

Dezhu Li

List of Publications by Year in descending order

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395
papers

15,837
citations

39113

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34195

103
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all docs

407
docs citations

407
times ranked

11867
citing authors

#	ARTICLE	IF	CITATIONS
1	Testing complete plastomes and nuclear ribosomal DNA sequences for species identification in a taxonomically difficult bamboo genus <i>Fargesia</i> . <i>Plant Diversity</i> , 2023, 45, 147-155.	1.8	5
2	Phylotranscriptomic analyses reveal multiple whole-genome duplication events, the history of diversification and adaptations in the Araceae. <i>Annals of Botany</i> , 2023, 131, 199-214.	1.4	7
3	New insights into intergeneric relationships of <i>Hickeliinae</i> (Poaceae: Bambusoideae) revealed by complete plastid genomes. <i>Plant Diversity</i> , 2023, 45, 125-132.	1.8	1
4	The identity of <i>Dinochloa</i> species and enumeration of <i>Melocalamus</i> (Poaceae: Bambusoideae) in China. <i>Plant Diversity</i> , 2023, 45, 133-146.	1.8	4
5	Phylogeny and biogeography of <i>Fagus</i> (Fagaceae) based on 28 nuclear single-copy loci. <i>Journal of Systematics and Evolution</i> , 2022, 60, 759-772.	1.6	15
6	Sexual dimorphism, temporal niche differentiation, and evidence for the Jack Sprat effect in an annual dioecious plant. <i>Journal of Systematics and Evolution</i> , 2022, 60, 1078-1091.	1.6	6
7	Testing genome skimming for species discrimination in the large and taxonomically difficult genus <i>Rhododendron</i> . <i>Molecular Ecology Resources</i> , 2022, 22, 404-414.	2.2	35
8	Genetic analysis of walnut cultivars from southwest China: Implications for germplasm improvement. <i>Plant Diversity</i> , 2022, 44, 530-541.	1.8	11
9	A revision of <i>Dryopteris</i> sect. <i>Diclidisodon</i> (Dryopteridaceae) based on morphological and molecular evidence with description of a new species. <i>Plant Diversity</i> , 2022, 44, 181-190.	1.8	2
10	Morphological trait-matching in plant-Hymenoptera and plant-Diptera mutualisms across an elevational gradient. <i>Journal of Animal Ecology</i> , 2022, 91, 196-209.	1.3	8
11	Fire-prone Rhamnaceae with South African affinities in Cretaceous Myanmar amber. <i>Nature Plants</i> , 2022, 8, 125-135.	4.7	24
12	A well-supported nuclear phylogeny of Poaceae and implications for the evolution of C4 photosynthesis. <i>Molecular Plant</i> , 2022, 15, 755-777.	3.9	47
13	<i>Dryopteris sukungiana</i> (Dryopteridaceae), a new species of the <i>D. sparsa</i> complex from Southwest China. <i>Phytotaxa</i> , 2022, 533, 256-266.	0.1	2
14	Cryptic Species Diversification of the <i>Pedicularis siphonantha</i> Complex (Orobanchaceae) in the Mountains of Southwest China Since the Pliocene. <i>Frontiers in Plant Science</i> , 2022, 13, 811206.	1.7	5
15	Epigenetic regulation of seed-specific gene expression by DNA methylation valleys in castor bean. <i>BMC Biology</i> , 2022, 20, 57.	1.7	7
16	A worldwide phylogenetic classification of the Poaceae (Gramineae) III: An update. <i>Journal of Systematics and Evolution</i> , 2022, 60, 476-521.	1.6	61
17	Plastid phylogenomics shed light on intergeneric relationships and spatiotemporal evolutionary history of Melocanninae (Poaceae: Bambusoideae). <i>Journal of Systematics and Evolution</i> , 2022, 60, 640-652.	1.6	12
18	Herbarium phylogenomics: Resolving the generic status of the enigmatic <i>Pseudobartsia</i> (Orobanchaceae). <i>Journal of Systematics and Evolution</i> , 2022, 60, 1218-1228.	1.6	2

