

Yiling Wang

List of Publications by Year in descending order

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Version: 2024-02-01

8
papers

139
citations

1478505

6
h-index

1588992

8
g-index

11
all docs

11
docs citations

11
times ranked

64
citing authors

#	ARTICLE	IF	CITATIONS
1	Localized environmental heterogeneity drives the population differentiation of two endangered and endemic <i>Opisthopappus</i> Shih species. <i>Bmc Ecology and Evolution</i> , 2021, 21, 56.	1.6	11
2	Demographic history and genetic differentiation of an endemic and endangered <i>Ulmus lamellosa</i> (<i>Ulmus</i>). <i>BMC Plant Biology</i> , 2020, 20, 526.	3.6	7
3	Population genetic variation characterization of the boreal tree <i>Acer ginnala</i> in Northern China. <i>Scientific Reports</i> , 2020, 10, 13515.	3.3	1
4	De novo Assembly and Transcriptome Characterization of <i>Opisthopappus</i> (<i>Asteraceae</i>) for Population Differentiation and Adaption. <i>Frontiers in Genetics</i> , 2018, 9, 371.	2.3	13
5	Leaves Micromorphological Characteristics of <i>Opisthopappus taihangensis</i> and <i>Opisthopappus longilobus</i> from Taihang Mountain, China. <i>Vegetos</i> , 2015, 28, 82.	1.5	14
6	Molecular Phylogeography and Population Genetic Structure of <i>O. longilobus</i> and <i>O. taihangensis</i> (<i>Opisthopappus</i>) on the Taihang Mountains. <i>PLoS ONE</i> , 2014, 9, e104773.	2.5	38
7	Chloroplast microsatellite diversity of <i>Opisthopappus</i> Shih (<i>Asteraceae</i>) endemic to China. <i>Plant Systematics and Evolution</i> , 2013, 299, 1849-1858.	0.9	31
8	Genetic diversity and population structure of <i>Opisthopappus longilobus</i> and <i>Opisthopappus taihangensis</i> (<i>Asteraceae</i>) in China determined using sequence related amplified polymorphism markers. <i>Biochemical Systematics and Ecology</i> , 2013, 49, 115-124.	1.3	24