

Whole-genome comparison of the 149 *M. tuberculosis* isolates resulted in a total of 11,516 high confidence polymorphisms including SNPs and indels (Supplementary Table S2):

- 1,693 SNPs/indels exclusive to STR resistant isolates
- 6,781 SNPs/indels exclusive to STR susceptible isolates
- 3,042 SNPs/indels in STR resistant and susceptible isolates

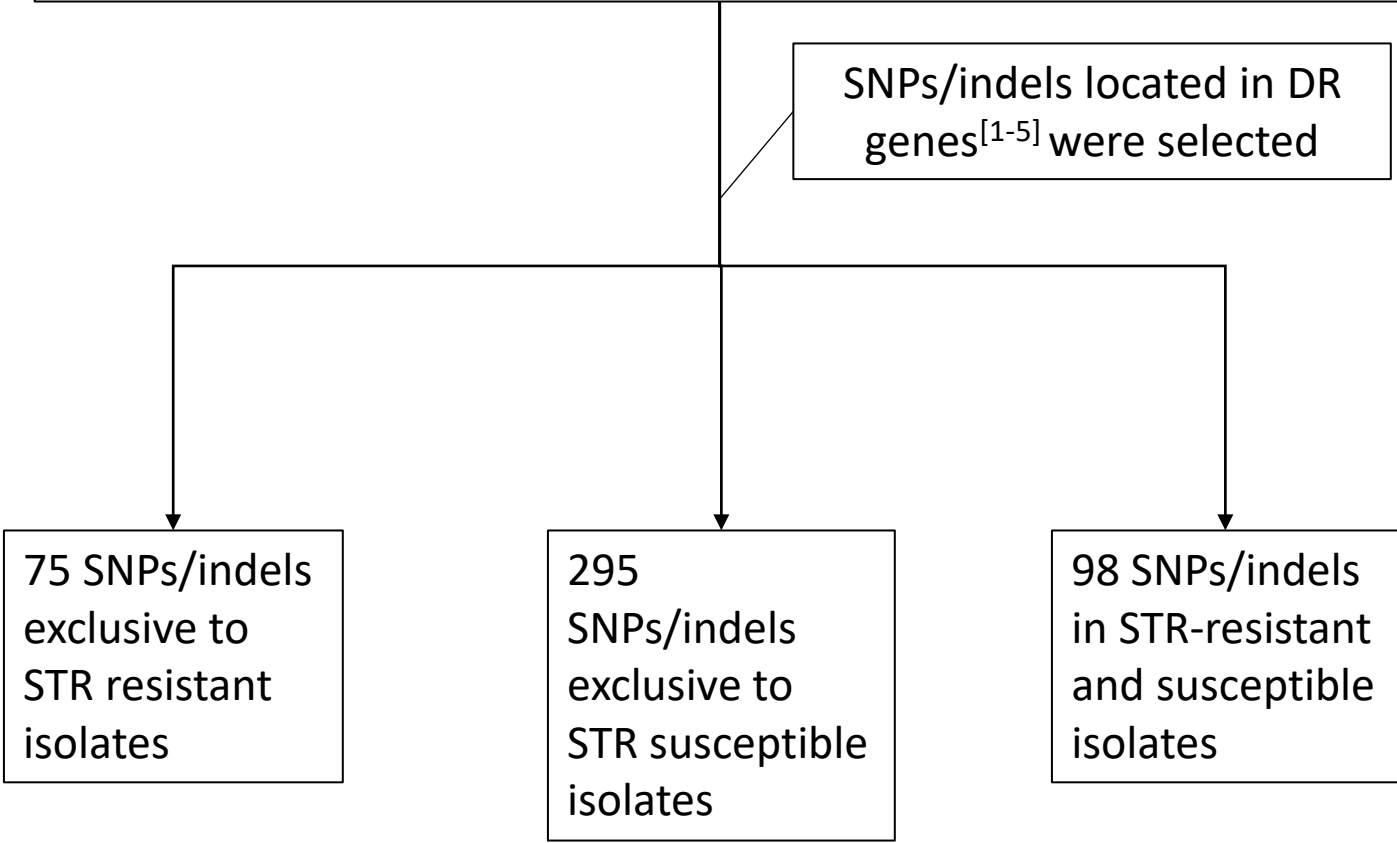


Figure S2: Criteria used to select polymorphisms (SNPs or indels) in genes previously associated with DR and present only in STR resistant *M. tuberculosis* isolates from the studied cohort.

References

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