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19	Species-specific partial gene duplication in <i>Arabidopsis thaliana</i> evolved novel phenotypic effects on morphological traits under strong positive selection. <i>Plant Cell</i> , 2022, 34, 802-817.	3.1	15
20	Determinants of Genetic Structure in a Highly Heterogeneous Landscape in Southwest China. <i>Frontiers in Plant Science</i> , 2022, 13, 779989.	1.7	5
21	Grasses through space and time: An overview of the biogeographical and macroevolutionary history of Poaceae. <i>Journal of Systematics and Evolution</i> , 2022, 60, 522-569.	1.6	35
22	Testing the Complete Plastome for Species Discrimination, Cryptic Species Discovery and Phylogenetic Resolution in <i>Cephalotaxus</i> (Cephalotaxaceae). <i>Frontiers in Plant Science</i> , 2022, 13, .	1.7	16
23	Structural Variation of Plastomes Provides Key Insight Into the Deep Phylogeny of Ferns. <i>Frontiers in Plant Science</i> , 2022, 13, .	1.7	11
24	Genetic Diversity and Structure of Persian Walnut (<i>Juglans regia</i> L.) in Pakistan: Implications for Conservation. <i>Plants</i> , 2022, 11, 1652.	1.6	12
25	<i>Fragaria</i> mitogenomes evolve rapidly in structure but slowly in sequence and incur frequent multinucleotide mutations mediated by microinversions. <i>New Phytologist</i> , 2022, 236, 745-759.	3.5	16
26	The Genomic Selfing Syndrome Accompanies the Evolutionary Breakdown of Heterostyly. <i>Molecular Biology and Evolution</i> , 2021, 38, 168-180.	3.5	23
27	Resolving robust phylogenetic relationships of core Brassicaceae using genome skimming data. <i>Journal of Systematics and Evolution</i> , 2021, 59, 442-453.	1.6	16
28	Comparative plastomic analysis and insights into the phylogeny of <i>Salvia</i> (Lamiaceae). <i>Plant Diversity</i> , 2021, 43, 15-26.	1.8	19
29	Parallel ddRAD and Genome Skimming Analyses Reveal a Radiative and Reticulate Evolutionary History of the Temperate Bamboos. <i>Systematic Biology</i> , 2021, 70, 756-773.	2.7	38
30	Diversity in seed oil content and fatty acid composition in <i>Acer</i> species with potential as sources of nervonic acid. <i>Plant Diversity</i> , 2021, 43, 86-92.	1.8	11
31	Genetic innovations: Transposable element recruitment and de novo formation lead to the birth of orphan genes in the rice genome. <i>Journal of Systematics and Evolution</i> , 2021, 59, 341-351.	1.6	14
32	Current status of herbarium specimens and geographical distribution of bamboos (Gramineae: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 22	0.2	0
33	Distinct late Pleistocene subtropical-tropical divergence revealed by fifteen low-copy nuclear genes in a dominant species in South-East China. <i>Scientific Reports</i> , 2021, 11, 4147.	1.6	2
34	The <i>Pharus latifolius</i> genome bridges the gap of early grass evolution. <i>Plant Cell</i> , 2021, 33, 846-864.	3.1	32
35	Two new species of <i>Yushania</i> (Poaceae: Bambusoideae) from South China, with a taxonomic revision of related species. <i>Plant Diversity</i> , 2021, 43, 492-501.	1.8	5
36	The impact of a native dominant plant, <i>Euphorbia jolkin</i> , on plantâ€flower visitor networks and pollen deposition on stigmas of coâ€flowering species in subalpine meadows of Shangriã€La, SW China. <i>Journal of Ecology</i> , 2021, 109, 2107-2120.	1.9	5

