

Table S9. Primers used for qRT-PCR assays in this study.

Gene ID	Gene name	Primer sequence(5'-3')	T _m (°C)
MD05G1283800	<i>GA2OX1</i>	F : ATAGCACCTCTGCCGTCAC TAA	57.3
		R : AATCAGCCAGCCTTGACCTGTA	57.7
MD09G1264800	<i>GAI</i>	F : AATTCCGGCGGGATTGAAGTCT	58.4
		R : ATGAGTGTGTGGACGAGTTGGA	57.7
MD11G1215600	<i>GAMYB1</i>	F : CTGGACGAACGGACAACGAGAT	58.7
		R : TGGGAATGGGAATGGGAATGGG	58.3
MD15G1060800	<i>PYL2</i>	F : GCACGTCGCTAATAACGCAGAA	58.1
		R : CGCCTTTCATGTTGCAGCTCTT	58.4
MD13G1017000	<i>AHK4</i>	F : TCGCAGGAAACCGTCTCTTACA	57.8
		R : ACCCAGCAGCCTAAATGGACTT	57.9
MD04G1212100	<i>AHP1</i>	F : ATGGATGGCGTTTCTCAGTTGC	57.9
		R : TGCAGTTGCTGAAGTTGGTTGA	57.3
MD16G1017900	<i>ARR2</i>	F : TCCACAACCCGTCATTTCTGCT	58.5
		R : GCACAACGCCCATCAAGAAGTT	58.3
MD06G1182400	<i>ARR6</i>	F : TCTTTCGGTGATTTCGCAGATG	57.1
		R : AGCCGTCACTTTGCAGGATGAT	58.6
MD16G1041700	<i>IPT1</i>	F : TCGACATCACCACCAACAAGCT	58.5
		R : CCGATTCGTAATGCCGGAAACA	57.7
MD15G1184800	<i>YUC2</i>	F : TATGGGAAGACGTGCTGAAACG	57.1
		R : TTTGTGGACCGTAACGGAAAGC	57.8
MD05G1309400	<i>ARF3</i>	F : TGAAGATGCAGCAGAGCGAAGA	58.4
		R : CTGCCATGCTTGCTTGTGTCTA	57.6
MD13G1164200	<i>SUS2</i>	F : TTGAGTGGTCAGTTCCGATGGA	57.4
		R : AGAAAGCAGGCTGCACAAAGAC	58.0
MD03G1095300	<i>ACTIN</i>	F : ACCACTCGTCTGTGACAATGGA	57.7
		R : ACCCATTTCCAACCATCACACCA	58.1
MD01G1035300	<i>HISTONE</i>	F : TGATCTGCGTTTCCAGAGCCAT	58.4
		R : ATGATGGTAACCCGCTTGGCAT	58.7
Mimida et al (2015)	<i>SAND</i>	F : CCCAGGACTTTGAGCTTTATGC	-
		R : TATCACCATGAAAAGGGGCTTG	-
Perini et al (2014)	<i>WL40</i>	F : GGATTTACTGTGTTGGTGAAG	-
		R : TGCCAATTACCTCCTTTTCGTG	-