



Fig. S2. Dendrogram of the 30 cutaneous tuberculosis and 58 pulmonary tuberculosis strains with 24-loci Mycobacterial interspersed repetitive unit-variable number tandem repeat (MIRU-VNTR) genotype and spoligotyping for Beijing strains and non-Beijing strains. The dendrogram of MIRU-VNTR results was generated by Bionumerics software using the unweighted pair group method of averages (UPGMA). The spoligotyping assignments are according to the international database SpolDB4.