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37	Genomic insights into the origin, domestication and genetic basis of agronomic traits of castor bean. <i>Genome Biology</i> , 2021, 22, 113.	3.8	32
38	Simultaneous diversification of Polypodiales and angiosperms in the Mesozoic. <i>Cladistics</i> , 2021, 37, 518-539.	1.5	38
39	Differential expressions of anthocyanin synthesis genes underlie flower color divergence in a sympatric <i>Rhododendron sanguineum</i> complex. <i>BMC Plant Biology</i> , 2021, 21, 204.	1.6	15
40	DNA methylation-mediated modulation of rapid desiccation tolerance acquisition and dehydration stress memory in the resurrection plant <i>Boea hygrometrica</i> . <i>PLoS Genetics</i> , 2021, 17, e1009549.	1.5	22
41	Phylogenomics of <i>Fargesia</i> and <i>Yushania</i> reveals a history of reticulate evolution. <i>Journal of Systematics and Evolution</i> , 2021, 59, 1183-1197.	1.6	9
42	Plastid NDH Pseudogenization and Gene Loss in a Recently Derived Lineage from the Largest Hemiparasitic Plant Genus <i>Pedicularis</i> (Orobanchaceae). <i>Plant and Cell Physiology</i> , 2021, 62, 971-984.	1.5	25
43	Rose without prickle: genomic insights linked to moisture adaptation. <i>National Science Review</i> , 2021, 8, nwab092.	4.6	15
44	Gene duplications and phylogenomic conflict underlie major pulses of phenotypic evolution in gymnosperms. <i>Nature Plants</i> , 2021, 7, 1015-1025.	4.7	68
45	Spatiotemporal maintenance of flora in the Himalaya biodiversity hotspot: Current knowledge and future perspectives. <i>Ecology and Evolution</i> , 2021, 11, 10794-10812.	0.8	38
46	Evolutionary and ecological factors structure a plant–bumblebee network in a biodiversity hotspot, the Himalaya–Hengduan Mountains. <i>Functional Ecology</i> , 2021, 35, 2523-2535.	1.7	10
47	Organelle Genomes and Transcriptomes of <i>Nymphaea</i> Reveal the Interplay between Intron Splicing and RNA Editing. <i>International Journal of Molecular Sciences</i> , 2021, 22, 9842.	1.8	7
48	New Genes Interacted With Recent Whole-Genome Duplicates in the Fast Stem Growth of Bamboos. <i>Molecular Biology and Evolution</i> , 2021, 38, 5752-5768.	3.5	28
49	Development of the petaloid bracts of a paleoherb species, <i>Saururus chinensis</i> . <i>PLoS ONE</i> , 2021, 16, e0255679.	1.1	1
50	Correlation Analysis Reveals an Important Role of GC Content in Accumulation of Deletion Mutations in the Coding Region of Angiosperm Plastomes. <i>Journal of Molecular Evolution</i> , 2021, 89, 73-80.	0.8	4
51	The complete chloroplast genome sequences of an endemic species of Urticaceae (<i>Debregeasia</i>) Tj ETQq1 1 0.784314 rgBT /Overlaid	0.2	0
52	Plastid phylogenomic insights into relationships of all flowering plant families. <i>BMC Biology</i> , 2021, 19, 232.	1.7	109
53	<i>Adiantum japonicum</i> , a new species of the <i>Adiantum pedatum</i> complex (Pteridaceae) from Japan. <i>Phytotaxa</i> , 2021, 525, 1-14.	0.1	0
54	Organelle Phylogenomics and Extensive Conflicting Phylogenetic Signals in the Monocot Order Poales. <i>Frontiers in Plant Science</i> , 2021, 12, 824672.	1.7	9

