

Table S9. Quantitative RT-PCR for RNA sequencing validation.

ID	Gene	Primer
PGSC0003DMT400014674	Elongation factor 1 alpha-like protein <i>EF1aL</i>	CTTGTACACCACGCTAAGGAG GTCAATGCAAACCATTCCTTG
PGSC0003DMT400033236	4,4-dimethyl-9beta,19-cyclopropylsterol-4alpha-methyl oxidase <i>SMO1</i>	GAGAGAACTGAGAATGGAGG GTGCTTGATAGAGAAACACAAC
PGSC0003DMT400020648	Cyclin D3 <i>CYCD3</i>	TGATAGGCTTAAACCTTGC TAACAGATGTAGCCACTGAC
PGSC0003DMT400064270	Peroxidase <i>POD</i>	CACTGTCCCTGCTGTTC TCCTGCCAATGATAGATTA
PGSC0003DMT400023554	Wax-ester synthase / diacylglycerol O-acyltransferase <i>WES</i>	GTGTTGACCCGATTACCCA CTTGTTTAGTGACAGCCTCTT
PGSC0003DMT400028227	Myb proto-oncogene transcription factor MYB48-like <i>MYB48</i>	GACGACACTTTGCACCAAC TCCAAGACAAGCCTACCTAAT
PGSC0003DMT400014568	General transcription factor IIE subunit 1-like <i>TFIIE</i>	AGGAAGGAGGGAGCAAA AGGCAGCAATAAGAATGAG
PGSC0003DMT400004046	Snakin2 <i>SN2</i>	CTCCTTGCTCCTTCTCG TAGCAAGGGCAAGTCTCAG
PGSC0003DMT400019919	Kip-related protein 4 <i>KRP4</i>	AAAAGACAGAATCGGAGTT GTCCTCTCTCTACCATCAAT
PGSC0003DMT400056352	WRKY transcription factor 75 <i>WRKY 75</i>	CAGCATCATCGTCGTCAT CATTAGTCCCAAGAACCC
PGSC0003DMT400015017	Transcription factor ATR2-like <i>ATR2</i>	CTCATAGTGCGTCTCTCCA AATCACACCTTCAGATCCAAT
PGSC0003DMG400001550	Pathogenesis-related protein STH-2 <i>STH-2</i> or TSI-1 protein	AAAACCAGGCATGGAAC ACGTGTAGACCTGATTCTTT
PGSC0003DMT400020263	Serine-rich protein <i>SRP</i>	CCAAACACGGCACAAAGA GGGACAGTGAACGGAGAA