Table S1. PCR primers for cDNA cloning.

Gene (product size, bp)	Primer (F, forward; R, reverse)
(1st PCR)	
CSF1 (943)	F: 5′-AAAGTGAAAGTTTGCCTCGGTGCTCTTGGC-3′ R: 5′-TAGAATTCCCTTTACACTGGCAATTCCACC-3′
CSF2 (513)	F: 5′-TCTTTTGCTAGTGAGCCCAGTGCTCAGAG-3′ R: 5′-TGGCTTCTGGCTGCTGCAGGCTGCTTTAC-3′
(2nd PCR)	
CSF1 (799)	F: 5′-ACGAATTCGCCACCATGACCGCGCCGGGCG-3′ R: 5′-TAGAATTCCCTTTACACTGGCAATTCCACC-3′
CSF2 (456)	F: 5′-AGAATTCGCCACCATGTGGCTGCAGAACCT-3′ R: 5′-TGAATTCTTACTTTTTGACTGGCCCCCAGC-3′

The *Eco*RI recognition sites are underlined.

Table S2. PCR primers for semiquantitative RT-PCR.

Gene (product size, bp)*	Accession no.	Primer (F, forward; R, reverse) *
CSF1 (323)	NM_001244523.1	F: 5′-AAAGTGAAAGTTTGCCTCGGTGCTCTTGGC-3′ R: 5′-TAGAATTCCCTTTACACTGGCAATTCCACC-3′
CSF1R (354)	XM_003124100.6	F: 5′-AAGATTGGGGACTTTGGTCTGGCAAG-3′ R: 5′-AAGGAGGGAGCAGATCTGCTGGAAGG-3′
CSF2 (343)	NM_214118.2	F: 5′-AGCATGTGGATGCCATCAAAGAAGCC-3′ R: 5′-ACTTTTTGACTGGCCCCCAGCAGTC-3′
CSF2R (379)	XM_001924779.5	F: 5′-AGTTTGAGGTGGTCTACAGGCGGCTTC-3′ R: 5′-TCCTCCTTCTGCACTGGGGAACATTC-3′
IL-4 (310)	NM_214123.1	F: 5′-ACCCTGGTCTGCTTACTGGCATGTAC-3′ R: 5′-TCTTCTTGGCTTCATGCACAGAACAGG-3′
IL-4R (382)	NM_214340	F: 5′-ATCTGCCTATCCGACTATGTCAGCAC-3′ R: 5′-AGGTGAGTTCCGAGTACAGGTAGCTC-3′
LIF (278)	NM_214402.2	F: 5′-TGTCACAGCAACCTCATGAACCAG -3′ R: 5′-TTCACAGCACCAGGATTGAGGCTC -3′
GP130 (340)	EF151500.1	F: 5′-AGATAGAGCCGTCTCATACTCATGG-3′ R: 5′-TCATTTGGAGCAGTCCATTCTACCC-3′

* Primers were designed to flank intron(s).