Chun-Neng Wang

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Stress associated proteins coordinate the activation of comprehensive antiviral immunity in <i>Phalaenopsis</i> orchids. New Phytologist, 2022, 233, 145-155.	7.3	4
2	Transcriptomic Analysis Suggests Auxin Regulation in Dorsal-Ventral Petal Asymmetry of Wild Progenitor Sinningia speciosa. International Journal of Molecular Sciences, 2022, 23, 2073.	4.1	2
3	Development of a petal protoplast transfection system for <i>Sinningia speciosa</i> . Applications in Plant Sciences, 2022, 10, .	2.1	1
4	Evolution of Terpene Synthases in Orchidaceae. International Journal of Molecular Sciences, 2021, 22, 6947.	4.1	19
5	Floral Color Diversity: How Are Signals Shaped by Elevational Gradient on the Tropical–Subtropical Mountainous Island of Taiwan?. Frontiers in Plant Science, 2020, 11, 582784.	3.6	11
6	Silencing of PhLA, a CIN-TCP gene, causes defected petal conical epidermal cell formation and results in reflexed corolla lobes in petunia. , 2020, 61, 24.		2
7	QUANTITATIVE ASSESSMENT OF ANISOCOTYLY IN HABERLEA RHODOPENSIS AND RAMONDA MYCONI. Edinburgh Journal of Botany, 2019, 76, 377-391.	0.4	0
8	Gene duplication and relaxation from selective constraints of GCYC genes correlated with various floral symmetry patterns in Asiatic Gesneriaceae tribe Trichosporeae. PLoS ONE, 2019, 14, e0210054.	2.5	5
9	Effective pollination of Aeschynanthus acuminatus (Gesneriaceae) by generalist passerines, in sunbird-absent East Asia. Scientific Reports, 2019, 9, 17552.	3.3	5
10	Quantifying colour and spot characteristics for the ventral petals in Sinningia speciosa. Biosystems Engineering, 2018, 167, 40-50.	4.3	5
11	Morphological characterization of infraâ€generic lineages in <i>Deparia</i> (Athyriaceae: Polypodiales). Cladistics, 2018, 34, 78-92.	3.3	9
12	Expression shifts of floral symmetry genes correlate to flower actinomorphy in East Asia endemic Conandron ramondioides (Gesneriaceae). , 2018, 59, 24.		9
13	Genetic Analysis of Floral Symmetry Transition in African Violet Suggests the Involvement of Trans-acting Factor for CYCLOIDEA Expression Shifts. Frontiers in Plant Science, 2018, 9, 1008.	3.6	13
14	Shoot regeneration process and optimization of Agrobacterium-mediated transformation in Sinningia speciosa. Plant Cell, Tissue and Organ Culture, 2018, 134, 301-316.	2.3	3
15	Integrating a comprehensive <scp>DNA</scp> barcode reference library with a global map of yews (<i>Taxus</i> L.) for forensic identification. Molecular Ecology Resources, 2018, 18, 1115-1131.	4.8	38
16	Phylogenetic analyses of Bradyrhizobium symbionts associated with invasive Crotalaria zanzibarica and its coexisting legumes in Taiwan. Systematic and Applied Microbiology, 2018, 41, 619-628.	2.8	5
17	Organelle Genome Inheritance in Deparia Ferns (Athyriaceae, Aspleniineae, Polypodiales). Frontiers in Plant Science, 2018, 9, 486.	3.6	21
18	Infrageneric Revision of the Fern Genus Deparia (Athyriaceae, Aspleniineae, Polypodiales). Systematic Botany, 2018, 43, 645-655.	0.5	4

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19	The Contribution of Neutral and Environmentally Dependent Processes in Driving Population and Lineage Divergence in Taiwania (Taiwania cryptomerioides). Frontiers in Plant Science, 2018, 9, 1148.	3.6	27
20	From shoot to leaf: step-wise shifts in meristem and KNOX1 activity correlate with the evolution of a unifoliate body plan in Gesneriaceae. Development Genes and Evolution, 2017, 227, 41-60.	0.9	10
