

Strain	Forward	Reverse
<i>rpoC</i> re-sequencing	5'-TTACTCGTTATCAGAACCGCCAG-3'	5'-TTATTAAAGTTTCTGAAAGCGC-3'
<i>araB</i> mRNA estimation	5'-CAAGCACGGTTTTTCAGTG-3'	5'-GATTCTGTGCGAGCTTTGGC-3'
<i>rpoA</i> mRNA estimation	5'-CATCACGTAAGTCAACGAAAGC -3'	5'-GCTGGTCATCGAAATGGAAAC-3'