

Liping Chen

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Regulation of Leaf Longevity by DML3-Mediated DNA Demethylation. <i>Molecular Plant</i> , 2020, 13, 1149-1161.	8.3	29
2	Heritability and Reversibility of DNA Methylation Induced by in vitro Grafting between <i>Brassica juncea</i> and <i>B. oleracea</i> . <i>Scientific Reports</i> , 2016, 6, 27233.	3.3	28
3	Heritable variation and small RNAs in the progeny of chimeras of <i>Brassica juncea</i> and <i>Brassica oleracea</i> . <i>Journal of Experimental Botany</i> , 2013, 64, 4851-4862.	4.8	27
4	Differential DNA methylation and gene expression in reciprocal hybrids between <i>Solanum lycopersicum</i> and <i>S. pimpinellifolium</i> . <i>DNA Research</i> , 2017, 24, 597-607.	3.4	24
5	Maintenance of grafting-induced epigenetic variations in the asexual progeny of <i>Brassica oleracea</i> and <i>B. juncea</i> chimera. <i>Plant Journal</i> , 2018, 96, 22-38.	5.7	20
6	Light- and temperature-regulated BjAPY2 may have a role in stem expansion of <i>Brassica juncea</i> . <i>Functional and Integrative Genomics</i> , 2015, 15, 753-762.	3.5	11
7	Production of <i>Brassica</i> tri-genomic vegetable germplasm by hybridisation between tuber mustard (<i>Brassica juncea</i>) and red cabbage (<i>B. oleracea</i>). <i>Euphytica</i> , 2015, 204, 323-333.	1.2	10
8	In vitro propagation of <i>Lychnis senno</i> Siebold et Zucc., a rare plant with potential ornamental value. <i>Scientia Horticulturae</i> , 2006, 107, 183-186.	3.6	8
9	Phenotypic and genetic variation occurred during wide hybridisation and allopolyploidisation between <i>Brassica rapa</i> and <i>Brassica nigra</i> . <i>Scientia Horticulturae</i> , 2014, 176, 22-31.	3.6	8
10	Linkage map construction using SSR markers and QTL analyses of stem expansion traits in <i>Brassica juncea</i> . <i>Scientia Horticulturae</i> , 2016, 209, 67-72.	3.6	6
11	The high-throughput sequencing of small RNAs profiling in wide hybridisation and allopolyploidisation between <i>Brassica rapa</i> and <i>Brassica nigra</i> . <i>Genomics Data</i> , 2015, 3, 1-3.	1.3	2