

Table S1. Information of candidate reference genes, target genes and primers for RT-qPCR.

Gene name ^a	Description	Primer Sequence (5'-3') (For./Rev.)	Amplicon length (bp)/Tm (°C)	PCR efficiency (E)	Correlation coefficient (R ²)	GenBank Accession ^b
<i>TBP1</i>	TATA binding protein 1	CGGAGTACAATCCCAAGCGT/ AGCATACTTCCTTGCTGCCAA	146/77.5	93.30	0.997	MZ210062
<i>CBP20</i>	Nuclear cap binding protein 20	AGCTATGCCACCGTCCTATG/ ACCACTCTCCCCGAAATCGC	208/78.5	118.60	0.982	MZ210063, MZ210064
<i>TIP41</i>	TIP41-like family protein	TCTTCTCTTGCATTCTGGCTT/ GGTCCTCTGTATGACGATGGG	229/79.5	86.90	0.999	MZ210065, MZ210066
<i>EF-1α</i>	Translation elongation factor EF1A	ACAAGATGGACGCCACTACC/ AGAGGCTTGTCTGAGGGTCT	244/80.0	94.20	0.997	MZ210067, MZ210068
<i>EIF4A1</i>	Eukaryotic translation initiation factor 4A1	GCAAGGTGGACTGGCTTACT/ TTTCAGGTTGGTTGGCAGA	216/79.5	92.90	0.999	MZ210069
<i>PTB1</i>	Polypyrimidine tract-binding protein 1	GTCAAACCTATGCGGCTCCT/ AAGGCGAAGACCGAACCAAT	240/81.5	92.50	0.998	MZ210070
<i>ACT7</i>	Actin 7	TTGCTGGCCGTGATCTAACT/ ACTTCTGGGCAACGGAATCT	241/79.0	94.80	1.000	MZ210071
<i>ACT1</i>	Actin 1	ACCCAAAGGCTAACCCTGAA/ CGTACGGCCACTAGCATACA	113/76.5	87.90	0.998	MZ210072
<i>GAPA</i>	Glyceraldehyde-3-phosphate dehydrogenase A subunit	GCAGAAGAAGTGAACGCTGC/ GCTGTAACCCCACTCGTTGT	201/80.5	106.50	0.990	MZ210073
<i>GAPB</i>	Glyceraldehyde-3-phosphate dehydrogenase B subunit	TGTTCCACTCGTGTCTGTGG/ CTGTAGCCCCACTCGTTGTC	129/78.0	103.80	0.986	MZ210074
<i>TUB1</i>	Tubulin beta-1 chain	CCAAGGAGGTTGACGAGCAA/ GCTCACCCCTTCGGAACATCT	179/77.5	90.40	0.995	MZ210076
<i>eIF2</i>	Eukaryotic translation initiation factor 2	TCCTTACACTCGAACCACGC/ ACACGGACCCATTCAACTCC	149/81.0	101.90	0.998	OL456245
<i>ACO1</i>	1-aminocyclopropane-1-carboxylate oxidase 1	CTTGGGGACCAACTTGAGGTTATT/ ATCGTTGCCTGGGTTGTAGAAG	120/77.0	95.80	0.999	OL456246
<i>PYL1</i>	PYR1-like 1	CAGTATAATTGGGGGCGAGCA/ ACGATTCCAGAACCACGCTC	104/81.0	89.60	0.988	OL456247
<i>CSD2</i>	Copper/zinc superoxide dismutase 2	TGTTGGTGATGACGGAAGTGT/ ATGCCACCTTCCCAAGAT	130/77.5	97.70	0.990	OL456248, OL456249

^a Gene name based on similarity to *Arabidopsis* proteins. ^b Full-length cDNA accession number.