

# **Proteome-wide Lysine Glutarylation Profiling of the *Mycobacterium tuberculosis* H37Rv**

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Figure S1.(A) MS/MS spectra of a tryptic peptide ion TK(kg)PTSAGQADDALVR from Rv1347c; (B) MS/MS spectra of a tryptic peptide ion VEFEIGHSPK(kg)GPQATGVR from Rv3648c; (C) MS/MS spectra of a tryptic peptide ion PVPGWGSK(kg)RPR from Rv1259; (D) MS/MS spectra of a tryptic peptide ion K(kg)YTAPEISAR from dnak; (E) MS/MS spectra of a tryptic peptide ion K(kg)AIQSAMR from rpsC; (F) MS/MS spectra of a tryptic peptide ion HAK(kg)GWNHFLDR from Rv0576.

