

Table S1. Primers used in the analysis and alignment of all studied genes.

Gene	Accession number	Primer sequence		PCR product length
		F – forward primer		
		R – reverse primer		
Reference genes				
EF1 α	LOC102600998	F – ATTGGAAATGGATATGCTCCA		101
	LOC102600107	R – TCCTTACCTGAACGCCTGTCA		
Cyclophilin	LOC102586427	F – GCTCTTCGCCGATACCACTC		120
		R – ACACGGTGGAAGGTTGAGC		
Target genes				
<i>P5CS1</i>	LOC102582423	F – CTGCTCCTACATTTCTCTCCTTCG R – CGCTCCTAATCTTCCAAGTGCT		160
<i>PDH</i>	LOC102588726	F – ATAAGTGAATTTGTACACCA		117
	LOC102588395	R – GGCGAAAAGTGGAAGCG		
<i>PAL</i>	LOC102596343	F – TCGCTGAAGTGATGAACGGAA R – ATTGTGGAGATGTTCTGGAGA		207
	LOC102585357			
	LOC102585026			
	LOC102587888			
	LOC102596891			
<i>CHS1a</i>	LOC102577871	F – CTTGTCCCCGATAGCGAA R – AAATAGAGAGTTCCAGTCAGAT		162
<i>APX1</i>	LOC102586473	F – CTGGTGTCTGTTGCTGTTGAA		259
	LOC102586145	R – AAGATAAGAGGATTGGCGGTC		
<i>APX3</i>	LOC102597356	F – CCTTCTTTAGGGATTACGCCAT R – AGCCACAACAACACCTACAGCA		131
<i>SOS1</i>	LOC102602690	F – TTGCCTCACCATTCCATTG R – TGATGCTTCTGTGTTGTGGAAG		138
<i>NHX1</i>	LOC102578176	F – TGTTGATCCCTTTTCGACCAT R – CCAAATAGGGGTCGCATAAA		111
<i>NHX2</i>	LOC102605641	F – TGTTACTGGCATTCTGGTTTAC R – AATAGACAGAACAGTGACAGGA		124
<i>NHX3</i>	LOC102589621	F – GTCCATTACTGCCCTTGTTAT R – AAATGATAGGTGGAAGAAGGTAA		131