 207-216.	0.1	4
41	Notholirion campanulatum is co-specific with N. bulbuliferum (Liliaceae) based on morphology and molecular data. Phytotaxa, 2020, 471, 234-246.	0.1	2
42	Comparative Complete Chloroplast Genome Analyses and Contribution to the Understanding of Chloroplast Phylogeny and Adaptive Evolution in Subgenus Anguinum. Russian Journal of Genetics, 2019, 55, 872-884.	0.2	2
43	The complete chloroplast genome of Semenovia gyirongensis (Tribe Tordylieae, Apiaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1863-1864.	0.2	4
44	The complete chloroplast genome of Chamaesium paradoxum. Mitochondrial DNA Part B: Resources, 2019, 4, 2069-2070.	0.2	4
45	The complete chloroplast genome of Haplosphaera phaea (Apiaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1969-1970.	0.2	1
46	Characterization of the complete plastid genome sequence of Allium Fasciculatum. Mitochondrial DNA Part B: Resources, 2019, 4, 1782-1783.	0.2	0
47	Complete plastome sequence of Tetrataenium Candicans (tribe Tordylieae, Apiaceae): a medicinal plant. Mitochondrial DNA Part B: Resources, 2019, 4, 3429-3431.	0.2	0
48	The complete chloroplast genome of Lilium Lankongense Franchet (Liliaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1824-1825.	0.2	2
49	Characterization of the complete chloroplast genome of Taibaisanqi (Tongoloa silaifolia). Mitochondrial DNA Part B: Resources, 2019, 4, 2912-2913.	0.2	0
50	Hansenia pinnativolucellata is conspecific with H. weberbaueriana (Apiaceae) based on morphology and molecular data. Phytotaxa, 2019, 418, 203-210.	0.1	7
51	The complete chloroplast genome of Allium macrostemon. Mitochondrial DNA Part B: Resources, 2019, 4, 1938-1939.	0.2	2
52	A Phytogeographic Divide Along the 500 mm Isohyet in the Qinghai-Tibet Plateau: Insights From the Phylogeographic Evidence of Chinese Alliums (Amaryllidaceae). Frontiers in Plant Science, 2019, 10, 149.	1.7	7
53	Phylogeny of Chinese Allium Species in Section Daghestanica and Adaptive Evolution of Allium (Amaryllidaceae, Alliioideae) Species Revealed by the Chloroplast Complete Genome. Frontiers in Plant Science, 2019, 10, 460.	1.7	64
54	The complete chloroplast genome of a wild onion species Allium monanthum (Alliaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 854-855.	0.2	3

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55	Sequence and phylogenetic analysis of complete plastid genome of a medicinal plant <i>Heracleum moellendorffii</i> . <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 1251-1252.	0.2	4
56	Characterization of the complete chloroplast genome of <i>Allium kingdonii</i> . <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 868-869.	0.2	3
57	The complete chloroplast genome of <i>Meeboldia yunnanensis</i> (Apiaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 4176-4177.	0.2	2
58	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
59	A phylogenetic study of Chinese <i>Polygonatum</i> (Polygonateae, Asparagaceae). <i>Nordic Journal of Botany</i> , 2019, 37, .	0.2	8
60	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
61	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
62	<i>Angelica onosepala</i> and <i>Heracleum yunnanense</i> are synonyms and refer to a species of <i>Tetrataenium</i> (Apiaceae). <i>Nordic Journal of Botany</i> , 2018, 36, njb-01563.	0.2	6
63	The complete chloroplast genome of <i>Notholition macrophyllum</i> . <i>Mitochondrial DNA Part B: Resources</i> , 2018, 3, 1102-1103.	0.2	1
64	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
65	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
66	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
67	The complete chloroplast genome of <i>Nomocharis pardanthina</i> . <i>Mitochondrial DNA Part B: Resources</i> , 2018, 3, 103-104.	0.2	5
68	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
69	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
70	The complete chloroplast genome of <i>Angelica nitida</i> . <i>Mitochondrial DNA Part B: Resources</i> , 2017, 2, 694-695.	0.2	3
71	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
72	<i>Chamaesium jiulongense</i> sp. nov. (Apiaceae) from Sichuan, China. <i>Nordic Journal of Botany</i> , 2017, 35, 676-680.	0.2	1

