

Supplementary data**Identification and abiotic stress analysis of calmodulin-binding transcription activator/signal responsive genes in non-heading Chinese cabbage (*Brassica campestris* ssp. *chinensis* Makino)**

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Table1. Sequences of primers for BcCAMTAs

Primer name	Primer sequences (5' to 3')
BcCAMTA1-F	AGTCACCGGAAGAACCTCAA
BcCAMTA1-R	TCAAGGAGAAGTGGACATCAA
BcCAMTA2.1-F	ATGGCGGATCGTGGAA
BcCAMTA2.1-R	TCATTCAAATGCAAGAGACA
BcCAMTA2.2-F	ATGGCGCAGATTAGCAA
BcCAMTA2.2-R	TCATTCAAATGCAAGAGACA
BcCAMTA3.1-F	ATGGCGGAAGCAAGACG
BcCAMTA3.1-R	TCAAGCGTCCACAAAGTT
BcCAMTA3.2-F	ATGGCGGAAGCAAGGAGA
BcCAMTA3.2-R	TCAATTGAAACCATTAGCGTAGG
BcCAMTA4-F	ATGCAGTCTGGTGAGTTCT
BcCAMTA4-R	CTACTTCTCCTTGCAGA
BcCAMTA5-F	ATGGCCGGCGTCGATT
BcCAMTA5-R	TCAGCTCTCAATATCCATCTGATTA
BcCAMTA6-F	ATGGATGGTGTGGTTCTC
BcCAMTA6-R	TTACAGATCAAGAGCATCAATAC
qBcCAMTA1-F	AATCGTCGTCGTCGTTAT
qBcCAMTA1-R	TGGTTGCTTATCGGAATCT
qBcCAMTA2.1-F	GTAAGCTATTGACTGTTG
qBcCAMTA2.1-R	TATTGCTATTGTTGTTGTTG
qBcCAMTA2.2-F	TGGAGGTTAAGGTATAGC
qBcCAMTA2.2-R	AATCAGCATCTTCACATAG
qBcCAMTA3.1-F	GTТАТGGATGCTTCAGGAA
qBcCAMTA3.1-R	CGTTCACTGGTGTAGACT
qBcCAMTA3.2-F	CTCATTCCAGTTCAACAG
qBcCAMTA4-F	TACGACCATATTGTTCTTGTTC
qBcCAMTA4-R	GATGAATGCTGCTGAAGTAA
qBcCAMTA5-F	TAACTACTCACCATCAAC
qBcCAMTA5-R	CCATCCTCTCTTCTTC
qBcCAMTA6-F	GAGGTGTTATTGGTTACTA
qBcCAMTA6-R	TTGATGATGAATGAGAGTT

Fig. S1 Multiple sequence alignment of CAMTA proteins in NHCC and Arabidopsis thaliana.

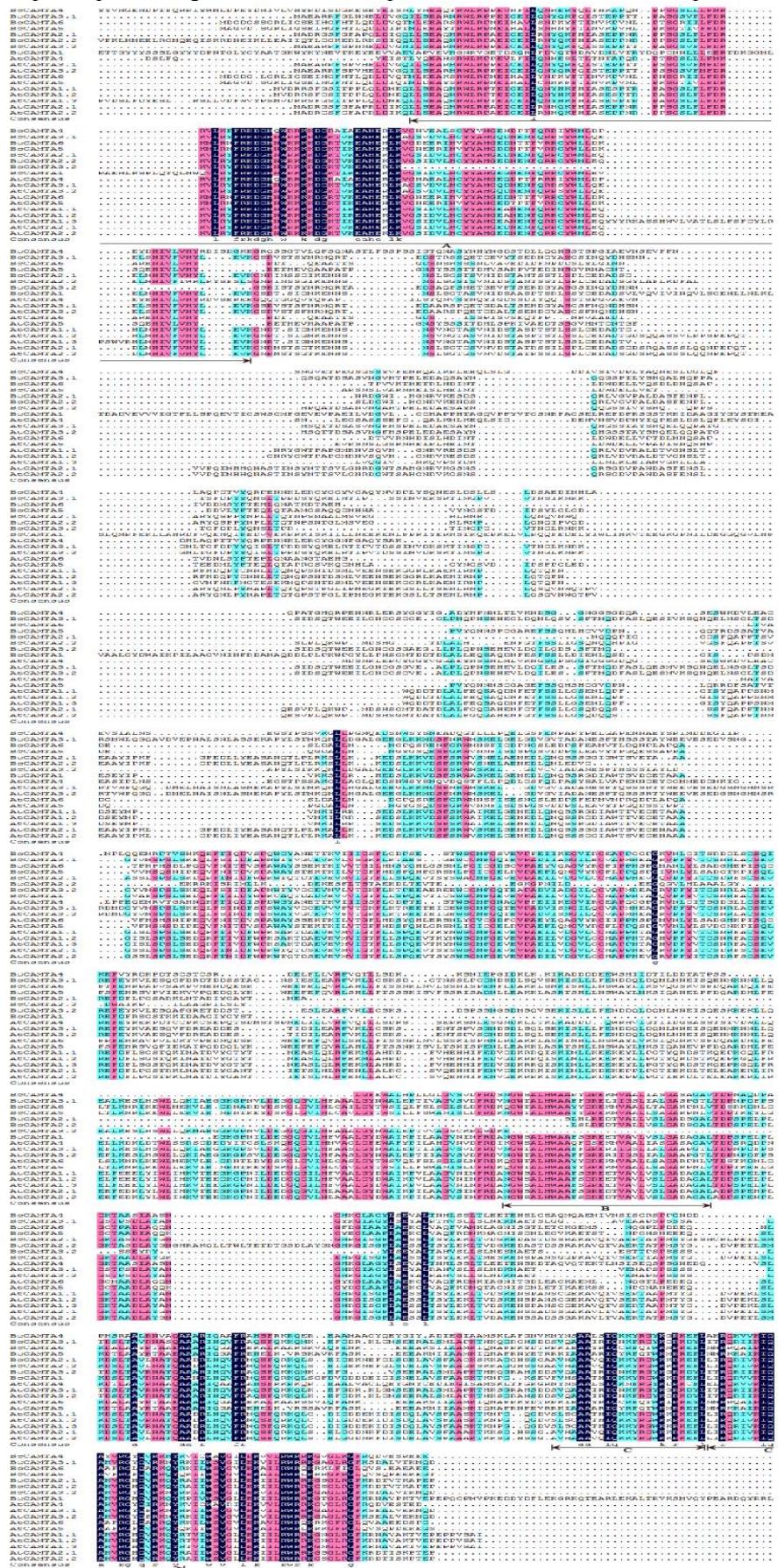


Fig 3.

