

Table S1. List of genes employed in the multiplexed RT-qPCR assay 1.

Protein name	Acronym	Accession number	Product length	Oligo sequences ^a	Function	References
Reference genes						
Glyceraldehyde 3 phosphate dehydrogenase	<i>Gapdh</i>	EZ026309	187	F: AGGTGGAGGCCAAGAAAGTCR: R: TTAGCTAGGAGGCCAGGCA	Housekeeping gene of demonstrated stable expression level	[24,26]
Ribosomal protein S7	<i>Rps7</i>	EZ031290	197	F: CAGGCATGTTACAACAAA R: TCAACCTCTTGTCCAGT		
Ribosomal protein L9	<i>Rpl9</i>	EZ026324	292	F: CGTGTAAACGTGTGGTTTGCR: R: TTGACACCTGAATTCCGAC		
Unknown transcript	<i>Ctg1913</i>	EZ040581	280	F: GATTTAACCCACGGCAGTGT R: ATGGTAGGGAGGAGGCTGTT		
Genes of interest						
Toll interleukin receptor	<i>TIR-1</i>	EF090256	137	F: AAAGCCGCAGTCATCAGTT R: GAAATTGGCGTTGAATTCTGT	Toll/TLR pathway	[19]
TNF receptor associated factor 6	<i>TRAF-6</i>	DY583189	127	F: TGATGAATGCTTTCGAG R: ACATGCTTGTCAAGCTGATG		
Mitogen-activated protein kinase/ERK kinase kinase-1	<i>MEKK-1</i>	DY581208 DY581138 DY582675	117	F: CTGCGGATATTGGTCTGT R: TTTCCTTGTGGTGTGATCCC		
Extracellular signal regulated kinases	<i>ERK-2</i>	EZ025389	217	F: CCAAAGGTTACAGCAAGGCT R: TCGGTGCTTTCATTCATA		
P38 mitogen-activated protein kinases.	<i>MAPK-p38</i>	EZ031759	237	F: AAAATCAGCAGTGAATCCGC R: TCGGGTCTGAATACGTAGC		
Part of AP-1 early response transcription factor	<i>AP-1/cFos</i>	EZ016042	177	F: CTGAAAGAGAATTGCTGGC R: GACGATTGACTTCGGACTT		
Part of AP-1 early response transcription factor	<i>AP-1/cJun</i>	EZ020860	366	F: TCGATCGAAGGGACAGTCT R: GTGCTAGTTGCGGTGTTCAA		
c-AMP Activating transcription factor 4	<i>ATF4/5</i>	DY577805	247	F: GGCCAGAACGTATCACCAAT R: TCTTCGAATCAAACCCCTG		
Nuclear factor kappa B	<i>NF-kB</i>	G0002043 DY582971 DY580118 G0000491	152	F: GATGTTGCAGGCTAGITCA R: CTCATATGCAGGTTGGTGGAA		
TRIF-related adaptor molecule	<i>TRAM</i>	EZ047194	157	F: AAGCTAACGGCTACCAAGA R: TGTCATGCAACAGAAAT	IFN pathway	[19]
Apical extracellular protein 60 kDa proteinaceous toxin	<i>Apextrin</i> <i>Tx60-A</i>	EF091848 DY579588	352 207	F: GGATTCTGACCAAAAGGC R: GAGGGCTCTGATATGGGTT F: TACTGCCCTGAGGTTGGCT R: CTAAACATCCCGCTGACTGT	MAC/PF - membrane attack complex/perforin domain.	[19]
Complement factor B precursor	<i>Bf</i>	GO001635	227	F: TTATCCATCCGAGCGCTAAC R: AGGATCATCTTCTCTGCAG	Alternative pathway of complement activation with 6e-77 on 30% coverage to <i>N. vectensis</i> (AB450042).	[42]
Complement C3	<i>C3</i>	EF090257	167	F: CCGCTACACGCTAGACAAAC R: CCGCAGAGTCGATGTCACAA	Alternative pathway of complement activation with 0e-value on 87% coverage to <i>N. vectensis</i> (AB450038).	[42]
Mannose-binding lectin	<i>millectin</i>	EU717895	257	F: AGCGAGTATCCACAAACCC R: GGCTTTCTCGATGTTTCCA	Innate immune system cell surface recognition receptors.	[15,16,23]
C type lectin	<i>CTL-1</i>	GO001638	312	F: GGGTTGTGTAACAGGCTT R: CTTTCCATTCTGGTCTCCCTG	Innate immune system cell surface recognition receptors.	[15,16]
C type lectin	<i>CTL-2</i>	GS01UH10	267	F: CAGGCTGGATCGGACTCAT R: CATGTCACAGTGGTGTACGC		
Hemolytic lectin-1	<i>HL-1</i>	EU863776 EU863777	335	F: TTGGCTCAGAGGGAAACTA R: GCAGAAATGCCCTTGTTGT	Innate immune system cell surface recognition receptors.	[15,16]
Hemolytic lectin-2	<i>HL-2</i>	EU863776	302	F: AACAGTTGAGATAACCGCCG R: TTGATTCCTGGTGCATTGAA		
Hemolytic lectin-3	<i>HL-3</i>	EU863777	379	F: TTCTGGAGATGGGTAACGC R: TCGTTCTCAGCGTGTGTT		

^a Forward (AGGTGACACTATAGAATA) and reverse (GTACGACTCACTATAGGGA) fluorescent universal tags, omitted from the oligo sequences in the table, were added at the 5' end of all primers.