Table S1. Primer sequences for semi-quantitative RT-PCR.

UBQ – F	GATCTTTGCCGGAAAACAATTGGAGG
UBQ – R	CGACTTGTCATTAGAAAGAAAGAGAT
FLC – F	TTCTCCAAACGTCGCAACGGTCTC
FLC – R	GATTTGTCCAGCAGGTGACATCTC
FT – F	CTCAGGAACTTCTATACTTTGGTTATG
FT - R	CTGACAATTGTAGAAAACTGCG
TSF – F	CTTGATCCTTTCACGAGGTTGG
TSF – R	GGCAGTTGAAGTAAGAGGCA
FLM – F	CTCCGGTGACGACATTTCCAAGATCAT
FLM – R	TAAAAATCCAATCCGTACATTCAGACACAA
MAF2 – F	CCTTCAAAAAACGGTGGGGAAG
MAF2 – R	CAATTCTTTCTCATGAGCTTCTG
MAF3 – F	GAAAGGGAGAAGTTGCTGATAG
MAF3 – R	AAGAACTCTGATATTTGTCTACTAAGGTAC
MAF4 – F	CAATGCAAGATTGAAGAAGCGAAAAGC
MAF4 – R	GTGACATTGCTCTTGCATCCTCTG
MAF5 – F	TTCCACCGGCAAACTCTACAACTC
MAF5 – R	CCACCTCGCTAGCTAGAACCTT
SVP – F	ATCTCTTGAGTTACAGCTGGTTGAGAACA
SVP – R	ATGCACATTGTTACATATTTGCATGCCAA

Cycle parameters were as follows:
94°C for 2:00
  94°C for 0:30
  gene-specific annealing temp for 0:30 [0:20 for *MAF4*]
  72°C for gene-specific extension time
  (repeated until total cycle number achieved)
72°C for 8:00

Table S2. Gene-specific cycle parameters for semi-quantitative RT-PCR.

Gene:	**Annealing Temp:**	**Extension Time:**	**Cycle Number:**
UBQ	63°C	0:30	23x
FLC	63°C	0:30	32x
FT	63°C	0:30	36x
TSF	63°C	0:30	36x
FLM	55°C	0:50	42x
MAF2	61°C	0:20	41x
MAF3	60°C	0:40	37x
MAF4	64°C	0:17	45x
MAF5	60°C	0:30	42x
SVP	60°C	0:30	34x

Table S3. Primer sequences for quantitative RT-PCR.

UBQ – F	GGCCTTGTATAATCCCTGATGAATAAG
UBQ – R	AAAGAGATAACAGGAACGGAAACATAGT
FT – F	CTCAGGAACTTCTATACTTTGGTTATG
FT – R	GTTCCAGTTGTACGAGGGATATCAG
CO – F	CATTAACCATAACGCATACATTTCATC
CO – R	TCCGGCACAACACCAGTTT
GI – F	ACTAGCAGTGGTCGACGGTTTATC
GI – R	GCTGGTAGACGACACTTCAATAGATT

Cycle parameters were as follows:
50°C for 2:00
95°C for 15:00
  95°C for 0:15
  59-61°C for 1:00
  (repeated 40x)
Standard dissociation curve to check product specificity