Wei Wu

List of Publications by Year in descending order

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31	1,063	18	32
papers	citations	h-index	g-index
39	39	39	1393
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	<i>SAUR36,</i> a SMALL AUXIN UP RNA Gene, Is Involved in the Promotion of Leaf Senescence in Arabidopsis Â. Plant Physiology, 2013, 161, 1002-1009.	2.3	173
2	Predicting the current and future cultivation regions of Carthamus tinctorius L. using MaxEnt model under climate change in China. Global Ecology and Conservation, 2018, 16, e00477.	1.0	96
3	Fungal endophyte-derived Fritillaria unibracteata var. wabuensis: diversity, antioxidant capacities in vitro and relations to phenolic, flavonoid or saponin compounds. Scientific Reports, 2017, 7, 42008.	1.6	75
4	Effects of Salt Stress on the Growth, Physiological Responses, and Glycoside Contents of Stevia rebaudiana Bertoni. Journal of Agricultural and Food Chemistry, 2013, 61, 5720-5726.	2.4	62
5	Genetic diversity and relationships among safflower (Carthamus tinctorius L.) analyzed by inter-simple sequence repeats (ISSRs). Genetic Resources and Crop Evolution, 2007, 54, 1043-1051.	0.8	61
6	RNA-Seq for gene identification and transcript profiling of three Stevia rebaudiana genotypes. BMC Genomics, 2014, 15, 571.	1.2	55
7	Variation of essential oil of Mentha haplocalyx Briq. and Mentha spicata L. from China. Industrial Crops and Products, 2013, 42, 251-260.	2.5	44
8	Fusarium redolens 6WBY3, an endophytic fungus isolated from Fritillaria unibracteata var. wabuensis, produces peimisine and imperialine-3β-d-glucoside. Fìtoterapìâ, 2015, 103, 213-221.	1.1	42
9	Integrating transcriptomics and metabolomics to studies key metabolism, pathways and candidate genes associated with drought-tolerance in Carthamus tinctorius L. Under drought stress. Industrial Crops and Products, 2020, 151, 112465.	2.5	40
10	Essential oil variations in different Perilla L. accessions: chemotaxonomic implications. Plant Systematics and Evolution, 2009, 281, 1-10.	0.3	34
11	Optimization of potassium for proper growth and physiological response of Houttuynia cordata Thunb Environmental and Experimental Botany, 2011, 71, 292-297.	2.0	34
12	Molecular Cloning and Expression Analysis of Genes Encoding Two Microsomal Oleate Desaturases (FAD2) from Safflower (Carthamus tinctorius L.). Plant Molecular Biology Reporter, 2012, 30, 139-148.	1.0	34
13	Peimisine and peiminine production by endophytic fungus Fusarium sp. isolated from Fritillaria unibracteata var. wabensis. Phytomedicine, 2014, 21, 1104-1109.	2.3	32
14	Endophytic fungus strain 28 isolated from Houttuynia cordata possesses wide-spectrum antifungal activity. Brazilian Journal of Microbiology, 2016, 47, 480-488.	0.8	31
15	Mining of simple sequence repeats (SSRs) loci and development of novel transferability-across EST-SSR markers from de novo transcriptome assembly of Angelica dahurica. PLoS ONE, 2019, 14, e0221040.	1.1	30
16	Isolation and Characterization of a Temperature-Regulated Microsomal Oleate Desaturase Gene (CtFAD2-1) from Safflower (Carthamus tinctorius L.). Plant Molecular Biology Reporter, 2012, 30, 391-402.	1.0	26
17	Extraction, purification and antioxidation of a polysaccharide from Fritillaria unibracteata var. wabuensis. International Journal of Biological Macromolecules, 2018, 112, 1073-1083.	3.6	26
18	Antioxidant Abilities, Phenolics and Flavonoids Contents in the Ethanolic Extracts of the Stems and Leaves of Different Stevia rebaudiana Bert Lines. Sugar Tech, 2013, 15, 209-213.	0.9	21

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19	Cladosins L-O, new hybrid polyketides from the endophytic fungus Cladosporium sphaerospermum WBS017. European Journal of Medicinal Chemistry, 2020, 191, 112159.	2.6	19
20	Bolting, an Important Process in Plant Development, Two Types in Plants. Journal of Plant Biology, 2019, 62, 161-169.	0.9	18
21	Characterizing glycosyltransferases by a combination of sequencing platforms applied to the leaf tissues of Stevia rebaudiana. BMC Genomics, 2020, 21, 794.	1.2	14
22	Exopolysaccharides from the fungal endophytic Fusarium sp. A14 isolated from Fritillaria unibracteata Hsiao et KC Hsia and their antioxidant and antiproliferation effects. Journal of Bioscience and Bioengineering, 2019, 127, 231-240.	1.1	13
23	Identification and Evaluation of Reference Genes for Accurate Transcription Normalization in Safflower under Different Experimental Conditions. PLoS ONE, 2015, 10, e0140218.	1.1	13
24	Mutations in the uridine diphosphate glucosyltransferase 76G1 gene result in different contents of the major steviol glycosides in Stevia rebaudiana. Phytochemistry, 2019, 162, 141-147.	1.4	12
25	Identification of the Key Residues of the Uridine Diphosphate Glycosyltransferase 91D2 and its Effect on the Accumulation of Steviol Glycosides in <i>Stevia rebaudiana</i> Journal of Agricultural and Food Chemistry, 2021, 69, 1852-1863.	2.4	12
26	C19-Diterpenoid alkaloids from the roots of Aconitum hemsleyanum var.Âhanyuanum and their chemotaxonomic significance. Biochemical Systematics and Ecology, 2010, 38, 1052-1055.	0.6	9
27	Cloning and sequence analysis of the safflower betaine aldehyde dehydrogenase gene. Genetics and Molecular Research, 2014, 13, 344-353.	0.3	7
28	Composition and variability of the essential oil of Houttuynia of China. Chemistry of Natural Compounds, 2008, 44, 778-783.	0.2	6
29	The effect of floral morphology on seed set in Carthamus tinctorius Linnaeus (Asteraceae) clones of Sichuan province in China. Plant Systematics and Evolution, 2012, 298, 59-68.	0.3	6
30	Transcriptome changes in Polygonum multiflorum Thunb. roots induced by methyl jasmonate. Journal of Zhejiang University: Science B, 2015, 16, 1027-1041.	1.3	6
31	Simultaneous determination of 10 nucleosides and nucleobases from different cultivation years of Fritillaria unibracteata var. wabuensis by HPLC-DAD. Journal of Chinese Pharmaceutical Sciences, 2017, 26	0.4	O