Jing Wang

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A Tourist-like MITE insertion in the upstream region of the BnFLC.A10 gene is associated with vernalization requirement in rapeseed (Brassica napus L.). BMC Plant Biology, 2012, 12, 238.	3.6	94
2	The evolution of Brassica napus FLOWERING LOCUST paralogues in the context of inverted chromosomal duplication blocks. BMC Evolutionary Biology, 2009, 9, 271.	3.2	86
3	Comparative Analysis of FLC Homologues in Brassicaceae Provides Insight into Their Role in the Evolution of Oilseed Rape. PLoS ONE, 2012, 7, e45751.	2.5	79
4	A <scp>CACTA</scp> â€like transposable element in the upstream region of <i>BnaA9</i> . <i>scp>CYP78A9</i> acts as an enhancer to increase silique length and seed weight in rapeseed. Plant Journal, 2019, 98, 524-539.	5.7	77
5	Promoter Variation and Transcript Divergence in Brassicaceae Lineages of FLOWERING LOCUS T. PLoS ONE, 2012, 7, e47127.	2.5	37
6	Transposon insertions within alleles of BnaFLC.A10 and BnaFLC.A2 are associated with seasonal crop type in rapeseed. Journal of Experimental Botany, 2020, 71, 4729-4741.	4.8	32
7	Sequence variation and functional analysis of a FRIGIDA orthologue (BnaA3.FRI) in Brassica napus. BMC Plant Biology, 2018, 18, 32.	3.6	24
8	Universal endogenous gene controls for bisulphite conversion in analysis of plant DNA methylation. Plant Methods, 2011, 7, 39.	4.3	15
9	Widespread and evolutionary analysis of a MITE family Monkey King in Brassicaceae. BMC Plant Biology, 2015, 15, 149.	3.6	9
10	Functional homoeologous alleles of CONSTANS contribute to seasonal crop type in rapeseed. Theoretical and Applied Genetics, 2021, 134, 3287-3303.	3.6	6