

Additional File S1.- Table. Designed primers for gene expression analyses by quantitative RT-PCR.

Citrus gene	Locus	Forward / Reverse	Sequence (5'→3')	Amplicon size (bp)
CsNCED1	orange1.1g007379m	F R	AATGCTTGGGAAGAGCCTGAG AGTGGACTCGCCGGTCTTTAG	147
CsCYP707A1	orange1.1g038621m	F R	TCAATGTTGCACTGCTCTCC CTTTGGCACCCATGAAAGAT	246
CsAOG	orange1.1g022744m	F R	CGGGTTCAGTGTGGTCTT GCCTCGAGAGAAATGGATGT	176
CsBG18	orange1.1g010588m	F R	CAAGGCAAAACAGGGTGGAT CAGCCTCAGAGCTGGTGAAT	212
CsRAB18	orange1.1g028210m	F R	CTGAAGCTGAACGGGAGATT TTGTGGTGGTAGAGGTGGTG	195
CsACT	orange1.1g037845m	F R	CCCTTCCTCATGCCATTCTTC CGGCTGTGGTGGTAAACATG	105
CsTUB	orange1.1g013335m	F R	GGGGCAAATGAGCACTAAA CGCCTGAACATCTCCTGAAT	187