Gene	putative gene function	oligonucleotide	oligonucleotide sequence (5´-3‘)
gapA	glyceraldehyde-3-phosphate dehydrogenase	gapA_fwd	gttttcccagtcacgacgttgta**tgaagtatgactccactcacgg**
		gapA_rev	ttgtgagcggataacaatttc**aacgcctttcattgcgccttcggaa**
infB	translation initiation factor 2	infB_fwd	gttttcccagtcacgacgttgta**ctctctgctggactacattcg**
		infB_rev	ttgtgagcggataacaatttc**cgctttcagctccagaacttc**
mdh	malate dehydrogenase	mdh_fwd	gttttcccagtcacgacgttgta**cccaactgccttcaggttcag**
		mdh_rev	ttgtgagcggataacaatttc**ccttccacgtaggcgcattcc**
pgi	phosphoglucose isomerase	pgi_fwd	gttttcccagtcacgacgttgta**gagaaaaacctgccggtgctgctg**
		pgi_rev	ttgtgagcggataacaatttc**cggttaatcaggccgttagtggagc**
phoE	phosphoporine E	phoE_fwd	gttttcccagtcacgacgttgta**acctggcgcaacaccgatttcttc**
		phoE_rev	ttgtgagcggataacaatttc**ttcagctggttgattttgtaatccac**
rpoB	RNA polymerase subunit β	rpoB_fwd	gttttcccagtcacgacgttgta**ggcgaaatggcggaaaacca**
		rpoB_rev	ttgtgagcggataacaatttc**gagtcttcgaagttgtaacc**
tonB	periplasmic energy transducer	tonB_fwd	gttttcccagtcacgacgttgta**ctctatacttcggtacatcaggtt**
		tonB_rev	ttgtgagcggataacaatttc**cctgtttggcggccagcacctggt**
	sequencing oligonucleotides	MLST_seq_fwd	gttttcccagtcacgacgttgta
		MLST_seq_rev	ttgtgagcggataacaatttc
		phoE_seq_fwd	tttcttcggcgtggtagatcc
		phoE_seq_rev	gtaatccacaaaggcattc

Specific oligonucleotides: bold part binds target gene, underlined part (overhang) serves as annealing site for sequencing primer (phoE was sequenced with distinct nested primers)