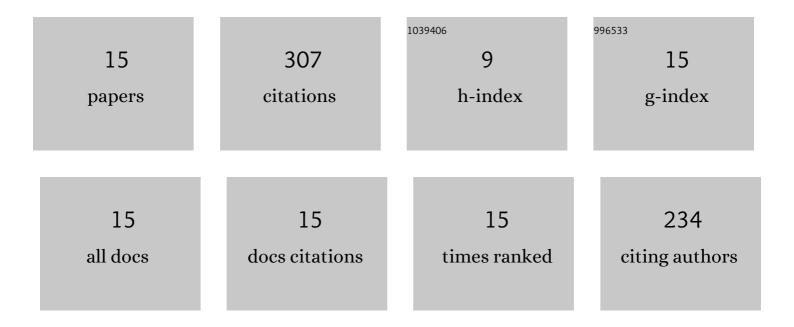
Yan Wan

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6247566/publications.pdf Version: 2024-02-01



ΥΛΝ ΜΛΛΝ

#	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 C-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 (<i>Fagopyrum tataricum</i>). 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 /Overlo