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73	<i>Semenovia torilifolia</i> is conspecific with <i>S. malcolmii</i> (Apiaceae) based on morphology and molecular data. <i>Phytotaxa</i> , 2017, 321, 225.	0.1	5
74	Morphological and micromorphological data support the independent specific status of <i>Chamaesium spatuliferum</i> (Apiaceae) from China. <i>Phytotaxa</i> , 2017, 314, 81.	0.1	1
75	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
76	<i>Semenovia gyirongensis</i> (Apiaceae), a new species from Xizang, China. <i>PhytoKeys</i> , 2017, 82, 57-72.	0.4	2
77	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
78	<i>Spiraea fangii</i> (Rosaceae), a new species from Sichuan, China. <i>Phytotaxa</i> , 2016, 268, 155.	0.1	0
79	Fragmented habitat drives significant genetic divergence in the Chinese endemic plant, <i>Urophysa henryi</i> (Ranunculaceae). <i>Biochemical Systematics and Ecology</i> , 2016, 69, 76-82.	0.6	4
80	<i>Spiraea longifolia</i> (Rosaceae), a New Species from Sichuan, China. <i>Annales Botanici Fennici</i> , 2016, 53, 238-242.	0.0	0
81	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
82	A tool for the analysis of chromosomes: KaryoType. <i>Taxon</i> , 2016, 65, 586-592.	0.4	110
83	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
84	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
85	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
86	<i>Pimpinella rhomboidea</i> var. <i>tenuiloba</i> is a synonym of <i>Melanosciadium bipinnatum</i> (Apiaceae). <i>Nordic Journal of Botany</i> , 2015, 33, 659-661.	0.2	5
87	Intraspecific differentiation of <i>Pleurospermum hookeri</i> (Apiaceae), and its interspecific relationships with two close relatives in the genus <i>Pleurospermum</i> . <i>Journal of Systematics and Evolution</i> , 2015, 53, 308-320.	1.6	5
88	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
89	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
90	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

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91	<i>Bupleurum shanianum</i> sp. nov. (Apiaceae) from China. <i>Nordic Journal of Botany</i> , 2015, 33, 71-73.	0.2	2
92	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
93	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
94	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
95	<i>Pternopetalum monophyllum</i> (Apiaceae), a New Species from Sichuan, China. <i>Annales Botanici Fennici</i> , 2014, 51, 414-418.	0.0	2
96	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
97	A cytotaxonomic analysis of Chinese <i>Polygonatum</i> (Asparagaceae) species. <i>Nordic Journal of Botany</i> , 2014, 32, 441-451.	0.2	6
98	<i>Polygonatum gongshanense</i> (Asparagaceae), a New Species from Gongshan, Yunnan, China. <i>Annales Botanici Fennici</i> , 2014, 51, 333-336.	0.0	5
99	Evolutionary events in <i>Lilium</i> (including <i>Nomocharis</i> , Liliaceae) are temporally correlated with orogenies of the Tibetan plateau and the Hengduan Mountains. <i>Molecular Phylogenetics and Evolution</i> , 2013, 68, 443-460.	1.2	97
100	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
101	<i>Bupleurum baimaense</i> (Apiaceae), a New Species from Hengduan Mountains, China. <i>Annales Botanici Fennici</i> , 2013, 50, 379-385.	0.0	5
102	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
103	Phylogeographic analysis of a temperate-deciduous forest restricted plant (<i>Bupleurum longiradiatum</i>) Tj ETQq1 1 0.784314 rgBT /Over Molecular Phylogenetics and Evolution, 2013, 68, 628-643.	1.2	57
104	<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
105	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
106	Chromosomal Study on Chinese <i>Bupleurum</i> L. (Apiaceae). <i>Zhi Wu Ke Xue Xue Bao</i> , 2013, 31, 11.	0.1	3
107	<i>Angelica dabashanensis</i> (Apiaceae), a New Species from Shaanxi, China. <i>Annales Botanici Fennici</i> , 2012, 49, 125-133.	0.0	2
108	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

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109	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
110	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
111	<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. <i>Nordic Journal of Botany</i> , 2011, 29, 424-430.	0.2	4
112	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
113	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
114	A taxonomic re-assessment 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
115	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
116	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
117	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
118	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
119	Yuzurimine from <i>Daphniphyllum macropodum</i> Miq.. <i>Acta Crystallographica Section E: Structure Reports Online</i> , 2010, 66, o3013-o3013.	0.2	0
120	Phylogeny and biogeography of <i>Allium</i> (Amaryllidaceae: Alliieae) based on nuclear ribosomal internal transcribed spacer and chloroplast <i>rps16</i> sequences, focusing on the inclusion of species endemic to China. <i>Annals of Botany</i> , 2010, 106, 709-733.	1.4	170
121	Epidermal Morphology of <i>Ligusticum</i> (Apiaceae) from China. <i>Annales Botanici Fennici</i> , 2010, 47, 261-279.	0.0	6
122	Karyotypes of Four Genera in Liliaceae (<i>s.str.</i>) from Hengduan Mountains of Southwestern China. <i>Acta Botanica Yunnanica</i> , 2010, 31, 399-405.	0.1	6
123	In Vivo 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
124	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
125	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
126	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

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127	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
128	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
129	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
130	Morphological cladistic analysis of <i>Ligusticum</i> (Umbelliferae) in China. <i>Nordic Journal of Botany</i> , 2008, 26, 118-128.	0.2	7
131	Molecular Authentication of the Traditional Chinese Medicinal Plant <i>Euphorbia pekinensis</i> . <i>Planta Medica</i> , 2007, 73, 91-93.	0.7	20
132	Molecular Authentication of the Traditional Dai Medicinal Plant <i>Croton caudatus</i> . <i>Planta Medica</i> , 2007, 73, 611-613.	0.7	8
133	Notes on two species of <i>Brotherella</i> (Bryopsida: Sematophyllaceae) from Asia. <i>Journal of Bryology</i> , 2006, 28, 268-271.	0.4	3
134	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