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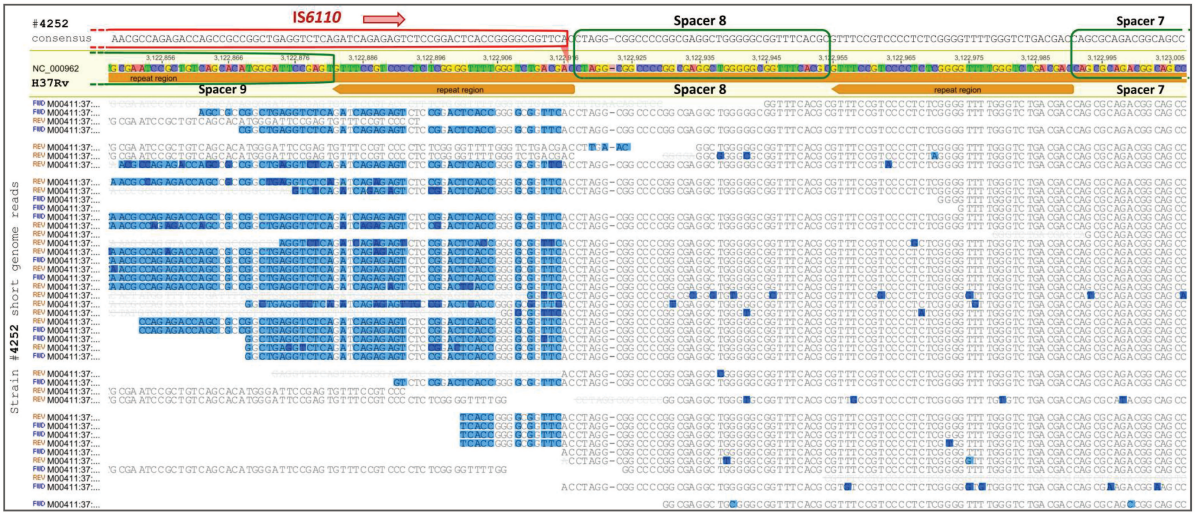
Next-Generation Sequencing of *Mycobacterium tuberculosis*

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 (IS*6110* asymmetrically inserted in the direct repeat unit immediately adjacent to a spacer) was discovered by Filliol et al. (*2*).

References

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