

Table S2. Quantitative RT-PCR analysis of amplified versus non-amplified samples

	Sample no. 33		Sample no. 55	
	Non-amplified . \log_2 mean expression (n=3)	Amplified mRNA. \log_2 mean expression (n=3)	Non-amplified . \log_2 mean expression (n=3)	Amplified (\log_2) Mean expression (\log_2)
<i>glyA</i>	0.87	0.87	0.68	0.74
<i>pykA</i>	1.13	1.13	1.02	1.02
<i>tpiA</i>	1.20	1.09	0.98	0.98
<i>luxS</i>	0.68	0.87	0.53	0.71
<i>cirA</i>	0.70	0.85	0.50	0.74
<i>hybB</i>	0.91	0.85	0.56	0.60
<i>fdxG</i>	0.92	1.02	0.81	0.87
<i>tonB1</i>	0.78	0.73	0.58	0.60
<i>hlyX</i>	0.85	0.87	0.89	0.78
<i>hyaA</i>	0.44	0.76	0.06	0.52
<i>tbpA1</i>	0.92	0.87	0.82	0.76
<i>tonB2</i>	0.26	0.78	0.12	0.62
<i>lldD</i>	0.32	0.73	0.21	0.63