

Supplementary Table S1. Nucleotide sequences of primer pairs employed in this study.

Gene		Sequence (5' → 3')	Efficiency (%)	Source
<i>FcMYB1</i>	Fw	GGTTGAGTTGAATCTC	97.3	Synthesis / [11]
	Rv	GCAACTTGAGGATCAGCC		
<i>FcPAL2</i>	Fw	GTGAACCTCTTCCGATATGCTA	96.0	Synthesis
	Rv	CGATTAAAGCTAGGAATCCTCATCA		
<i>FcPAL4</i>	Fw	TGCAATTCCTATAGAGGGTCTTGA	100.7	Synthesis
	Rv	GGCACAAATGTGTCTTTACCCA		
<i>FcC4H</i>	Fw	CCTCCCCCTCATAAAACCCACCA	110.0	Synthesis
	Rv	CGATGATTGCGACTATAGCGA		
<i>Fc4CL</i>	Fw	ACTTGGTGAGGGATATGGGATG	108.0	[11]
	Rv	GCACCAGTTTCAGGTCTACG		
<i>FcCHS</i>	Fw	CGTCTCTCGACACTTCTCCG	100.8	Synthesis
	Rv	GGGTACGTGCTCTGGTCAAT		
<i>FcCHI</i>	Fw	TGGTAGGAGGTAGATGGCCGC	99.7	Synthesis
	Rv	GGTGGTAGTTGGGACTTATCGT		
<i>FcF3H</i>	Fw	GGGAGGGGTTTGGTATAGCTT	96.1	Synthesis
	Rv	GGTGTGGGGTGAAGGGGTGGTATAT		
<i>FcF3'H</i>	Fw	CAGAGAGTGGTGCGGAAAAG	97.9	Synthesis
	Rv	CCTTCAACACCATATCGACTCA		
<i>FcDFR1</i>	Fw	TAACTAAGGCATGGGGTTGG	95.9	Synthesis
	Rv	CCTGTGGTAACTCCAGCAGA		
<i>FcANS</i>	Fw	ACCCATAACTATAACTGTTTAAGGGCT	97.9	Synthesis
	Rv	TTGATCTTCGCAGCCAGTGT		
<i>FcUFGT</i>	Fw	ACCTAGGGTAGCAGCGTACA	94.7	Synthesis
	Rv	GTCATGCTCTTGCTCGGCAAC		
<i>FcFLS</i>	Fw	TTATCTTTGGGGTTAGGGCTTGAA	100.0	Synthesis
	Rv	GAGAATGGTGAGGGCGGACA		
<i>FcLAR</i>	Fw	GGTGATGGCACGGTTAAAGC	100.4	[11]
	Rv	CTCCCACAGTGAAGCAAGTCC		
<i>FcANR</i>	Fw	AACTAGAAACCGGTGGTCTTTGT	90.0	Synthesis
	Rv	GCTGAAGTTCTCAGCATGTTATGAT		
<i>Fc18S</i>	Fw	ATTTCGGTCCTATTCTGTTGGC	96.0	[30]
	Rv	GCTTTCGCAGTTGTTCTGCTTTT		
<i>FcGAPDH</i>	Fw	GATCTACCTTGCGCAAACCA	96.0	[31]
	Rv	GAGGCGGATCATGTGAACTT		