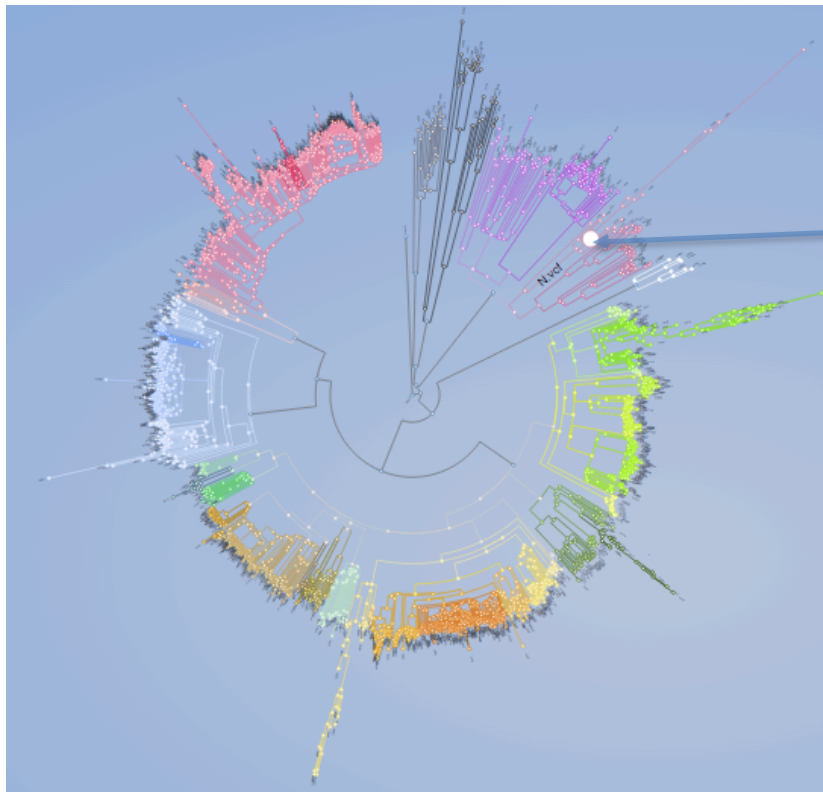
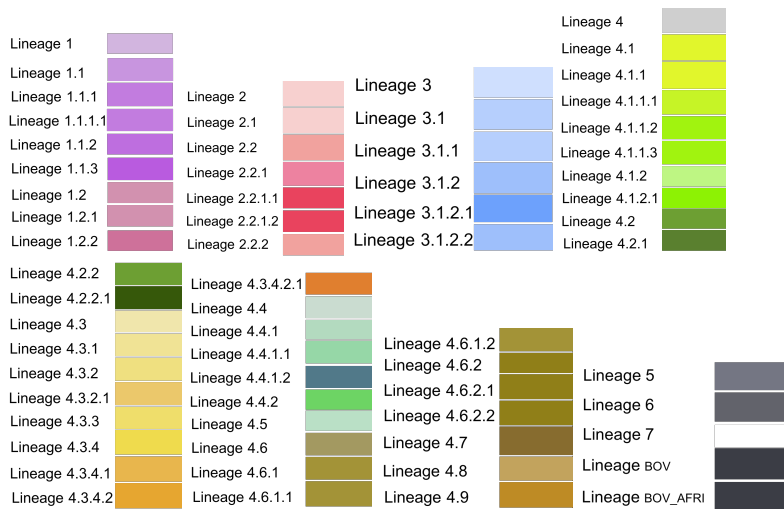


Supplementary Figure S1



Position of isolates CSF-3053, 46-5069 and 43-13838 compared to 1,601 genome of *M. tuberculosis* complex members.



Coll F, McNerney R, Guerra-Assuncao JA, Glynn JR, Perdigo J, Viveiros M, Portugal I, Pain A, Martin N, and Clark TG. 2014a. A robust SNP barcode for typing *Mycobacterium tuberculosis* complex strains. *Nature Communications* 5:4812. 10.1038/ncomms5812