Yan Wan

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/6247566/publications.pdf

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15	307	1039406 9	996533 15
papers	citations	h-index	g-index

15	15	15	234
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Prospects for Proanthocyanidins from Grape Seed: Extraction Technologies and Diverse Bioactivity. Food Reviews International, 2023, 39, 349-368.	4.3	8
2	Prospects of cereal protein-derived bioactive peptides: Sources, bioactivities diversity, and production. Critical Reviews in Food Science and Nutrition, 2022, 62, 2855-2871.	5.4	34
3	Integrating transcriptome and physiological analyses to elucidate the molecular responses of buckwheat to graphene oxide. Journal of Hazardous Materials, 2022, 424, 127443.	6.5	11
4	Evaluation of morphology, nutrients, phytochemistry and pigments suggests the optimum harvest date for high-quality quinoa leafy vegetable. Scientia Horticulturae, 2022, 304, 111240.	1.7	3
5	Identification of the specific long-noncoding RNAs involved in night-break mediated flowering retardation in Chenopodium quinoa. BMC Genomics, 2021, 22, 284.	1.2	8
6	Nitrate dose-responsive transcriptome analysis identifies transcription factors and small secreted peptides involved in nitrogen response in Tartary buckwheat. Plant Physiology and Biochemistry, 2021, 162, 1-13.	2.8	7
7	Quinoa sprouts as potential vegetable source: Nutrient composition and functional contents of different quinoa sprout varieties. Food Chemistry, 2021, 357, 129752.	4.2	34
8	Genome-wide identification of genes involved in heterotrimeric G-protein signaling in Tartary buckwheat (Fagopyrum tataricum) and their potential roles in regulating fruit development. International Journal of Biological Macromolecules, 2021, 171, 435-447.	3.6	4
9	The complete mitochondrial genomes of two model ectomycorrhizal fungi (Laccaria): features, intron dynamics and phylogenetic implications. International Journal of Biological Macromolecules, 2020, 145, 974-984.	3.6	52
10	Transcriptome profiling identifies transcription factors and key homologs involved in seed dormancy and germination regulation of Chenopodium quinoa. Plant Physiology and Biochemistry, 2020, 151, 443-456.	2.8	22
11	Investigation into the underlying regulatory mechanisms shaping inflorescence architecture in Chenopodium quinoa. BMC Genomics, 2019, 20, 658.	1.2	16
12	The complete mitochondrial genomes of five important medicinal Ganoderma species: Features, evolution, and phylogeny. International Journal of Biological Macromolecules, 2019, 139, 397-408.	3.6	62
13	Post-Anthesis Photosynthetic Properties Provide Insights into Yield Potential of Tartary Buckwheat Cultivars. Agronomy, 2019, 9, 149.	1.3	15
14	Relationship between stem characteristics and lodging resistance of Tartary buckwheat (Fagopyrum tataricum). Plant Production Science, 2019, 22, 202-210.	0.9	27
15	Isoflavonoid Accumulation Pattern as Affected by Shading from Maize in Soybean (Glycine max (L.)) Tj ETQq1 1 0.784314 rgBT /Overl	0.9	4