

Supplementary Table 1

L. infantum primer sequences and qPCR efficiency. Gene accession numbers, forward/reverse primer sequences, and primer efficiencies for each gene target for the *L. infantum* qPCR assays are presented. Primer efficiency was calculated by assaying diluted cDNA.

Gene Target	Accession	Forward Primer	Reverse Primer	Primer Efficiency
<i>UBH</i>	LinJ.29.2410	5'-AACGTGAACAACCTGGATGTGCGTC-3'	5'-ATGGTGCCAAGCTTGACACACGCA-3'	103.7%
<i>AML1</i>	LinJ.08.0680	5'- AAGTGCCTGGAGCTGAGTGACGGC-3'	5'-ACGGGAGCATCAGGAAGATAATGT-3'	94.5%
<i>AML2</i>	LinJ.08.0780	5'-CATCGTCGTGTACGGCTTGGCTTT-3'	5'- TGTAGTAGTCCACCATCATGAGCG-3'	94.2%
<i>P27</i>	LinJ.28.1070	5'-TTACCGCTCCGCTACAACGTGCT-3'	5'- ACCACAGGGAGTAGATCACGACAA-3'	91.6%
<i>ATG8</i>	LinJ.19.1660	5'- ACAAGAGCAAGTTCTTGGTACCGC-3'	5'- AAACAGCACTCTCATGAGCGACCT-3'	98.4%
<i>A2</i>	LinJ.22.0670	5'-GTCCGCTTGTGGTGTGCTG-3'	5'- CAACGGCCGCCTTGTGC-3'	92.6%
<i>Pr2C</i>	LinJ.16.1510	5'- ATGAGCATCGCTGCGGACAT-3'	5'- GGTTGTGGATCTTCTGCTTGC-3'	90%