



Figure S1. Amplification specificity of primers in RT-PCR and qRT-PCR.

A: single amplicon of the expected size for each gene was visualized on a 1 % agarose gel. M, marker. B: single peaks in melt curve analysis.

Table S1. Oligonucleotide primers for candidate qRT-PCR reference genes in *H.cunea*

Gene	GenBank Accession Number	Primer Sequence (5'→3')	Amplicon Size (bp)	Slope	Efficiency (%)	R ²
β -actin (<i>β-actin</i>)	OL409114	F: AATGGCTCCGGTATGTGCAA R: ACGAGTCCTTCCGTCCCATA	104	-2.9051	120.91	0.9978
Ribosomal protein L13 (<i>RPL13</i>)	OL409115	F: TCCATTAAGGCCGGTTGTCC R: TCCATTAAGGCCGGTTGTCC	142	-2.8027	127.40	0.9741
Argininase kinase (<i>AK</i>)	OL409116	F: GTTGAGAGGGAAGCGAGGAT R: GAGTGTGAAAGCGAGCAGAG	130	-3.2262	104.15	0.9997
eukaryotic initiation factor 4A (<i>EIF4A</i>)	OL409117	F: TGCGTCTCGGAAGTGTCTGTTA R: GCAGACGGCTTTTCGAATCC	165	-3.1411	108.14	0.9993
α -tubulin (<i>α-TUB</i>)	OL409118	F: TCCATTAAGGCCGGTTGTCC R: AATGCACGGGCTTACAAGGA	92	-3.0535	112.56	0.9996
Polyubiquitin (<i>UBI</i>)	OL409119	F: AATCTTCGCCGGCAAACAAC R: CAGAAGGCTCCACTTCCAGG	121	-3.1757	106.48	0.9998
Ribosomal protein S26 (<i>RPS26</i>)	OL409120	F: CAGGGACATCAACGAAGCCT R: GGTGCTCTTGGGTGGAGTAC	114	-3.2577	102.75	0.9981
Ribosomal protein S15 (<i>RPS15</i>)	OL409121	F: GCGTCACTCTCAAGATCAGC R: ATGGTGCAGATCCTCCAGTC	101	-3.1853	106.03	0.9994
Serine protease1 (<i>HcSPI</i>)	MH663425	F: CTGGTGGCATGGATGCTAAG R: TTACAGTCCCTCCACACACC	102	-3.6566	87.70	0.9971