Next-Generation Sequencing of *Mycobacterium tuberculosis*

Technical Appendix

Technical Appendix Figure. Whole-genome, short-sequencing reads of *Mycobacterium tuberculosis* strain 4542 mapped to complete genome of reference strain H37Rv (NC_00962.3) by using Geneious 9.0 package: genome locus of spacers 7–9 in DR/CRISPR locus. Spacer numbers are given according to 43-spoligotyping (1). DR/CRISPR locus in *M. tuberculosis* is located on complementary strand, whereas spacers are consecutively numbered within the locus, not genome. Such a situation (IS6110 asymmetrically inserted in the direct repeat unit immediately adjacent to a spacer) was discovered by Filliol et al. (2).

References
