

Table 1 Suppl. Primers used in this study. RT-qPCR - reverse transcription quantitative PCR.

	Primer name	Primer 5'-3'	Size [bp]
Subcellular localization	CsCAMTA1-F	CC ATCGAT ATGGCTCAGACTAGACGATATG	3054
	CsCAMTA1-R	TCC CCCGGG TCATTGATCAGCTACATATGCG	
	CsCAMTA2-F	GCG TCTAGA ATGGCGGATCGAGGGTCGTA	3252
	CsCAMTA2-R	TCC CCCGGG TCATTCAAATGCCACAGACATG	
	CsCAMTA3-F	GAT GGATCC ATGGCAGACTCTAGACGCTTTG	3279
	CsCAMTA3-R	TCC CCCGGG TCAAGAAGCATTAGGCATGAGA	
	CsCAMTA4-F	GCG TCTAGA ATGATGCTTGCAAGGATATGATG	2541
	CsCAMTA4-R	TCC CCCGGG CTACACAAATCGGTATGTCGAT	
RT-qPCR	CsCAMTA5-F	GAT GGATCC ATGGCTGAGCTGCTTGTCGG	2754
	CsCAMTA5-R	TCC CCCGGG TCAATCAGCCATTTCCATATCT	
	CsCAMTA1-2F	ATAAAAAGGTTGTTTGGTTCGGT	152
	CsCAMTA1-2R	TCGCAAGAACTCATATTCGTCT	
	CsCAMTA2-2F	TAGAGGCAAACCCTGAGAAGAA	100
	CsCAMTA2-2R	TTACGAAGAAGCCATGAGTACA	
	CsCAMTA3-2F	ATAATCCGGAAGCAAATCATCA	154
	CsCAMTA3-2R	CCGACTTAAATCCACGCAAACC	
	CsCAMTA4-2F	AGGAGTTGGTTTGCGAGGTT	172
	CsCAMTA4-2R	CCGGCGATATTGGTTACGTG	
	CsCAMTA5-2F	AAAAGATACTTTGTCAGCTTACCG	118
	CsCAMTA5-2R	CTCTGGACTAGAAAACCTAATTGC	
	CsActin-F	CATCCCTCAGCACCTTCC	176
	CsActin-R	CCAACCTTAGCACTTCTCC	
Transcription activation	CsCAMTA1-3F	GC CCCGGG ATGGCTCAGACTAGACGATATG	3054
	CsCAMTA1-3R	GCGCTGCAG TCATTGATCAGCTACATATGCG	
	CsCAMTA2-3F	GC CCCGGG ATGGCGGATCGAGGGTCGTA	3252
	CsCAMTA2-3R	GCGCTGCAG TCATTCAAATGCCACAGACATG	
	CsCAMTA3-3F	GC CCCGGG ATGGCAGACTCTAGACGCTTTG	3279
	CsCAMTA3-3R	GCGCTGCAG TCAAGAAGCATTAGGCATGAGA	
	CsCAMTA4-3F	GC CCCGGG ATGATGCTTGCAAGGATATGATG	2541
	CsCAMTA4-3R	GCGCTGCAG CTACACAAATCGGTATGTCGAT	
	CsCAMTA5-3F	GC CCCGGG ATGGCTGAGCTGCTTGTCGG	2754
	CsCAMTA5-3R	GCGCTGCAG TCAATCAGCCATTTCCATATCT	

