

Jingjuan Li

List of Publications by Year in descending order

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11
papers

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1307594

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375
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#	ARTICLE	IF	CITATIONS
1	Exogenous metabolites spray, which identified from metabolomics analysis and transcriptomic analysis, can improve salt tolerance of Chinese cabbages (<i>Brassica rapa</i> L.ssp <i>pekinensis</i>)*. Journal of Plant Interactions, 2021, 16, 452-461.	2.1	3
2	Comparative transcriptome analysis between a resistant and a susceptible Chinese cabbage in response to <i>Hyaloperonospora brassicae</i> . Plant Signaling and Behavior, 2020, 15, 1777373.	2.4	7
3	Genome-wide gene expression profiles in response to downy mildew in Chinese cabbage (<i>Brassica rapa</i>) Tj ETQq1 1,0,784314 rgBT /O	1.7	10
4	Identification of miRNAs and their targets in regulating tuberous root development in radish using small RNA and degradome analyses. 3 Biotech, 2018, 8, 311.	2.2	7
5	Ectopic expression of a <i>Brassica rapa</i> AINTEGUMENTA gene (BrANT-1) increases organ size and stomatal density in <i>Arabidopsis</i> . Scientific Reports, 2018, 8, 10528.	3.3	7
6	Physiological and Transcriptomic Responses of Chinese Cabbage (<i>Brassica rapa</i> L. ssp. <i>pekinensis</i>) to Salt Stress. International Journal of Molecular Sciences, 2017, 18, 1953.	4.1	28
7	Comparative Transcriptome Analysis Reveals Effects of Exogenous Hematin on Anthocyanin Biosynthesis during Strawberry Fruit Ripening. International Journal of Genomics, 2016, 2016, 1-14.	1.6	8
8	Genome-Wide Identification and Analysis of the VQ Motif-Containing Protein Family in Chinese Cabbage (<i>Brassica rapa</i> L. ssp. <i>pekinensis</i>). International Journal of Molecular Sciences, 2015, 16, 28683-28704.	4.1	43
9	Integrative Analysis of mRNA and miRNA Expression Profiles of the Tuberous Root Development at Seedling Stages in Turnips. PLoS ONE, 2015, 10, e0137983.	2.5	21
10	Genome-wide identification and analysis of the growth-regulating factor family in Chinese cabbage (<i>Brassica rapa</i> L. ssp. <i>pekinensis</i>). BMC Genomics, 2014, 15, 807.	2.8	80
11	MicroRNA expression analysis of rosette and folding leaves in Chinese cabbage using high-throughput Solexa sequencing. Gene, 2013, 532, 222-229.	2.2	20