21	Cytokinin biosynthesis ISOPENTENYLTRANSFERASE genes are differentially expressed during phyllomorph development in the acaulescent Streptocarpus rexii (Gesneriaceae). South African Journal of Botany, 2017, 109, 96-111.	2.5	4
22	A Taxonomic Revision of the Kerivoula hardwickii Complex (Chiroptera: Vespertilionidae) with the Description of a New Species. Acta Chiropterologica, 2017, 19, 19.	0.6	14
23	The evolution of genome size and distinct distribution patterns of rDNA in Phalaenopsis (Orchidaceae). Botanical Journal of the Linnean Society, 2017, 185, 65-80.	1.6	17
24	Association between Petal Form Variation and CYC2-like Genotype in a Hybrid Line of Sinningia speciosa. Frontiers in Plant Science, 2017, 8, 558.	3.6	31
25	Genetic Diversity of a Novel Oil Crop, Camellia brevistyla, Revealed by ISSR DNA Markers. Horticultural Science and Technology, 2017, 35, 588-598.	0.6	3
26	A Revised Framework of <i>Dryopteris</i> Subg. <i>Nothoperanema</i> (Dryopteridaceae) Inferred from Phylogenetic Evidence, with Descriptions of Two New Sections. Systematic Botany, 2016, 41, 596-605.	0.5	9
27	Historical biogeography of the fern genus Deparia (Athyriaceae) and its relation with polyploidy. Molecular Phylogenetics and Evolution, 2016, 104, 123-134.	2.7	36
28	Quantifying floral shape variation in 3D using microcomputed tomography: a case study of a hybrid line between actinomorphic and zygomorphic flowers. Frontiers in Plant Science, 2015, 6, 724.	3.6	45
29	MicroRNA396-Targeted <i>SHORT VEGETATIVE PHASE</i> Is Required to Repress Flowering and Is Related to the Development of Abnormal Flower Symptoms by the Phyllody Symptoms1 Effector. Plant Physiology, 2015, 168, 1702-1716.	4.8	44
30	Quantitative analysis of floral symmetry and tube dilation in an F2 cross of Sinningia speciosa. Scientia Horticulturae, 2015, 188, 71-77.	3.6	23
31	Three-dimensional Shape Variation Analysis on Sinningia Speciosa Flowers. , 2014, , .		0
32	Flower development of <i>Phalaenopsis</i> orchid involves functionally divergent <i><scp>SEPALLATA</scp></i> â€like genes. New Phytologist, 2014, 202, 1024-1042.	7.3	113
33	CA2 and GA20-oxidase expressions are associated with the meristem position in Streptocarpus rexii (Gesneriaceae). Plant Growth Regulation, 2014, 72, 123-140.	3.4	6
34	The Hybrid Origin of <1>Adiantum meishanianum 1 (Pteridaceae): A Rare and Endemic Species in Taiwan. Systematic Botany, 2014, 39, 1034-1041.	0.5	16
35	Historical connectivity, contemporary isolation and local adaptation in a widespread but discontinuously distributed species endemic to Taiwan, Rhododendron oldhamii (Ericaceae). Heredity, 2013, 111, 147-156.	2.6	60
36	Quantitative Evaluation of the Floral Shape Variation in Sinningia Speciosa Domestication. , 2013, , .		0

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37	Gibberellin as a suppressor of lateral dominance and inducer of apical growth in the unifoliateStreptocarpus wendlandii(Gesneriaceae). New Zealand Journal of Botany, 2012, 50, 267-287.	1.1	6
38	Development of PCR primer sets for intron 1 of the low-copy gene LEAFY in Davalliaceae. American Journal of Botany, 2012, 99, e223-e225.	1.7	11
39	Light as environmental regulator for germination and macrocotyledon development in Streptocarpus rexii (Gesneriaceae). South African Journal of Botany, 2012, 81, 50-60.	2.5	13
