Supplementary material to article by H. Jiang et al. "Molecular Characteristics of Mycobacterium tuberculosis Strains Isolated from Cutaneous Tuberculosis Patients in China"

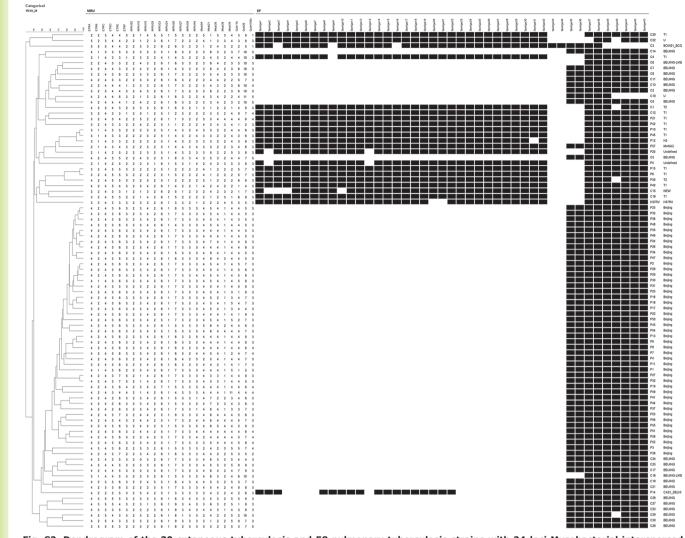


Fig. S2. Dendrogram of the 30 cutaneous tuberculosis and 58 pulmonary tuberculosis strains with 24-loci Mycobacterial interspersed repetitive unit-variable number tandem repeat (MIRU-VNTR) genotype and spoligotyping for Beijing strains and non-Beijing strains. The dendrogram of MIRU-VNTR results was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

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