

# Zhiyuan Li

## List of Publications by Year in descending order

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12  
papers

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1163117  
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docs citations

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times ranked

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