

Lin Chen

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1347315/publications.pdf>

Version: 2024-02-01

17
papers

64
citations

1937685

4
h-index

1720034

7
g-index

18
all docs

18
docs citations

18
times ranked

56
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic Diversity and Population Structure Revealed by SSR Markers on Endemic Species <i>Osmanthus serrulatus</i> Rehd. from Southwestern Sichuan Basin, China. <i>Forests</i> , 2021, 12, 1365.	2.1	4
2	SWATH-MS based quantitative proteomics reveal regulatory metabolism and networks of androdioecy breeding system in <i>Osmanthus fragrans</i> . <i>BMC Plant Biology</i> , 2021, 21, 468.	3.6	1
3	The genome of Chinese flowering cherry (<i>Cerasus serrulata</i>) provides new insights into <i>Cerasus</i> species. <i>Horticulture Research</i> , 2020, 7, 165.	6.3	22
4	The complete chloroplast genome of <i>Changnienia amoena</i> S. S. Chien (Orchidaceae) and its phylogenetic implication. <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 238-239.	0.4	0
5	Revisiting the phylogeny and taxonomy of <i>Osmanthus</i> (Oleaceae) including description of the new genus <i>Chengiodendron</i> . <i>Phytotaxa</i> , 2020, 436, 283-292.	0.3	8
6	<i>Stellaria multipartita</i> (Caryophyllaceae), a new species from Chongqing, China. <i>Phytotaxa</i> , 2020, 442, 196-204.	0.3	1
7	The complete chloroplast genome of <i>Fraxinus pennsylvanica</i> (Oleaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 1932-1933.	0.4	2
8	The complete chloroplast genome of <i>Pinus yunnanensis</i> Franchet (Pinaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 2600-2601.	0.4	2
9	The complete chloroplast genome of <i>Osmanthus serrulatus</i> rehd., a natural sweet olive endemic to China. <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 3396-3397.	0.4	0
10	The complete chloroplast genome of <i>Sorbus folgneri</i> (C.K.Schneid.) Rehd. (Rosaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 728-729.	0.4	4
11	Characterization of the complete chloroplast genome sequence of <i>Tsuga longibracteata</i> W. C. Cheng (Pinaceae). <i>Conservation Genetics Resources</i> , 2019, 11, 117-120.	0.8	7
12	Microsatellites characteristics analysis and SSR marker development for <i>Paphiopedilum hirsutissimum</i> based on transcriptome sequencing. <i>Plant Genetic Resources: Characterisation and Utilisation</i> , 2018, 16, 394-396.	0.8	1
13	Spatial distribution and environmental interpretation of wild <i>Sinojackia xylocarpa</i> communities based on self-organizing map (SOM). <i>Biodiversity Science</i> , 2018, 26, 1268-1276.	0.6	0
14	De novo transcriptome analysis of <i>Osmanthus serrulatus</i> Rehd. flowers and leaves by Illumina sequencing. <i>Biochemical Systematics and Ecology</i> , 2015, 61, 531-540.	1.3	7
15	<i>Zingiber nanlingensis</i> sp. nov. (Zingiberaceae) from Guangdong, China. <i>Nordic Journal of Botany</i> , 2011, 29, 431-434.	0.5	2
16	<i>Impatiens nanlingensis</i> sp. nov. (Balsaminaceae) from Guangdong, China. <i>Nordic Journal of Botany</i> , 2009, 27, 475-477.	0.5	2
17	<i>Microtropis shenzhenensis</i> sp. nov. (Celastraceae) from Guangdong, China. <i>Nordic Journal of Botany</i> , 2009, 27, 469-471.	0.5	1