40	Refugia and phylogeography of Taiwania in East Asia. Journal of Biogeography, 2011, 38, 1992-2005.	3.0	82
41	First insights into fern matK phylogeny. Molecular Phylogenetics and Evolution, 2011, 59, 556-566.	2.7	127
42	A complex case of simple leaves: indeterminate leaves co-express ARP and KNOX1 genes. Development Genes and Evolution, 2010, 220, 25-40.	0.9	22
43	Tissueâ€direct PCR, a rapid and extractionâ€free method for barcoding of ferns. Molecular Ecology Resources, 2010, 10, 92-95.	4.8	37
44	Identifying a mysterious aquatic fern gametophyte. Plant Systematics and Evolution, 2009, 281, 77-86.	0.9	44
45	15-P013 Inheritance and molecular genetics of floral symmetry in Darwin's Gloxinia peloria (Sinningia) Tj ETÇ	2q110.78	4314 rgBT 0
46	Molecular population genetics and gene expression analysis of duplicated CBF genes of Arabidopsis		
	thaliana. BMC Plant Biology, 2008, 8, 111.	3.6	22
47	thaliana. BMC Plant Biology, 2008, 8, 111. A Heat-Inducible Transcription Factor, HsfA2, Is Required for Extension of Acquired Thermotolerance in Arabidopsis. Plant Physiology, 2007, 143, 251-262.	3.6 4.8	22 551
47 48	 thaliana. BMC Plant Biology, 2008, 8, 111. A Heat-Inducible Transcription Factor, HsfA2, Is Required for Extension of Acquired Thermotolerance in Arabidopsis. Plant Physiology, 2007, 143, 251-262. Population Genetic Structure of Titanotrichum oldhamii (Gesneriaceae), a Subtropical Bulbiliferous Plant with Mixed Sexual and Asexual Reproduction. Annals of Botany, 2004, 93, 201-209. 	3.6 4.8 2.9	22 551 30
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47 48 49 50	 thaliana. BMC Plant Biology, 2008, 8, 111. A Heat-Inducible Transcription Factor, HsfA2, Is Required for Extension of Acquired Thermotolerance in Arabidopsis. Plant Physiology, 2007, 143, 251-262. Population Genetic Structure of Titanotrichum oldhamii (Gesneriaceae), a Subtropical Bulbiliferous Plant with Mixed Sexual and Asexual Reproduction. Annals of Botany, 2004, 93, 201-209. Aspects of sexual failure in the reproductive processes of a rare bulbiliferous plant, Titanotrichum oldhamii (Gesneriaceae), in subtropical Asia. Sexual Plant Reproduction, 2004, 17, 23-31. Altered expression of GFLO , the Gesneriaceae homologue of FLORICAULA/LEAFY , is associated with the transition to bulbil formation in Titanotrichum oldhamii. Development Genes and Evolution, 2004, 214, 122-127. 	 3.6 4.8 2.9 2.2 0.9 	22 551 30 11 36
47 48 49 50 51	 thaliana. BMC Plant Biology, 2008, 8, 111. A Heat-Inducible Transcription Factor, HsfA2, Is Required for Extension of Acquired Thermotolerance in Arabidopsis. Plant Physiology, 2007, 143, 251-262. Population Genetic Structure of Titanotrichum oldhamii (Gesneriaceae), a Subtropical Bulbiliferous Plant with Mixed Sexual and Asexual Reproduction. Annals of Botany, 2004, 93, 201-209. Aspects of sexual failure in the reproductive processes of a rare bulbiliferous plant, Titanotrichum oldhamii (Gesneriaceae), in subtropical Asia. Sexual Plant Reproduction, 2004, 17, 23-31. Altered expression of GFLO, the Gesneriaceae homologue of FLORICAULA/LEAFY, is associated with the transition to bulbil formation in Titanotrichum oldhamii. Development Genes and Evolution, 2004, 214, 122-127. Phylogenetic Position of Titanotrichum oldhamii (Gesneriaceae) Inferred From Four Different Gene Regions. Systematic Botany, 2004, 29, 407-418. 	 3.6 4.8 2.9 2.2 0.9 0.5 	22 551 30 11 36 55