Fengde Wang

List of Publications by Year in descending order

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1163117 1058476 14 396 8 14 citations h-index g-index papers 14 14 14 594 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Transcriptome analysis of the roots at early and late seedling stages using Illumina paired-end sequencing and development of EST-SSR markers in radish. Plant Cell Reports, 2012, 31, 1437-1447.	5.6	96
2	Genome-wide identification and analysis of the growth-regulating factor family in Chinese cabbage (Brassica rapa L. ssp. pekinensis). BMC Genomics, 2014, 15, 807.	2.8	80
3	Transcriptome analysis of rosette and folding leaves in Chinese cabbage using high-throughput RNA sequencing. Genomics, 2012, 99, 299-307.	2.9	48
4	Genome-Wide Identification and Analysis of the VQ Motif-Containing Protein Family in Chinese Cabbage (Brassica rapa L. ssp. Pekinensis). International Journal of Molecular Sciences, 2015, 16, 28683-28704.	4.1	43
5	Physiological and Transcriptomic Responses of Chinese Cabbage (Brassica rapa L. ssp. Pekinensis) to Salt Stress. International Journal of Molecular Sciences, 2017, 18, 1953.	4.1	28
6	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
7	MicroRNA expression analysis of rosette and folding leaves in Chinese cabbage using high-throughput Solexa sequencing. Gene, 2013, 532, 222-229.	2.2	20
8	Characterization and Development of EST-SSRs by Deep Transcriptome Sequencing in Chinese Cabbage (<i>Brassica rapa</i> L. ssp. <i>pekinensis</i>). International Journal of Genomics, 2015, 2015, 1-11.	1.6	20
9	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
10	Genome-Wide Identification and Analysis of the Cytochrome B5 Protein Family in Chinese Cabbage (<i>Brassica rapa</i> L. ssp. <i>Pekinensis</i>). International Journal of Genomics, 2019, 2019, 1-16.	1.6	8
11	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
12	Ectopic expression of a Brassica rapa AINTEGUMENTA gene (BrANT-1) increases organ size and stomatal density in Arabidopsis. Scientific Reports, 2018, 8, 10528.	3.3	7
13	Transcriptome Analysis of Orange Head Chinese Cabbage (Brassica rapal. ssp.pekinensis) and Molecular Marker Development. International Journal of Genomics, 2017, 2017, 1-8.	1.6	5
14	Construction of an Intragenic SSR-Based Linkage Map and QTL Mapping for Agronomic Traits in Chinese Cabbage (Brassica rapa L. ssp. pekinensis). Horticulturae, 2022, 8, 165.	2.8	5