Jing Wang

List of Publications by Year in descending order

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		1163117	1372567	
10	459	8	10	
papers	citations	h-index	g-index	
10 all docs	10 docs citations	10 times ranked	465 citing authors	

#	Article	IF	CITATIONS
1	Functional homoeologous alleles of CONSTANS contribute to seasonal crop type in rapeseed. Theoretical and Applied Genetics, 2021, 134, 3287-3303.	3.6	6
2	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
3	A <scp>CACTA</scp> â€like transposable element in the upstream region of <i>BnaA9</i> . <i><cp>CYP78A9</cp></i> acts as an enhancer to increase silique length and seed weight in rapeseed. Plant Journal, 2019, 98, 524-539.	5.7	77
4	Sequence variation and functional analysis of a FRIGIDA orthologue (BnaA3.FRI) in Brassica napus. BMC Plant Biology, 2018, 18, 32.	3.6	24
5	Widespread and evolutionary analysis of a MITE family Monkey King in Brassicaceae. BMC Plant Biology, 2015, 15, 149.	3.6	9
6	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
7	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
8	Promoter Variation and Transcript Divergence in Brassicaceae Lineages of FLOWERING LOCUS T. PLoS ONE, 2012, 7, e47127.	2.5	37
9	Universal endogenous gene controls for bisulphite conversion in analysis of plant DNA methylation. Plant Methods, 2011, 7, 39.	4.3	15
10	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