

Gene	Contig no.	Forward primer sequence (5'-3')	Reverse primer sequence (5'-3')	Primer efficiency [%]
<i>Cytochrome p450</i>	Contig_13761	CAGCGGTAGAAGAATGTGCC	GTTTCATCTGACGGGTTTGCA	100
<i>Galactinol synthase family protein</i>	Contig_6371	ATGGAGGTACTGTTGGGAGG	GCACCAACAGAAGCCAATGT	105
<i>Low-temperature-induced 65 kda</i>	Contig_13090	CGGTTACTGGGACTGAGT	TCATCACCAGGCCTCAACTT	101
<i>Nitrate transporter -like</i>	Contig_10659	CATGAAGGCGGGATCACTCT	CCTAAGGGTTGTGGGTGCAA	99
<i>Octicosapeptide phox bem1p family isoform 1</i>	Contig_5573	CTCTCAACAACAGCGTTGGC	CTGCCGGGTCGACCTTATTT	100
<i>Protein p21-like</i>	Contig_6745	CACATTTACGTGCCCTGGTG	AGCGTCTGCGCAACTAAT	97
<i>Serine-threonine protein plant</i>	Contig_14505	CTTTCCCAGCCGACATAAGC	GTCAGAAGATTCCACGGTGC	97
<i>Protein yls9-like</i>	Contig_2897	GGGAGTCAGATAAGTGGGTGG	TGTCCGGTCAAAGCCAAGTT	96
<i>UDP-glycosyltransferase74b1-like</i>	Contig_1957	AATGGGTGGTGGCTAGAGTG	TGCTCACTTATCCTGCCCAA	100
<i>Receptor-like protein 12</i>	Contig_21713	TTTGCGGATCTTGGACCTCT	CACCTTCTCCACGTTCGTC	94
<i>Probable lrr receptor-like serine threonine-protein kinase at4g36180-like</i>	Contig_11937	CTGTAACGCGCCAATGTCTG	TCTGGGACAATTCCGAGGGA	95
<i>CCT motif family protein isoform partial</i>	Contig_1104	CATGGTCAATCCCTGGCCTT	GACATGGGTAAGGACCACCA	95