Supplementary Table 3

Primer name	Function	Primer sequence (5' to 3')
hasA qPCR F1	Quantification of *hasA* transcription	CATTGTTCATCGGTCAGAG
hasA qPCR R1		GTATCTGAGTCAACG
hasB qPCR F2	Quantification of *hasB* transcription	GGTCTAGATGCTCAGC
hasB qPCR R2		CTTTGGCAGGCAATAG
hasC qPCR F1	Quantification of *hasC* transcription	CGAGCTCGAATACAATC
hasC qPCR R1		CTCTAGGGTGGCTCTG
gyrA forward	Quantification of *gyrA* transcription	CTATGAAGCGATGGTCCGTATGG
gyrA reverse		CCATAGAACCAAAGTTTCCATGACC
has tan F1	Detection of tandem duplication of the *has* locus	GCTGCCAATAGAGCTAAGTC
has tan R1		CTGATCAAGCGCGATAC