

Supplementary Information for:

Esx paralogs are functionally equivalent to ESX-1 proteins but are dispensable for virulence in *M. marinum*

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Running Head: Gene duplications of ESX-1 virulence factors

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Supplementary Methods

Crystal violet assay. 5×10^5 RAW 264.7 cells were seeded overnight in a 24 well TC treated plate (CELLSTAR, Greiner Bio-one, Monroe, NC). Bacteria were grown in 7H9 broth for 3 days, washed three times with sterile PBS and syringe passaged to break up clumps. RAW 264.7 cells were incubated with 1×10^6 bacteria (MOI 2) for 2 hours at 37°C. Each infection was performed in technical triplicate. Wells were treated with 100µg/mL gentamycin for 1 hour at 37°C to kill extracellular bacteria. Each well was washed three times with PBS and then replenished with DMEM media containing FBS. After 2 days, the media was removed and the cells were fixed with 4% paraformaldehyde (PFA) in PBS for 10 min. at room temperature. The wells were stained with 0.1% crystal violet (Alfa Aesar) in dH₂O for 30 min. at room temperature, with gentle rocking. The plate was washed by carefully submerging in a tray of distilled water, under running water until the rinses were clear. The plate was allowed to dry completely and then imaged with a camera.

Table S1. *M. marinum* strains used in this study.

Strain	Description	Reference
M	Wild-type strain (WT)	ATCC BAA-535
Δ RD1	Unmarked deletion of <i>eccCb1'-espK'</i>	(1)
Δ RD1 Δ esxB_1	Δ RD1, unmarked deletion of <i>esxB_1</i> (<i>MMAR_0187</i>)	This study
Δ esxBA	Δ esxBA, Kan ^r	(2)
Δ esxBA/p _{Mops} esxBA	Δ esxBA, p _{Mops} esxBA integrated at the <i>attB</i> site; Kan ^r , Hyg ^r	This study
Δ esxBA/p _{Mops} esxB_1-esxA_3	Δ esxBA, p _{Mops} esxB_1-esxA_3 integrated at the <i>attB</i> site; Kan ^r , Hyg ^r	This study
Δ esxBA/p _{Mops} esxB_1 M98A -esxA_3	Δ esxBA, p _{Mops} esxB_1 M98A -esxA_3 integrated at the <i>attB</i> site; Kan ^r , Hyg ^r	This study
Δ esxBA/p _{Mops} esxB_2-esxA_2	Δ esxBA, p _{Mops} esxB_2-esxA_2 integrated at the <i>attB</i> site; Kan ^r , Hyg ^r	This study
Δ esxBA/pMH406	Δ esxBA, pMH406 integrated at the <i>attB</i> site; Kan ^r , Hyg ^r . The pMH406 plasmid includes the <i>esxBA</i> genes from <i>M. tb</i> behind the mycobacterial optimal promoter.	Gift from Jeff Cox, and (3)
Δ esxBA/pMH406 <i>esxB</i> L94A	Δ esxBA, pMH406 <i>esxB</i> L94A integrated at the <i>attB</i> site; Kan ^r , Hyg ^r	This study
Δ esxBA/pMH406 <i>esxB</i> S95A	Δ esxBA, pMH406 <i>esxB</i> S95A integrated at the <i>attB</i> site; Kan ^r , Hyg ^r	This study
Δ esxBA/pMH406 <i>esxB</i> S96A	Δ esxBA, pMH406 <i>esxB</i> S96A integrated at the <i>attB</i> site; Kan ^r , Hyg ^r	This study
Δ esxBA/pMH406 <i>esxB</i> Q97A	Δ esxBA, pMH406 <i>esxB</i> Q97A integrated at the <i>attB</i> site; Kan ^r , Hyg ^r	This study
Δ esxBA/pMH406 <i>esxB</i> M98A	Δ esxBA, pMH406 <i>esxB</i> M98A integrated at the <i>attB</i> site; Kan ^r , Hyg ^r	This study
Δ esxBA/pMH406 <i>esxB</i> G99A	Δ esxBA, pMH406 <i>esxB</i> G99A integrated at the <i>attB</i> site; Kan ^r , Hyg ^r	This study
Δ esxBA/pMH406 <i>esxB</i> F100A	Δ esxBA, pMH406 <i>esxB</i> F100A integrated at the <i>attB</i> site; Kan ^r , Hyg ^r	This study
Δ esxB_1	Unmarked deletion of <i>esxB_1</i> (<i>MMAR_0187</i>)	This study
Δ esxB_1/p _{Mops} esxB_1-esxA_3	Δ esxB_1, p _{Mops} esxB_1-esxA_3 integrated at the <i>attB</i> site; Hyg ^r	This study
Δ MMAR_0184-esxA_2 (Δ extESX-6)	Unmarked deletion of <i>MMAR_0184</i> - <i>MMAR_0196</i>	This study