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55	Revisiting the phylogeny of Dipsacales: New insights from phylogenomic analyses of complete plastomic sequences. <i>Journal of Systematics and Evolution</i> , 2020, 58, 103-117.	1.6	30
56	The complete chloroplast genome of <i>Rhododendron delavayi</i> (Ericaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 37-38.	0.2	13
57	Born migrants: Historical biogeography of the cosmopolitan family Cannabaceae. <i>Journal of Systematics and Evolution</i> , 2020, 58, 461-473.	1.6	21
58	Evolutionary history of a relict conifer, <i>Pseudotaxus chienii</i> (Taxaceae), in south-east China during the late Neogene: old lineage, young populations. <i>Annals of Botany</i> , 2020, 125, 105-117.	1.4	27
59	Extreme plastid RNA editing may confound phylogenetic reconstruction: A case study of <i>Selaginella</i> (lycophytes). <i>Plant Diversity</i> , 2020, 42, 356-361.	1.8	13
60	Evolutionary legacy of a forest plantation tree species (<i>Pinus armandii</i>): Implications for widespread afforestation. <i>Evolutionary Applications</i> , 2020, 13, 2646-2662.	1.5	15
61	Complementary Transcriptome and Proteome Analyses Provide Insight into the Floral Transition in Bamboo (<i>Dendrocalamus latiflorus</i> Munro). <i>International Journal of Molecular Sciences</i> , 2020, 21, 8430.	1.8	3
62	Repeated intercontinental migrations and recurring hybridizations characterise the evolutionary history of yew (<i>Taxus</i> L.). <i>Molecular Phylogenetics and Evolution</i> , 2020, 153, 106952.	1.2	10
63	GetOrganelle: a fast and versatile toolkit for accurate de novo assembly of organelle genomes. <i>Genome Biology</i> , 2020, 21, 241.	3.8	1,538
64	Comparative analysis of plastid genomes within the Campanulaceae and phylogenetic implications. <i>PLoS ONE</i> , 2020, 15, e0233167.	1.1	11
65	Characteristics and Mutational Hotspots of Plastomes in <i>Debregeasia</i> (Urticaceae). <i>Frontiers in Genetics</i> , 2020, 11, 729.	1.1	18
66	Episodic and guanine-cytosine-biased bursts of intragenomic and interspecific synonymous divergence in <i>Ajugoideae</i> (Lamiaceae) mitogenomes. <i>New Phytologist</i> , 2020, 228, 1107-1114.	3.5	13
67	Development of 32 novel microsatellite loci in <i>Juglans sigillata</i> using genomic data. <i>Applications in Plant Sciences</i> , 2020, 8, e11328.	0.8	6
68	Exploration of Plastid Phylogenomic Conflict Yields New Insights into the Deep Relationships of Leguminosae. <i>Systematic Biology</i> , 2020, 69, 613-622.	2.7	131
69	ddRAD analyses reveal a credible phylogenetic relationship of the four main genera of <i>Bambusa-Dendrocalamus-Gigantochloa</i> complex (Poaceae: Bambusoideae). <i>Molecular Phylogenetics and Evolution</i> , 2020, 146, 106758.	1.2	19
70	A new subtribal classification of Arundinarieae (Poaceae, Bambusoideae) with the description of a new genus. <i>Plant Diversity</i> , 2020, 42, 127-134.	1.8	27
71	Evolution of Angiosperm Pollen: 8. Lamiids. <i>Annals of the Missouri Botanical Garden</i> , 2020, 105, 323-376.	1.3	10
72	Discovery of the first succulent bamboo (Poaceae, Bambusoideae) in a new genus from Laos karst areas, with a unique adaptation to seasonal drought. <i>PhytoKeys</i> , 2020, 156, 125-137.	0.4	5

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73	Documentation of plant diversity of Southeast Asia: the new role of Belt and Road Initiative. <i>PhytoKeys</i> , 2020, 138, 1-2.	0.4	1
74	Characterization of 30 microsatellite markers for distylous <i>Primula denticulata</i> (Primulaceae) using HiSeq sequencing. <i>Genes and Genetic Systems</i> , 2020, 95, 275-279.	0.2	1
75	Complete plastome of an endemic fern species from China: <i>Neocheiropteris palmatopedata</i> (Polypodiaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 2394-2395.	0.2	2
76	Complete chloroplast genome sequences of <i>Debregeasia orientalis</i> (Urticaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 1830-1831.	0.2	4
77	Pollination-Induced Transcriptome and Phylogenetic Analysis in <i>Cymbidium tortisepalum</i> (Orchidaceae). <i>Russian Journal of Plant Physiology</i> , 2019, 66, 618-627.	0.5	3
78	Forest community assembly is driven by different strata-dependent mechanisms along an elevational gradient. <i>Journal of Biogeography</i> , 2019, 46, 2174-2187.	1.4	32
79	Genome Sequences Provide Insights into the Reticulate Origin and Unique Traits of Woody Bamboos. <i>Molecular Plant</i> , 2019, 12, 1353-1365.	3.9	116
80	Greater than the sum of the parts: how the species composition in different forest strata influence ecosystem function. <i>Ecology Letters</i> , 2019, 22, 1449-1461.	3.0	51
81	Rapid diversification of alpine bamboos associated with the uplift of the Hengduan Mountains. <i>Journal of Biogeography</i> , 2019, 46, 2678-2689.	1.4	52
82	Development of 20 chloroplast microsatellite primers in wuyao (<i>Lindera aggregata</i> , Lauraceae). <i>Applications in Plant Sciences</i> , 2019, 7, e01213.	0.8	2
83	Plastid phylogenomics and biogeographic analysis support a trans-Tethyan origin and rapid early radiation of Cornales in the Mid-Cretaceous. <i>Molecular Phylogenetics and Evolution</i> , 2019, 140, 106601.	1.2	37
84	Characterization of 30 microsatellite markers in distylous <i>Primula sinolisteri</i> (Primulaceae) using HiSeq sequencing. <i>Applications in Plant Sciences</i> , 2019, 7, e01208.	0.8	3
85	Genome assembly of a tropical maize inbred line provides insights into structural variation and crop improvement. <i>Nature Genetics</i> , 2019, 51, 1052-1059.	9.4	202
86	Phylogenomic analyses reveal intractable evolutionary history of a temperate bamboo genus (Poaceae: <i>Tylotachya</i>). <i>Evolution</i> , 2019, 73, 1874-1884.	1.8	14
87	Using nuclear loci and allelic variation to disentangle the phylogeny of <i>Phyllostachys</i> (Poaceae). <i>Evolution</i> , 2019, 73, 1783-1793.	1.2	13
88	PGA: a software package for rapid, accurate, and flexible batch annotation of plastomes. <i>Plant Methods</i> , 2019, 15, 50.	1.9	660
89	Phylogenomic analysis reveals multiple evolutionary origins of selfing from outcrossing in a lineage of heterostylous plants. <i>New Phytologist</i> , 2019, 224, 1290-1303.	3.5	35
90	Origin of angiosperms and the puzzle of the Jurassic gap. <i>Nature Plants</i> , 2019, 5, 461-470.	4.7	467

