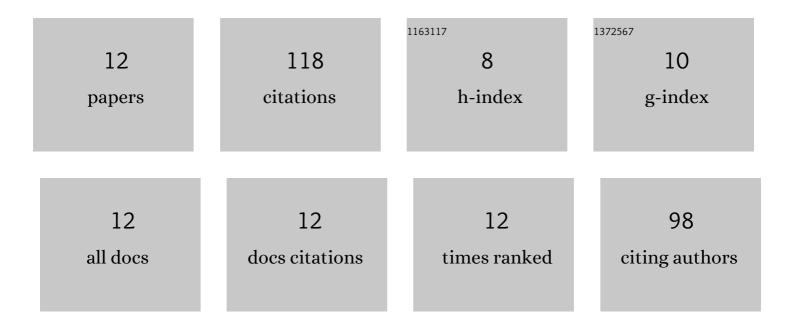
Zhiyuan Li

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8642093/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	NtNAC053, A Novel NAC Transcription Factor, Confers Drought and Salt Tolerances in Tobacco. Frontiers in Plant Science, 2022, 13, .	3.6	7
2	Systematic Analysis of the bZIP Family in Tobacco and Functional Characterization of NtbZIP62 Involvement in Salt Stress. Agronomy, 2021, 11, 148.	3.0	13
3	Bioinformatics and Expression Analysis of IDA-Like Genes Reveal Their Potential Functions in Flower Abscission and Stress Response in Tobacco (Nicotiana tabacum L.). Frontiers in Genetics, 2021, 12, 670794.	2.3	3
4	Utilization of Ogura CMS germplasm with the clubroot resistance gene by fertility restoration and cytoplasm replacement in Brassica oleracea L. Horticulture Research, 2020, 7, 61.	6.3	10
5	Kompetitive allele-specific PCR (KASP) genotyping and heterotic group classification of 244 inbred lines in cabbage (Brassica oleracea L. var. capitata). Euphytica, 2020, 216, 1.	1.2	10
6	Pigment variation and transcriptional response of the pigment synthesis pathway in the S2309 triple-color ornamental kale (Brassica oleracea L. var. acephala) line. Genomics, 2020, 112, 2658-2665.	2.9	10
7	Creation of fertility-restored materials for Ogura CMS in Brassica oleracea by introducing Rfo gene from Brassica napus via an allotriploid strategy. Theoretical and Applied Genetics, 2020, 133, 2825-2837.	3.6	11
8	Characterization and Mapping of a Novel Premature Leaf Senescence Mutant in Common Tobacco (Nicotiana tabacum L.). Plants, 2019, 8, 415.	3.5	6
9	Heterotic Group Classification of 63 Inbred Lines and Hybrid Purity Identification by Using SSR Markers in Winter Cabbage (Brassica Oleracea L. var. capitata). Horticultural Plant Journal, 2018, 4, 158-164.	5.0	13
10	Differentially Expressed Genes Associated with the Cabbage Yellow-Green-Leaf Mutant in the ygl-1 Mapping Interval with Recombination Suppression. International Journal of Molecular Sciences, 2018, 19, 2936.	4.1	9
11	Morphological and molecular characterization of the second backcross progenies of Ogu-CMS Chinese kale and rapeseed. Euphytica, 2017, 213, 1.	1.2	20
12	Systematic Analysis of Tobacco CrRLK1L Family Genes and Functional Identification of NtCrRLK1L47 in Environmental Stresses. Frontiers in Plant Science, 0, 13, .	3.6	6