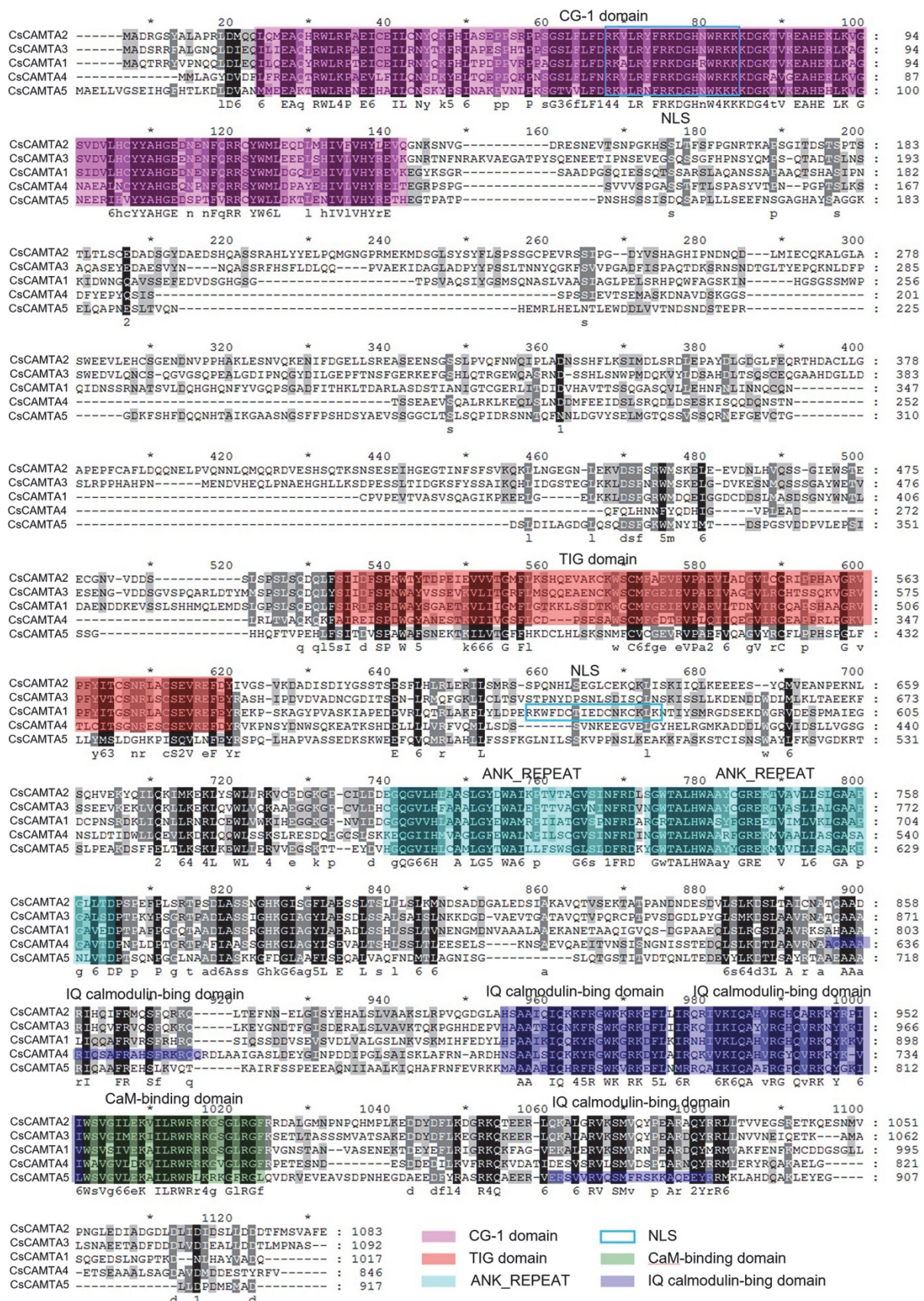


Fig. 1 Suppl. Multiple sequence alignments of the 5 *Citrus* CAMTA proteins obtained by *ClustalX* and viewed by *GeneDoc*. Protein sequences for the CsCAMTA members are as follow: CsCAMTA1 (orange1.lg001759), CsCAMTA2 (orange1.lg001406), CsCAMTA3 (orange1.lg001365), CsCAMTA4 (orange1.lg044976), CsCAMTA5 (orange1.lg001759). Conserved domains analyses of CsCAMTA proteins were predicted using the *Pfam* and *Motif Scan* programs.