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91	Differential Quaternary dynamics of evergreen broadleaved forests in subtropical China revealed by phylogeography of <i>Lindera aggregata</i> (Lauraceae). <i>Journal of Biogeography</i> , 2019, 46, 1112-1123.	1.4	20
92	Allopolyploidy in the Wintergreen Group of tribe Gaultherieae (Ericaceae) inferred from low-copy nuclear genes. <i>Nordic Journal of Botany</i> , 2019, 37, .	0.2	6
93	Why is fruit colour so variable? Phylogenetic analyses reveal relationships between fruit-colour evolution, biogeography and diversification. <i>Global Ecology and Biogeography</i> , 2019, 28, 891-903.	2.7	30
94	Upward elevation and northwest range shifts for alpine <i>Meconopsis</i> species in the Himalaya-Hengduan Mountains region. <i>Ecology and Evolution</i> , 2019, 9, 4055-4064.	0.8	52
95	Plastid phylogenomic insights into the evolution of Caryophyllales. <i>Molecular Phylogenetics and Evolution</i> , 2019, 134, 74-86.	1.2	101
96	Prevalence of isomeric plastomes and effectiveness of plastome super-barcodes in yews (<i>Taxus</i>) worldwide. <i>Scientific Reports</i> , 2019, 9, 2773.	1.6	54
97	Distributional responses to climate change for alpine species of <i>Cyananthus</i> and <i>Primula</i> endemic to the Himalaya-Hengduan Mountains. <i>Plant Diversity</i> , 2019, 41, 26-32.	1.8	30
98	Specificity and seasonal prevalence of anther smut disease <i>Microbotryum</i> on sympatric Himalayan <i>Silene</i> species. <i>Journal of Evolutionary Biology</i> , 2019, 32, 451-462.	0.8	5
99	Genomic analysis reveals rich genetic variation and potential targets of selection during domestication of castor bean from perennial woody tree to annual semi-woody crop. <i>Plant Direct</i> , 2019, 3, e00173.	0.8	12
100	Incomplete reproductive isolation between <i>Rhododendron</i> taxa enables hybrid formation and persistence. <i>Journal of Integrative Plant Biology</i> , 2019, 61, 433-448.	4.1	20
101	Evolutionary constraints on disparity of ericaceous pollen grains. <i>Annals of Botany</i> , 2019, 123, 805-813.	1.4	2
102	The topological differences between visitation and pollen transport networks: a comparison in species rich communities of the Himalaya-Hengduan Mountains. <i>Oikos</i> , 2019, 128, 551-562.	1.2	24
103	Advances in the evolution of plastid genome structure in lycophytes and ferns. <i>Biodiversity Science</i> , 2019, 27, 1172-1183.	0.2	3
104	Evolution of Angiosperm Pollen. 7. Nitrogen-fixing Clade. <i>Annals of the Missouri Botanical Garden</i> , 2019, 104, 171-229.	1.3	7
105	<i>Bulbophyllum reflexipetalum</i> (Orchidaceae, Epidendroideae, Malaxideae), a new species from Xizang, China. <i>PhytoKeys</i> , 2019, 130, 33-39.	0.4	4
106	<i>Yushania tongpeii</i> (Poaceae, Bambusoideae), a new bamboo species from north-eastern Yunnan, China. <i>PhytoKeys</i> , 2019, 130, 135-141.	0.4	2
107	Taxonomic and nomenclatural notes on <i>Pedicularis</i> (Orobanchaceae): I. One new species from northwest Yunnan, China. <i>PhytoKeys</i> , 2019, 130, 205-215.	0.4	3
108	China's biodiversity hotspots revisited: A treasure chest for plants. <i>PhytoKeys</i> , 2019, 130, 1-24.	0.4	22

