

Supplementary table S1. Oligonucleotides used as primers for real-time RT-PCR.

Gene	Accession no.	Primer orientation	Primer ^a	Primer amount ^b	Amplicon size	% identity at the amino acid level with the best score (accession no.)
Pheophorbide a oxygenase	DY261492 ^c	F	5'-GCGACAAATGACGGTAAAAAGC-3'	3.6 µl	103 bp	81% <i>Solanum lycopersicum</i> (AF321984)
		R	5'-CGTTTCTTCATCTGATGCTGCA-3'	3.6 µl		
RCC reductase	CX306368 ^e	F	5'-CTCCAGCTCTCTCATCCATCG-3'	1.2µl	104 bp	64% <i>Arabidopsis thaliana</i> (NM119863)
		R	5'-GTGCGGCAGAACAATTGACTTT -3'	1.2µl		
Geranylgeranyl reductase	DY275174 ^e	F	5'-GCGTGATTAACGGGTTAT-3'	3.6 µl	353 bp	75% <i>Medicago truncatula</i> (AY960125)
		R	5'-CACTGTGCCTGTACCA-3'	3.6 µl		
Phytoene synthase	AF220218 ^d	F	5'-GGTCGTCCATTTGATATGCTTG-3'	1.2µl	111 bp	100% <i>Citrus unshiu PSY</i>
		R	5'-CCTAAGGTCCATCCTCATTCCCT-3'	0.2µl		
Stay green	CX308230 ^e	F	5'-CATCTTCTCCAAGGAAGCTCCCC-3	1.2µl	101 bp	86% <i>Solanum lycopersicum</i> (DQ100158)
		R	5'-CCCAAACCAAAGCCTCCTGTA-3	1.2µl		
Secretory peroxidase	CX308252 ^e	F	5'-GTTCTTGAGAGGTTTGCAGCCA-3	1.2µl	101 bp	89% <i>Nicotiana tabacum</i> (AF149251)
		R	5'-TGCACCAGCTTCACACAATGA-3	1.2µl		
Cysteine proteinase	CX305503 ^e	F	5'-CGAGAGAACTCCAAGGACCAGA-3	1.2µl	113 bp	100% <i>Citrus sinensis</i> (Z47793)
		R	5'-GCACATGCGTTCTCTTGCAA-3	1.2µl		
Early light induced protein	CX289917 ^e	F	5'-CATACCGAAAGCTGAGCCTCA -3'	1.2µl	113 bp	80% <i>Brassica rapa</i> (AY433944)
		R	5'-TCGCAAGCCTCCCGTTTAT-3'	1.2µl		
Chlorophyllase	AF160869 ^e	F	5'-TCATCGTTGTTGCTCCTCAGC-3'	0.5µl	69 bp	100% <i>Citrus sinensis CLH</i>
		R	5'-TGCGCTGTTAAGTTCATTGG-3'	0.5µl		

^aOligonucleotides were designed using the Primer express software (Applied Biosystems). ^bOptimized amounts of a 5µM oligonucleotide solution. ^cCitrus ESTs sequences. ^dCitrus full length cDNA sequence (Kim et al., 2001). ^eCitrus full length cDNA sequence (Jacob-Wilk et al. 1999).