

Table 1 Suppl. The sequences of the primers used in these experiments. The nucleotide sequence analysis was performed using *BLASTN* and *BLASTX* programs of *NCBI* (<http://www.ncbi.nlm.nih.gov/BLAST>). Chromosomal location of *VvZFP11* gene was predicted by *BLAT* server at the *Genoscope Genome Browser* (<http://www.genoscope.cns.fr/blat-server/cgi-bin/vitis/webBlat>). The protein conserved domain was analyzed using *Smart* (http://smart.emblheidelberg.de/smart/change_mode.pl) and *ExPASy* (<http://au.expasy.org/tools/>). The amino acid sequences of the ZFP proteins were aligned using *DNAMAN*.

Primer name	Oligonucleotide primers	Purpose or vector
VvZFP11-BamHI-F	5'-AGCAGGGATCCTATAATATGAAGAGAAGCAGAGTTGATGGA-3'	pCAMBIA3301
VvZFP11-SacI-R	5'-ATGCTGAGCTCTTACAAAATACAATCCACCATATGAGGCAC-3'	
VvZFP11-F	GTCACAGAGCCAGCCACAAGAA	qRT-PCR
VvZFP11-R	TTGCCCAATCGCGAACTCCAAC	
VvAct-F	TCCTGTGGACAATGGATGGA	qRT-PCR
VvAct-R	CTTGCATCCCTCAGCACCTT	
AtPR1-F	GGAGCTACGCAGAACAATAAGA	qRT-PCR
AtPR1-R	CCCACGAGGATCATAGTTGCAACTGA	
AtPDF1.2-F	TCATGGCTAAGTTTGCTTCC	qRT-PCR
AtPDF1.2-R	AATACACACGATTTAGCACC	
β-TUB 4-F	GAGGGAGCCATTGACAACATCTT	qRT-PCR
β-TUB 4-R	GCGAACAGTTCACAGCTATGTTCA	
VvZFP11-BglII-F	GGGAGATCTAATATGAAGAGAAGCAGAGTTGATGGAG	pGBKT7
VvZFP11-PstI-R	GGGCTGCAGCAAAAATACAATCCACCATATGAGGCACC	
VvZFP11ΔA-BglII-F	GGGAGATCTATGAAGAGAAGCAGAGTTGATGGAG	pGBKT7
VvZFP11ΔA-PstI-R	GGGCTGCAGATCCAAGGACAAAACCTCTCTCGTC	
VvZFP11ΔB-BglII-F	GGGAGATCTATGAAGAGAAGCAGAGTTGATGGAG	pGBKT7
VvZFP11ΔB-PstI-R	GGGCTGCAGGGGCTTCTTGTGGCTGGCTCTGTGA	
VvZFP11ΔC-BglII-F	GGGAGATCTATGAAGAGAAGCAGAGTTGATGGAG	pGBKT7
VvZFP11ΔC-PstI-R	GGGCTGCAGCCTAGACAGGAGCATCAAGCAGTT	
VvZFP11ΔD-BglII-F	GGGAGATCTGAAAATCCGAATCCACCAACCAGCT	pGBKT7
VvZFP11ΔD-PstI-R	GGGCTGCAGCAAAAATACAATCCACCATATGAGGCACC	
VvZFP11ΔE-BglII-F	GGGAGATCTCCTACGCCGGTGTGCGTTGTGAAGAA	pGBKT7
VvZFP11ΔE-PstI-R	GGGCTGCAGCAAAAATACAATCCACCATATGAGGCACC	
VvZFP11-XbaI-F	5'-AGCATCTAGATATAATATGAAGAGAAGCAGAGTTGATGGA-3'	pBI221-GFP
VvZFP11-KpnI-R	5'-ATGCTGGTACCCAAAATACAATCCACCATATGAGGCAC-3'	