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109	<i>Dendrocalamus menghanensis</i> (Poaceae, Bambusoideae), a new woody bamboo from Yunnan, China. <i>PhytoKeys</i> , 2019, 130, 143-150.	0.4	2
110	<i>Marsdenia yarlungzangboensis</i> (Apocynaceae, Asclepiadoideae), a new species from Xizang, China. <i>PhytoKeys</i> , 2019, 130, 85-92.	0.4	2
111	Functional trade-offs and the phylogenetic dispersion of seed traits in a biodiversity hotspot of the Mountains of Southwest China. <i>Ecology and Evolution</i> , 2018, 8, 2218-2230.	0.8	10
112	Impact of pre- and post-pollination barriers on pollen transfer and reproductive isolation among three sympatric <i>Pedicularis</i> (Orobanchaceae) species. <i>Plant Biology</i> , 2018, 20, 662-673.	1.8	16
113	Comparative intra- and interspecific sexual organ reciprocity in four distylous <i>Primula</i> species in the Himalaya-Hengduan Mountains. <i>Plant Biology</i> , 2018, 20, 643-653.	1.8	2
114	Plastome Phylogenetics: 30 Years of Inferences Into Plant Evolution. <i>Advances in Botanical Research</i> , 2018, , 293-313.	0.5	64
115	Evolution of Angiosperm Pollen. 5. Early Diverging Superasteridae (Berberidopsidales, Caryophyllales). <i>Trends in Plant Science</i> , 2018, 103, 106-161.	1.3	5
116	A comparison of different methods for preserving plant molecular materials and the effect of degraded DNA on ddRAD sequencing. <i>Plant Diversity</i> , 2018, 40, 106-116.	1.8	8
117	Plastome characteristics of Cannabaceae. <i>Plant Diversity</i> , 2018, 40, 127-137.	1.8	31
118	Does reproductive isolation reflect the segregation of color forms in <i>Spiranthes sinensis</i> (Pers.) Ames complex (Orchidaceae) in the Chinese Himalayas?. <i>Ecology and Evolution</i> , 2018, 8, 5455-5469.	0.8	19
119	Taxonomic studies on <i>Zingiber</i> (Zingiberaceae) in China IV: <i>Z. pauciflorum</i> sp. nov. from Yunnan. <i>Nordic Journal of Botany</i> , 2018, 36, njb-01534.	0.2	3
120	Warming-induced upward migration of the alpine treeline in the Changbai Mountains, northeast China. <i>Global Change Biology</i> , 2018, 24, 1256-1266.	4.2	81
121	Genetic structure and differentiation in <i>Dendrocalamus sinicus</i> (Poaceae: Bambusoideae) populations provide insight into evolutionary history and speciation of woody bamboos. <i>Scientific Reports</i> , 2018, 8, 16933.	1.6	18
122	Protect Third Pole's fragile ecosystem. <i>Science</i> , 2018, 362, 1368-1368.	6.0	76
123	Comparative transcriptomics identifies patterns of selection in roses. <i>BMC Plant Biology</i> , 2018, 18, 371.	1.6	13
124	Taxonomic studies on <i>Zingiber</i> (Zingiberaceae) in China VI: <i>Z. leucochilum</i> , a new species with running rhizome from Sichuan. <i>Nordic Journal of Botany</i> , 2018, 36, e01840.	0.2	4
125	Phylogenetic approaches resolve taxonomical confusion in <i>Pedicularis</i> (Orobanchaceae): Reinstatement of <i>Pedicularis delavayi</i> and discovering a new species <i>Pedicularis milliana</i> . <i>PLoS ONE</i> , 2018, 13, e0200372.	1.1	4
126	Evolution of Angiosperm Pollen. 6. The Celastrales, Oxalidales, and Malpighiales (Com) Clade and Zygophyllales. <i>Annals of the Missouri Botanical Garden</i> , 2018, 103, 393-442.	1.3	8