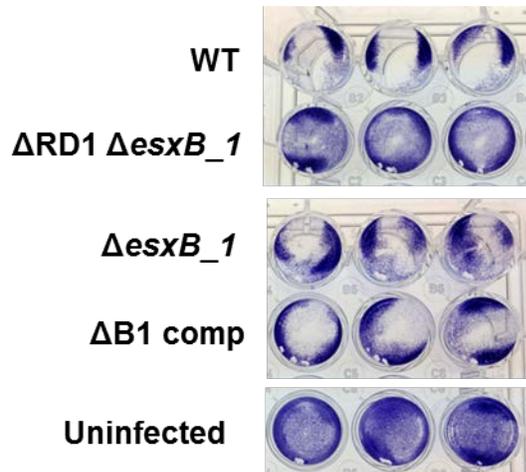
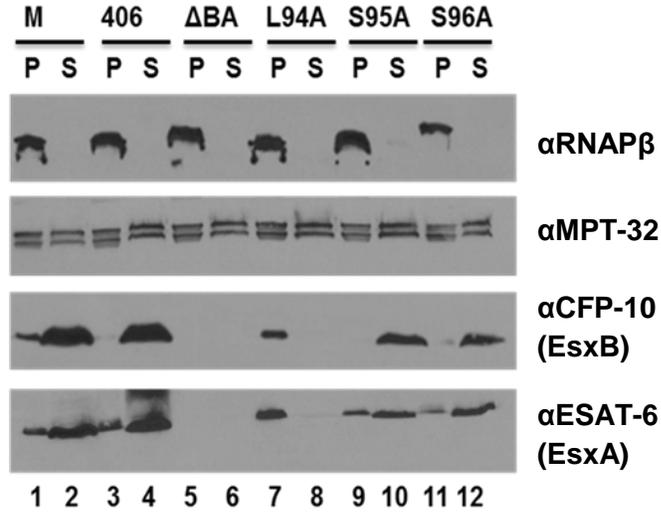


Figure S1. The *esxB_1* gene is dispensable for cytotoxicity to RAW cells. RAW264.7 cells were infected with *M. marinum* strains at an MOI of 2. 2 days post infection, the cells were fixed and stained with crystal violet. Clearing of the RAW cell monolayer indicates cytotoxicity. The figure is representative of 3 independent infections with each strain tested in triplicate.

A.



B.

