

Supplemental Table 1. Primer sets used in this study.

Primer name	Purpose	Sequence (5'-3')*	Annealing site in <i>JcCSH</i>
CSH_F1	cDNA cloning	ArTAGATGATACAwwTGATGCdTATG	1117-1136
CSH_R1	cDNA cloning	GAAACHCCATrTTsdTTCATrTAGCA	1613-1638 (complementary)
CSH_F2	cDNA cloning	CAyArTGGCrTrCmrAGrCTAGAGGC	818-843
CSH_R2	cDNA cloning	GGTACAGGATTTATGCAAATCCAG	1946-1969 (complementary)
CSH_F3	3'RACE	AGAGGACAykkTGCTTCwkGCATCG	1586-1610
CSH_R3	5'RACE	CAGTTTCATTTTCGACTTTCATCGACC	868-893 (complementary)
CSH_F4**	YFP fusion	<u>CACCATGGCAATGCAACCTGCAATTGTTCAAGC</u>	65-93 (translation start)
CSH_R4	YFP fusion	AGTGGCAATAGGTTCAATGAACAATCC	1841-1867 (lacking stop codon, complementary)
CSH_F5	qPCR	AGAGCAACAAATCGAAGTTGGTAA	1-24
CSH_R5	qPCR	TAATAATTCAAATGCCGGTTGATTG	307-331 (complementary)
18SrRNA_F	qPCR	AGAAACGGCTACCACATC	
18SrRNA_R	qPCR	CCAAGGTCCAACACTACGAG	

*: d, A/G/T; h, A/C/T; k, G/T; m, A/C; r, A/G; s, C/G; w, A/T; y, C/T.**: CACC was added for directional cloning in the pENTR™/D-TOPO Cloning System (Life Technologies).