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127	Transcriptome analysis reveals crucial genes involved in the biosynthesis of nervonic acid in woody <i>Malania oleifera</i> oilseeds. <i>BMC Plant Biology</i> , 2018, 18, 247.	1.6	31
128	Complete chloroplast genome sequences of two <i>Boehmeria</i> species (Urticaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2018, 3, 937-938.	0.2	3
129	Integrating a comprehensive <sc>DNA</sc> barcode reference library with a global map of yews (<i>Taxus</i> L.) for forensic identification. <i>Molecular Ecology Resources</i> , 2018, 18, 1115-1131.	2.2	38
130	<sc>DNA</sc> barcoding herbaceous and woody plant species at a subalpine forest dynamics plot in Southwest China. <i>Ecology and Evolution</i> , 2018, 8, 7195-7205.	0.8	14
131	Plastid Genome Evolution in the Early-Diverging Legume Subfamily Cercidoideae (Fabaceae). <i>Frontiers in Plant Science</i> , 2018, 9, 138.	1.7	97
132	The Hemiparasitic Plant <i>Phtheirospermum</i> (Orobanchaceae) Is Polyphyletic and Contains Cryptic Species in the Hengduan Mountains of Southwest China. <i>Frontiers in Plant Science</i> , 2018, 9, 142.	1.7	38
133	Genome skimming herbarium specimens for DNA barcoding and phylogenomics. <i>Plant Methods</i> , 2018, 14, 43.	1.9	132
134	The first complete plastid genome of <i>Burmannia disticha</i> L. from the mycoheterotrophic monocot family Burmanniaceae. <i>Plant Diversity</i> , 2018, 40, 232-237.	1.8	5
135	Differential expression networks and inheritance patterns of long non-coding <sc>RNA</sc>s in castor bean seeds. <i>Plant Journal</i> , 2018, 95, 324-340.	2.8	43
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139	Plastid phylogenomics and adaptive evolution of <i>Gaultheria</i> series <i>Trichophyllae</i> (Ericaceae), a clade from sky islands of the Himalaya-Hengduan Mountains. <i>Molecular Phylogenetics and Evolution</i> , 2017, 110, 7-18.	1.2	26
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144	Asymmetrical natural hybridization varies among hybrid swarms between two diploid <i>Rhododendron</i> species. <i>Annals of Botany</i> , 2017, 120, 51-61.	1.4	28

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147	Breeding system and pollination of two closely related bamboo species. <i>AoB PLANTS</i> , 2017, 9, plx021.	1.2	13
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157	Comparative chloroplast genomes of eleven <i>Schima</i> (Theaceae) species: Insights into DNA barcoding and phylogeny. <i>PLoS ONE</i> , 2017, 12, e0178026.	1.1	34
158	Negative correlation between rates of molecular evolution and flowering cycles in temperate woody bamboos revealed by plastid phylogenomics. <i>BMC Plant Biology</i> , 2017, 17, 260.	1.6	27
159	Domestication Origin and Breeding History of the Tea Plant (<i>Camellia sinensis</i>) in China and India Based on Nuclear Microsatellites and cpDNA Sequence Data. <i>Frontiers in Plant Science</i> , 2017, 8, 2270.	1.7	71
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180	The Tree of Life: China project. Journal of Systematics and Evolution, 2016, 54, 273-276.	1.6	7

