Table S2. Primers used for quantitative reverse transcriptase-polymerase chain reaction (qRT-PCR) analysis.

Gene description	Gene[a]	Forward primer sequence	Reverse primer sequence
Control	*gyrB*	5'-TATGAGTGGCTTGGCGTTAG-3'	5'-ACACCGCTTCTGGATAATAGC-3'
Glycogen synthesis	*malQ*	5'-TGAACAGGCGACGAAGTATG-3'	5'-CCGACGAGAAGATGAGCTATAAG-3'
Glycogen synthesis	*glgB*	5'-CAGTAAAGACGGAGAGCGAATTA-3'	5'-CCACATAGTTACCCGCCATATT-3'
Glycogen synthesis	*glgX*	5'-AGCGATGTTTGCGCTAGAA-3'	5'-TCGATACCCGCTTGATGTAATG-3'

Glycogen synthesis	***glgC***	**5'-GGGAACCCAGTACAAACCTATC-3'**	**5'-TAATCACGCAACCACCAGAA-3'**
Glycogen synthesis	***glgA***	**5'-ACATGCTTGCTCGGAGTTATAC-3'**	**5'-GGATACGCCGGTAAGACAATAC-3'**
Glycogen synthesis	***glgP***	**5'-CGGAATTTATCCGTCGTGTCT-3'**	**5'-GCCACGCCATTCACTTTATTC-3'**

[a]Genes selected were those present in the putative glycogen synthesis locus of *P. multocida*.