

Table S2: Accession numbers of the transcripts quantified by qRT-PCR and their primer sequences.

Target Gene	GenBank (GB) ID	GenModel (ITAG2.3)	Primer Sequence
<i>ACT</i>	XM_004235020.1	Solyc03g078400.2	QF 5'-TTAGCACCTTCCAGCAGATGT-3' QR 5'-AACAGACAGGACACTCGCACT-3'
<i>PPO-F</i>	AK247126.1	Solyc08g074630.1.1	QF 5'-CGGAGTTTGCAGGGAGTTATAC-3' QR 5'-TTGATCTCCACACTTTCAATGG-3'
<i>TD-II</i>	M61915.1	Solyc09g008670.2	QF 5'-TGCCGTTAAAAATGTCACCA-3' QR 5'-ACTGGCGATGCCAAAATATC-3'
<i>JIP-21</i>	AJ295638.1	Solyc03g098790.1	QF 5'-ACTCGTCCTGTGCTTTGTCC-3' QR 5'-CCCAAGAGGATTTTCGTTGA-3'
<i>WIPI-II</i>	AY129402.1	Solyc03g020080.2	QF 5'-GACAAGGTACTAGTAATCAATTATCC-3' QR 5'-GGGCATATCCCGAACCCAAGA-3'
<i>PR-P6</i>	M69248.1	Solyc00g174340.1	QF 5'-GTACTGCATCTTCTTGTTC-3' QR 5'-TAGATAAGTGCTTGATGTCCA-3'