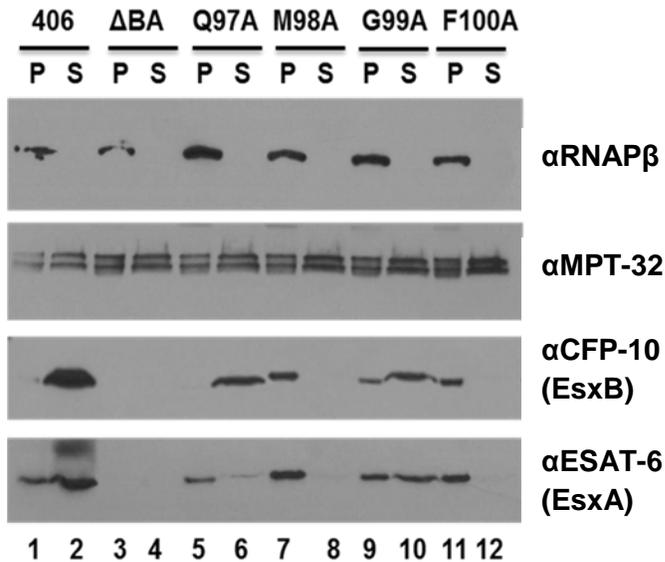


Figure S2. *M. tb* EsxB in *M. marinum* is recognized for ESX-1 export by its C-terminus. EsxB_{MT} are secreted from *M. marinum* as demonstrated by an ESX-1 secretion assay. 10μg of protein was loaded. “M” is the *M. marinum* WT strain. “406” refers to the Δ*esxB*/pMH406 strain. “ΔBA” refers to the Δ*esxB* strain. The remaining strains are Δ*esxB*/pMH406 strains which encode a mutagenized EsxB_{MT} protein from an integrated plasmid (e.g. Δ*esxB*/pMH406 *esxB* L94A). RNAP-β serves as a lysis control. Mpt-32, a Sec-secreted protein, is the loading control. *In vitro* production (P, pellet) and secretion (S, supernatant) of ESX-1 substrates, CFP-10 (EsxB) and ESAT-6 (EsxA) were analyzed. The western blot is representative of 3 independent experiments.

A.

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Mtb EsxB 1 MAEMKTDAAATLAQEAGNFERISGDLKTQIDQVESTAGSLQCGQWRGAAGTAAQAAVVRFQE
Mm EsxB 1 MAEMKTDAAATLAQEAGNFERISGDLKTQIDQVESTAGSLQAQWRGAAGTAAQAAVVRFQE
Mm EsxB_1 1 MAEMKTDAAATLAQEAGNFERISGDLKTQIDQVESTAGSLQAQWRGAAGTAAQAAVVRFQE
Mm EsxB_2 1 MAEMKTDAAATLTGOAHQFERIADDLKASIRRVETTAAGLLLAWRGDAGHAAQAAIALFHQ

61 AANKQKQELDEISTNIRQAGVQYSRADEEQQQALSSQMGF
61 AANKQKAEELDEISTNIRQAGVQYSRADDEEQQQALSSQMGF
61 AANKQKAEELDEISTNIRQAGVQYSRADDEEQQQALSSQMGF
61 AATQQVKKLNEISTKIWTAQDYTDTDHETGLSNQMNF
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B.

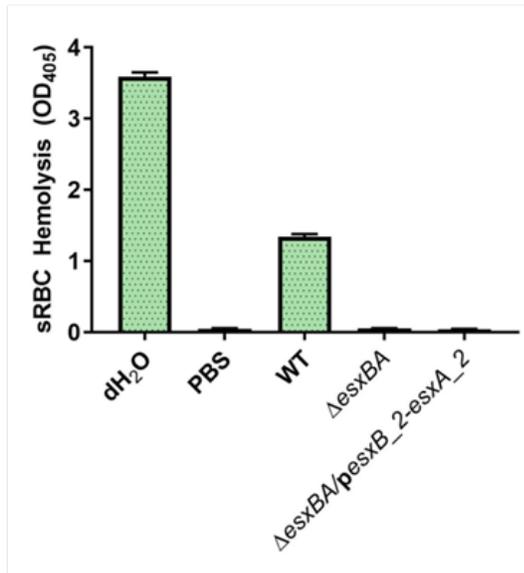


Figure S3. The EsxB₂ protein differs from and is not functionally equivalent to EsxB, (A) Alignment of the *M. tb* EsxB, *M. marinum* EsxB, *M. marinum* EsxB₁, and *M. marinum* EsxB₂ protein sequences. *M. marinum* EsxB and EsxB₁ are 100% identical at the protein level. The *M. marinum* EsxB and EsxB₁ proteins differ from *M. tb* EsxB by 3 amino acid residues. While the first 11 amino acids of the EsxB₂ protein are completely conserved among all EsxB proteins shown here, the remaining protein sequence shows variation. There is lack of conservation of the ESX-1 C-terminus (last 7 amino acids). **(B)** Introduction of an *esxB₂* expression plasmid does not restore hemolytic activity to the Δ*esxB*A strain, demonstrating the EsxB₂ and EsxA₂ proteins are not functionally equivalent to the EsxB_A proteins. 3 isolates were tested. Error bars represent the standard deviation of three technical replicates.

