

Table S2 Quantitative PCR primers used in this study

Genes	Reference/Accession Nr.	Forward (5'>3')	Reverse (5'>3')
<i>GAPDH</i>	Yan <i>et al.</i> , 2014	TTGTCTCCTGCGACTTCA	CCACCACCCTGTTACTGTT
<i>TLR1</i>	NM_001285605.1	ACAATCCATTCCAATGTTCC	ACAATGGTGACAATCAGCAG
<i>TLR2</i>	XM_013970466.1	CTGTGTGCGTCTTCCTCAGA	TCAGGGAGCAGAGTAACCAGA
<i>TLR4</i>	NM_001285574.1	GGTTTCCACAAAAGCCGTAA	AGGACGATGAAGATGATGCC
<i>TLR5</i>	NM_001285699.1	ACCTGGGTGGAAGTCAGATA	GGTTGAGGGAAAAATCAATG
<i>TLR6</i>	NM_001285540.1	AGGCCAAGTATCAAGAGACG	AGAGGACAGTCACAGCAACA
<i>TLR10</i>	NM_001285541.1	TTGCATGATGGAATCAAAAC	AACCAATTGGAAGATGAGGA
MyD88	XM_013973392.1	ACAAGCCAATGAAGAAAGAG	GAGGCGAGTCCAGAACC
<i>IFN-γ</i>	NM_001285682.1	TGATTCAAATTCGGTGGAT	GCAGGCAGGAGAACCATTAC
<i>IL-1β</i>	XM_013967700.1	CATGTGTGCTGAAGGCTCTC	AGTGTCGGCGTATCACCTTT
<i>IL-6</i>	NM_001285640.1	CCAATCTGGGTTCAATCAGG	ACCCACTCGTTTGAGGACTG
<i>IL-10</i>	XM_005690416.2	TTAAGGGTTACCTGGGTTGC	CCCTCTCTTGAGCATATTGA
<i>NHE1</i>	Yan <i>et al.</i> , 2014	CCTCTACAGCTACATGGCCTAC	GGGAGATGTTGGCTTCCA
<i>NHE2</i>	Yan <i>et al.</i> , 2014	TTGGAGAGTCCCTGCTGAAC	GGCCGTGATGTAGGACAAAT
<i>NHE3</i>	Yan <i>et al.</i> , 2014	AGCTACGTGGCCGAGGG	AGACAGAGGCCTCCACGGT

Genes	Reference/Accession Nr.	Forward (5'>3')	Reverse (5'>3')
<i>Na⁺/K⁺ ATPase</i>	Yan <i>et al.</i> , 2014	TGAGCATCCCAGTGTTGT	CCTTGTCAGATACTTCCT
<i>vH ATPase</i>	Yan <i>et al.</i> , 2014	TTTTATTGAACAAGAAGCCAATGA	GATTCATCAAATTGGACATCTGAA
<i>PAT1</i>	Yan <i>et al.</i> , 2014	CCTTGAGGCACGGCTAC	GCACCAGACTCCGAGACATA
<i>AE2</i>	Yan <i>et al.</i> , 2014	AGCAGCAACAACCTGGAGT	GGTGAAACGGGAGACGAA
<i>DRA</i>	Yan <i>et al.</i> , 2014	TTTAAAGTCCTAGAGTCCGTA	CGCTGATTTATTTCTTTAACCAC
<i>MCT1</i>	Yan <i>et al.</i> , 2014	ACCAGTTTTAGGTCGTCTCA	GGCTTCTCAGCAACATCTACA
<i>MCT4</i>	Yan <i>et al.</i> , 2014	GTTTGGGATAGGCTACAGTGACACA	GCAGCCAAAGCGATTCA