

Supplemental material - table S1. GenBank accession numbers used in development of assay B.

Bacterial species	GenBank accession number	Bacterial species	GenBank accession number
<i>M. abscessus</i>	AY271329	<i>M. microti</i>	AY271315
<i>M. africanum</i>	AY271330	<i>M. scrofulaceum</i>	AY271317
<i>M. avium</i>	AY271322	<i>M. smegmatis</i>	AY271345
<i>M. bovis</i>	AY271323	<i>M. szulgai</i>	AY271319
<i>M. celatum type I</i>	AY271337	<i>M. tuberculosis</i>	CP000611
<i>M. fortuitum I</i>	AY271325	<i>M. ulcerans</i> Agy99	CP000325
<i>M. gordonae type I</i>	AY271342	<i>M. peregrinum</i>	AY271324
<i>M. haemophilum</i>	AY271331	<i>M. xenopi</i>	AY271332
<i>M. kansasii type I</i>	AY271333	<i>E. coli</i>	NC009800
<i>M. leprae</i>	NC002677	<i>Staph. aureus</i>	CP003194
<i>M. malmoense</i>	AY271321	<i>Staph. epidermidis</i>	NC004461
<i>M. marinum</i>	AY271318	<i>Str. pyogenes</i>	CP000260

Table S1 shows the GenBank accession numbers (PubMed, NCBI) that were used for in-silico development of assay B by alignment of (myco-) bacterial *rpoB*-genes using DNASIS Max software (MiraiBio, San Francisco, California).