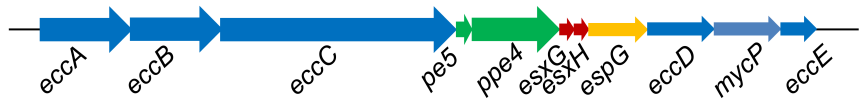


Supplementary Figure 1

A



B



Mtb *rv0282* upstream sequence:

```

1 cccgggctc cataatgaaaa tcattgttcagtaagc acac tctgcataatcgggctaccaa cgaaatggagtacgggtcat
81 gatcttgccagccgtgccta aaagcttggccgcagggccg agtcgattggtcgcggtcgc ctcgacac ttagcttatgca
161 atgctaatttcggggcaag ttcaggcgatcggccg ATG GCG
  
```

Msmeg *msmeg0615* upstream sequence:

```

1 cagtacagtcctgatgtggg cccgcaaggggtgagccgct tcacggcccagagccccac gtgcgctctcgacag ttaact
81 tatgtaatgctaacttctcg aaagttagcttaggcataca taaggagagagggcggggc gtgcgctcccgcgcacgcat
161 atgggaagtgcacgcttgc ggetccgcctcagggagcgc cccgggtcgaccgcgat GTG GTC
  
```

C

```

Mtb: ttagcttatgcaatgctaa
    |||*|||*|||*|||*|||
Msmeg: ttaacttatgtaatgctaa
  
```

Supplementary Figure 1. Comparison of the *esx3* Locus in Mtb and Msmeg

(A) The *esx3* locus spans from *eccA*₃ (*Rv0282*) to *eccE*₃ (*Rv0292*) in Mtb, and *eccA*₃ (*msmeg0615*) to *eccE*₃ (*msmeg0625*) in Msmeg. (B) Upstream promoter regions from *esx3*_{Mtb} and *esx3*_{Msmeg}; IdeR (orange) and Zur (green) recognition sites are indicated. The beginning of the protein coding sequence is in bold. (C) Nucleotide alignment of IdeR boxes from Mtb and Msmeg.

Supplementary Table 1. Primers Used	
Name	Sequence
TBEsx3JMF	AAC ATG AGG CCC TGC AGG AAC ACG GTC ATC CGC CGC AGA TAG TCC AAC TGG GAA CTG GCG CAG TTC CTC
TBEsx3JMR	CAT CGC TCG TTG TAG ACC AGT TGA AAA AAC GGG TGC TCC TCG GCT CCC TCC TCC ATA TGC ACT GAG CGT CAG
Int-Ab-F	CTA AGG AAG CTA AAA TGG AGA AAA AAA TCA CTG GAT ATA CCA CCG TTG ATA TAT CCC AAT GGT CTG ACG CTC AGT GGA AC
Int-Ab-R	GCC CAT GGT GAA AAC GGG GGC GAA GAA GTT GTC CAT ATT GGC CAC GTT TAA ATC AAA ACT GTA CGA TCG ACT GCC AGG C
<i>TB esxG-F</i>	TAT CCC ACA GTT GGT GGC CT
<i>TB esxG-R</i>	GAC TCC CCC TGG TGA AAC G
<i>TB ppe4-F</i>	TGG CTC TTA GCG TTG CTT CA
<i>TB ppe4-R</i>	CGT GAG GAA GGC GAT GAT GA
<i>TB mbtI-F</i>	TTC TGT CCC GTT GTG TCG AA
<i>TB mbtI-R</i>	CTG TAA CCC AGA GCA CGG AT
<i>TB bfrB-F</i>	TAA CGA ATT CAC AGC GGC AC
<i>TB bfrB-R</i>	CGA CCG CTT GGC TGT AAA AA
<i>TB 16S-F</i>	TCG TGT CGT GAG ATG TTG GG
<i>TB 16S-R</i>	TAG CAT GTG TGA AGC CCT GG
<i>Smeg esxG-F</i>	TCT TCT CGA CGC TCA CAT CC
<i>Smeg esxG-R</i>	GGA AGG CCT GCG AAC TCA T
<i>Smeg mbtI-F</i>	CCG GTA TCG AGA CTG CAT CC
<i>Smeg mbtI-R</i>	GTA GAC GAC GTA CTC CTC GC
<i>Smeg bfrB-F</i>	ACG AAA TTC CAC GCC CTC AT
<i>Smeg bfrB-R</i>	TAG AAG TGC TTG GCG AGC TG
<i>Smeg dnaA-F</i>	CCG GGT ATG CGT GTC AAG TA
<i>Smeg dnaA-R</i>	AGA ACT CCT CCT GGA TGC CT