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Supporting information for article:

Structure of the ectodomain of the electron transporter Rv2874 from Mycobacterium tuberculosis reveals a thioredoxin-like domain combined with a carbohydrate-binding module

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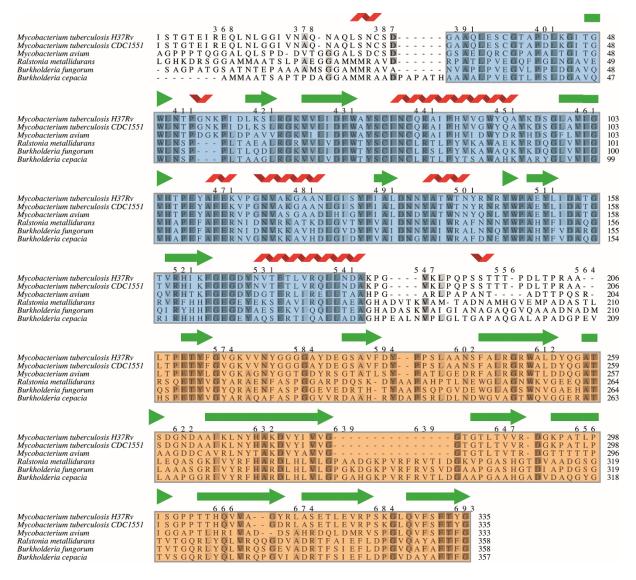


Figure S1 Topology diagram of Rv2874-C. The core components of the thioredoxin fold are indicated by a dashed box.

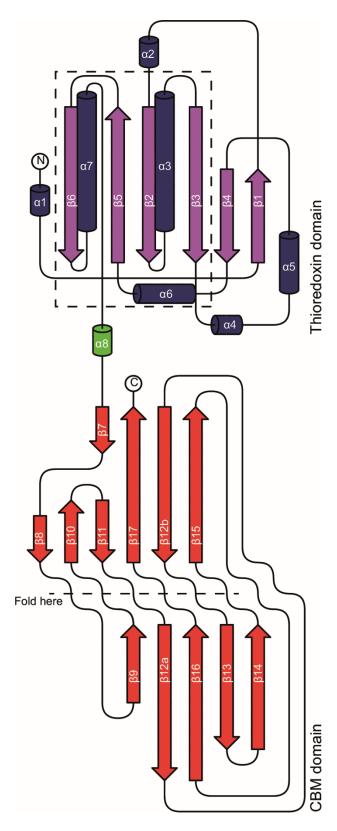


Figure S2 Sequence alignment of Rv2874-C with similar sequences. Secondary structure elements are marked. Residues that form part of the thioredoxin domain are shaded in blue. Residues forming the CBM domain are shaded orange