

SUPPLIMENTARY TABLE S1. Primer sequences used for qRT-PCR analysis.

Gene	Accession No.	Forward sequence (5'-3')	Reverse sequence (5'-3')
CAB	AY288914	GGCAGCCCATGGTACGGATC	CCTCCITCGCTGAAGATCTGT
SAG12	XM_013821610.2	AGAGAATACCAAACCAAACCGAA	GCAACTCCCAAAATCTCAGGG
AAP1	AY188953.1	TGCTTACGCCACGGTTCTCA	GCTGCGCGAACACCTGATAG
AAP2	AY188954.1	CGGTACTGTTTGGACCGCAA	GACACGGGTCTTTTCCTCCG
AAP4	AY188955.1	CGCTGGACCTGCAGTGATGT	CCGCTCTCGTGAAAGCAGTTT
AAP6	AJ565848.1	GCGGCCGTAATGTCCTTTTC	ACCCACACACCCGCATAACA
<i>ACTIN</i>	AF111812	GATTCCGTTGCCCTGAAGTA	GCGACCACCTTGATCTTCAT