

Supplementary Table 3: Oligonucleotides used in this study

Gene identifier ^a	Gene name ^a	Primer	
		Direction	Sequence (5' to 3') ^b
SACOL0583	<i>rplK</i>	Forward	GCACATCGTGGCTAAAAAAG
		Reverse	<u>CTAATACGACTCACTATAGGGAGATTATTCTACAACGATACCCA</u>
SACOL0594	<i>tuf</i>	Forward	GTATCTGCTGCTGACGGTCC
		Reverse	<u>CTAATACGACTCACTATAGGGAGAGGCATTACCATTTCAGTACC</u>
SACOL0838	<i>gapA1</i>	Forward	CGTGCTCGTGCAGCGGCAG
		Reverse	<u>CTAATACGACTCACTATAGGGAGATTATTTAGAAAAGTTCAGCTAAG</u>
SACOL1734	<i>gapA2</i>	Forward	GATTTAAGACGTGCACGTTTCATG
		Reverse	<u>CTAATACGACTCACTATAGGGAGATTAACTTGCACTTACAGTTTC</u>
SACOL1838	<i>pckA</i>	Forward	ATGTCAGTAGACACATACAC
		Reverse	<u>CTAATACGACTCACTATAGGGAGAATTGCACCATATTTGATTGC</u>

^a Based on TIGR annotation (<http://www.tigr.org>).

^b The underlined nucleotides at the 5' end represent the binding site for the T7 RNA polymerase