

Table S1: Spoligotyping information used to classify the samples into the various *M. tuberculosis* genotype families and lineages.

[illegible]

[illegible]

[illegible]

[illegible]

*Inna Vitol, Jeffrey Driscoll, Barry Kreiswirth, Natalia Kurepina, Kristin P. Bennett, "Identifying *Mycobacterium tuberculosis* Complex Strain Families using Spoligotypes", Infection, Genetics and Evolution, Volume 6, Issue 6, November 2006, Pages 491-504." in publications that benefit from this tool. https://tbinsight.cs.rpi.edu/run_tb_lineage.html