

Table S1 Primers used for RT-qPCR

Gene	Primer	Sequence	Size (bp)	Accession numbers
<i>Clqa</i>	Forward	CAGGAGAACGTGTACCAGAGCAAC	122	XM_012152629.2
	Reverse	CTCCGAGAGGACCTGATGGACAG		
<i>Clr</i>	Forward	CCCAGACTACCGCCAGGAAGAG	109	XM_012175492.2
	Reverse	TGGGAGGCAGATTGGCAGGAG		
<i>ClS</i>	Forward	CCTGGCAAGTCTTCTTCTCGAACC	130	XM_004006917.4
	Reverse	ACCACTGAGGAGGACCCAACATAC		
<i>C2</i>	Forward	CCACCAATCCCATCCAGCAGAAG	95	XM_012100922.4
	Reverse	GGCGTCCAGGAGCAGGTAGAG		
<i>C3</i>	Forward	CGCCACCAGCAGACTATAACGATC	105	XM_027969774.1
	Reverse	AGCAGCCTTGACCTCCACCTC		
<i>C4a</i>	Forward	TTCAGGACAGGTGGTGAGAGGATC	167	XM_027958803.1
	Reverse	GGAGGAGATGGAGGCGACAGAG		
<i>C5b</i>	Forward	GCTACGCTGGTGTTACTCTGGATC	157	XM_004003966.3
	Reverse	GCAGACATGACCTCGCCTATAAGC		
<i>C9</i>	Forward	GCCGCAACAGAGTGGTGGAAG	138	XM_004017026.3
	Reverse	TGCCATCCCTAACTCGGTCACAG		
<i>GAPDH</i>	Forward	GGGTCATCATCTCTGCACCT	176	NM_001190390.1
	Reverse	GGTCATAAGTCCCTCCACGA		