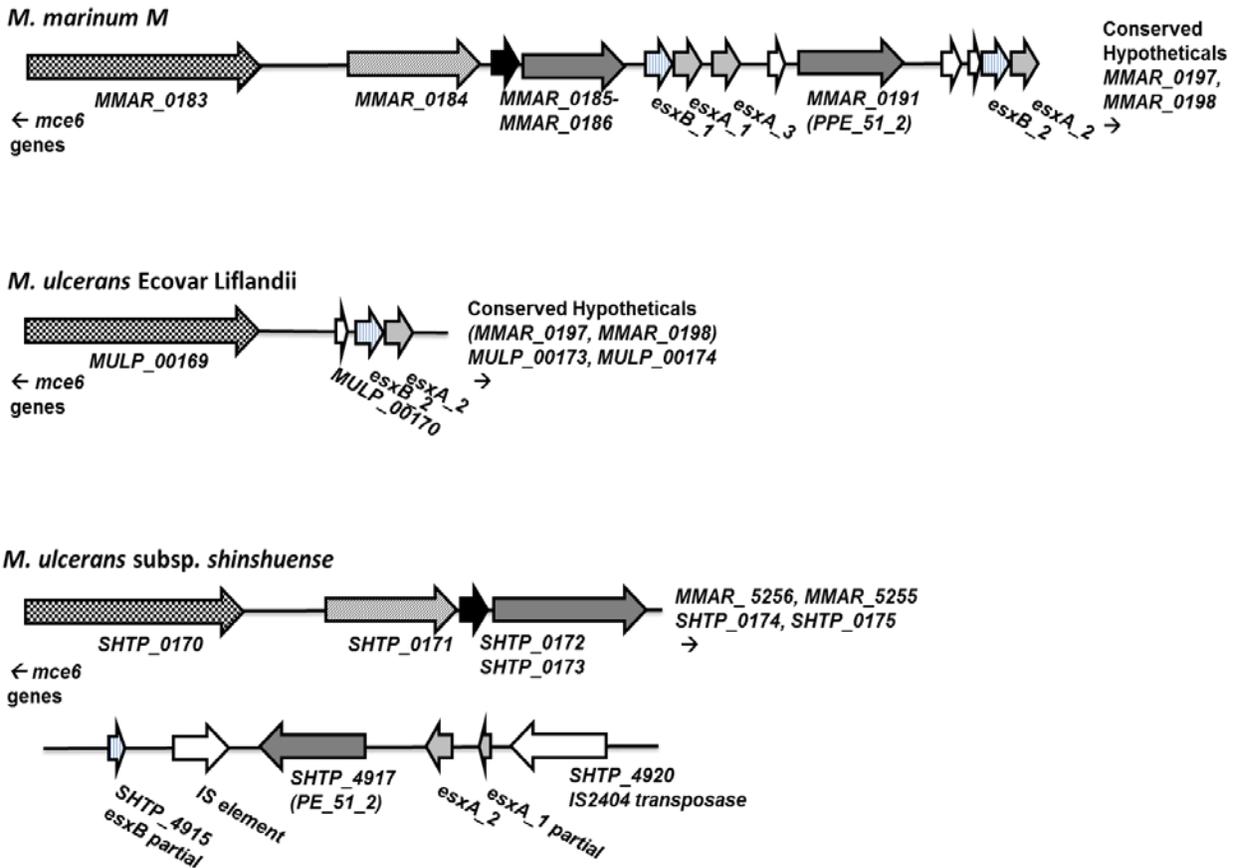


Figure S4. The *esx-6* locus is partially present in some *M. ulcerans* strains. Upstream of the *Esx-6* region is *MMAR_0183*, which encodes a PE-PGRS family protein. *MMAR_0183* is found in both *M. ulcerans* strains depicted. The *M. ulcerans* ecovar Liflandii strain has the *esxB_2* locus. *MMAR_0184*, the *eccB* paralog and *pe35/pe68_1* orthologs are present in *M. ulcerans* susp. *shinshuense*. The *esxB_1* and *esxB_2* loci are incomplete in this strain and found in a different area of the chromosome

References

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2. **Gao LY, Guo S, McLaughlin B, Morisaki H, Engel JN, Brown EJ.** 2004. A mycobacterial virulence gene cluster extending RD1 is required for cytolysis, bacterial spreading and ESAT-6 secretion. *Mol Microbiol* **53**:1677-1693.
3. **Guinn KM, Hickey MJ, Mathur SK, Zakei KL, Grotzke JE, Lewinsohn DM, Smith S, Sherman DR.** 2004. Individual RD1-region genes are required for export of ESAT-6/CFP-10 and for virulence of *Mycobacterium tuberculosis*. *Mol Microbiol* **51**:359-370.