

Supplementary Figure. Cluster dendrogram for *M. tuberculosis* isolates from children with definite TB. Whole genome sequencing (100 bp Paired End) was performed on isolates from 6 of the 7 children with definite TB by Genotypic Technology, Bangalore, India using the Genome Analyzer IIx (Illumina, USA). The sequences for each strain were compared to the reference strain [*M. tuberculosis* H37Rv (NC_000962)]. *M. tuberculosis* strains causing TB meningitis cluster together. This is consistent with several clinical reports and animal studies that have observed the association of specific *M. tuberculosis* strains and lineages with TB meningitis, and suggest that *M. tuberculosis* may possess virulence factors which promote the development of central nervous system disease.