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183	<i>Gaultheria marronina</i> sp. nov. (Ericaceae) from Sichuan, China. <i>Nordic Journal of Botany</i> , 2016, 34, 545-549.	0.2	2
184	Nuclear genetic variation of <i>Rosa odorata</i> var. <i>gigantea</i> (Rosaceae): population structure and conservation implications. <i>Tree Genetics and Genomes</i> , 2016, 12, 1.	0.6	13
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196	<i>Pseudobartsia glandulosa</i> , a new combination to replace <i>Pseudobartsia yunnanensis</i> (Orobanchaceae). <i>Phytotaxa</i> , 2015, 217, 197.	0.1	7
197	Typification of seven Chinese species of <i>Pedicularis</i> (Orobanchaceae) described by Bureau and Franchet with taxonomic notes. <i>Plant Ecology and Evolution</i> , 2015, 148, 144-148.	0.3	0
198	Floral nectary morphology and evolution in <i>Pedicularis</i> (Orobanchaceae). <i>Botanical Journal of the Linnean Society</i> , 2015, 178, 592-607.	0.8	9

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203	Investigating the MicroRNAs of Two Developmental Phases of <i>Dendrocalamus latiflorus</i> (Poaceae: Tj ETQq1 1 0,784314 ₆ gBT /Over	1.0	1
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218	Factors affecting stress tolerance in recalcitrant embryonic axes from seeds of four <i>Quercus</i> (Fagaceae) species native to the USA or China. <i>Annals of Botany</i> , 2014, 114, 1747-1759.	1.4	39
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220	The evolution of floral deception in <i>Epipactis veratrifolia</i> (Orchidaceae): from indirect defense to pollination. <i>BMC Plant Biology</i> , 2014, 14, 63.	1.6	27
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223	Should genes with missing data be excluded from phylogenetic analyses?. <i>Molecular Phylogenetics and Evolution</i> , 2014, 80, 308-318.	1.2	109
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243	Molecular systematics of <i>Dendrobium</i> (Orchidaceae, Dendrobieae) from mainland Asia based on plastid and nuclear sequences. <i>Molecular Phylogenetics and Evolution</i> , 2013, 69, 950-960.	1.2	96
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245	Molecular phylogenetics and character evolution of Cannabaceae. <i>Taxon</i> , 2013, 62, 473-485.	0.4	85
246	Comparative phylogeography of two sympatric beeches in subtropical China: Species-specific geographic mosaic of lineages. <i>Ecology and Evolution</i> , 2013, 3, 4461-4472.	0.8	34
247	Floral ontogeny of <i>Pedicularis</i> (Orobanchaceae), with an emphasis on the corolla upper lip. <i>Journal of Systematics and Evolution</i> , 2013, 51, 435-450.	1.6	7
248	<i>Gymnosporia thyrsoflora</i> comb. nov. (Celastraceae), a correct name to replace <i>G. graciliramula</i> from southwest China. <i>Nordic Journal of Botany</i> , 2013, 31, 746-747.	0.2	0
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250	(2204) Proposal to conserve <i>Pedicularis stenocorys</i> against <i>P. stenantha</i> (Orobanchaceae). <i>Taxon</i> , 2013, 62, 1066-1067.	0.4	2
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254	Developmental Genetics of the Perianthless Flowers and Bracts of a Paleoherb Species, <i>Saururus chinensis</i> . PLoS ONE, 2013, 8, e53019.	1.1	4
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257	Sampling Strategy and Potential Utility of Indels for DNA Barcoding of Closely Related Plant Species: A Case Study in <i>Taxus</i> . International Journal of Molecular Sciences, 2012, 13, 8740-8751.	1.8	46
258	Genetic Diversity and Population Structure: Implications for Conservation of Wild Soybean (<i>Glycine</i>) of Molecular Sciences, 2012, 13, 12608-12628.	1.8	60
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262	De Novo Sequencing and Characterization of the Floral Transcriptome of <i>Dendrocalamus latiflorus</i> (Poaceae: Bambusoideae). PLoS ONE, 2012, 7, e42082.	1.1	111
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269	Monophyly or Paraphyly? The Taxonomy of <i>Holcoglossum</i> (Aeridinae: Orchidaceae). PLoS ONE, 2012, 7, e52050.	1.1	13
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281	<i>Elatostema densistriolatum</i> sp. nov., <i>E. latistipulum</i> sp. nov. and <i>E. cyrtandrifolium</i> var. <i>hirsutum</i> var. nov. (Urticaceae) from southwest China. <i>Nordic Journal of Botany</i> , 2011, 29, 227-232.	0.2	9
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