

Table S6. Primer sequences used in Quantitative RT-PCR for iTRAQ validation.

ID	Gene	Primer
PGSC0003DMT400006894	Uncharacterized protein	TGGTGGACAGAGTGAAGAA CGAGTGTAGCATCGGAGTT
PGSC0003DMT400003773	Pathogenesis-related protein STH-2-like	CAACGACCACAATTTCCC TCCAACACCACCATCTCC
PGSC0003DMT400039734	Protochlorophyllide reductase-like	GGGATGCCTAAGGAGAAT CGAACCGACAATGATGAG
PGSC0003DMT400042741	4-alpha-glucanotransferase	TTCTTCTGTAGCCCGTAT AACTAGGTATCCTCCAACTC
PGSC0003DMT400015734	Heat shock protein 90-6	CAGTTTGTTTCATTCCCTAT ATCTTGCCCTTCCTTATT
PGSC0003DMT400037918	Glutathione s-transferase	GCCCATTTAGTCACAGAG CAAGGAACTTAGCCCAGA
PGSC0003DMT400005104	Glutathione s-transferase	ATGGAAGAATGGACCTCA CCAAATGCTACGACGATC
PGSC0003DMT400042214	Ubiquitin-conjugating enzyme E2	G TTCAGGAGGATCTAGTGTT CAGAA TTGTGAGGACCAA
PGSC0003DMT400090964	Protein transport protein Sec61	GTGGTAACGCTGTAGTAGGC GTAGTAACGAATCCGATGAAG
PGSC0003DMT400033585	14-3-3 protein	TTGAACAGAAGGAGGAAT GGTAGTAGCAGACGGAAC
PGSC0003DMT400018083	dnaJ protein homolog	TCAATACAAGGCGATAAATG GCTGCTCCTCGTGTCTCA
PGSC0003DMT400042739	Glycoside hydrolase	TTCTTCTGTAGCCCGTAT AACTAGGTATCCTCCAACTC