

Genetic Determinants of Tigecycline Resistance in *Mycobacteroides abscessus*

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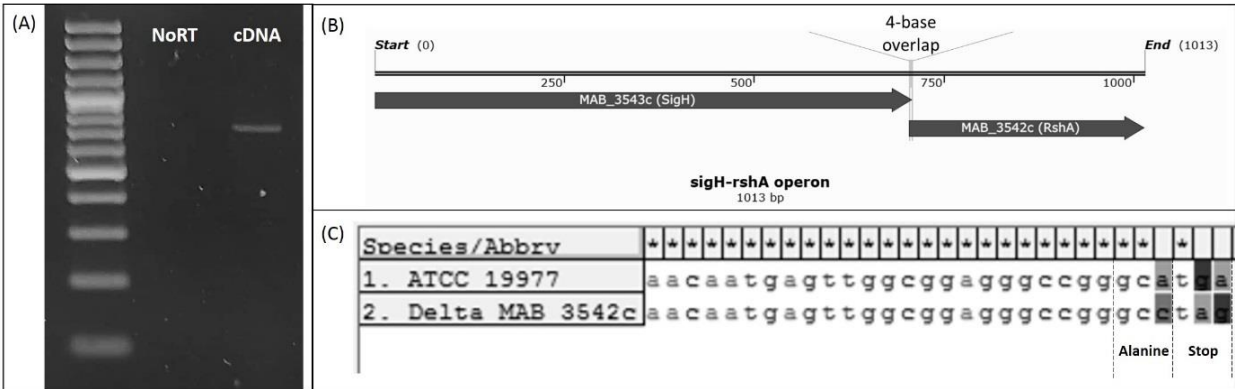


Figure S1. (A) The *sigH* (*MAB_3543c*) and *rshA* (*MAB_3542c*) genes are transcribed as an operon. RT-PCR analysis with the forward primer annealed to the *MAB_3543c* gene and the reverse primer annealed to the *MAB_3542c* gene. cDNA was prepared from the RNA of ATCC 19977. NoRT: no-reverse transcription control. (B) Both genes are neighbor genes in the ATCC 19977 genome with a 4-base overlap. (C) Partial DNA sequences of *MAB_3543c* from ATCC 19977 and Δ MAB_3542c.