**Supplementary data S4** - figure illustrating Single amino acid polymorphisms (SAPs) divergence observed in the Beijing clinical isolates analyzed in this work as compared to and absent (abs) from *M. tuberculosis* H37Rv, *M. tuberculosis* H37Ra, *M. tuberculosis* CDC1551, *M. tuberculosis* F11, *M. tuberculosis* KZN-1495, *M. bovis* AF2122-97, *M. bovis* BCG Pasteur 1173P2, *M. bovis* BCG Tokyo 172

Red Bars:
- TB-lab abs = Number of single amino acid polymorphisms (SAPs) observed in *M. tuberculosis* Beijing, but absent from *M. tuberculosis* H37Rv/*M. tuberculosis* H37Ra
- TB-clin abs = Number of single amino acid polymorphisms (SAPs) observed in *M. tuberculosis* Beijing, but absent from *M. tuberculosis* KZN-1495/*M. tuberculosis* F11/*M. tuberculosis* CDC1551
- *M. bovis* abs = Number of single amino acid polymorphisms (SAPs) observed in *M. tuberculosis* Beijing, but absent from *M. bovis* AF2122-97/*M. bovis* BCG Pasteur 1173P2/*M. bovis* BCG Tokyo 172

Black Bars:
The arrow indicates that the details of TB-clin abs are shown by the black bars
- KZN abs = Number of single amino acid polymorphisms (SAPs) observed in *M. tuberculosis* Beijing, but absent from *M. tuberculosis* KZN-1495
- F11 abs = Number of single amino acid polymorphisms (SAPs) observed in *M. tuberculosis* Beijing, but absent from *M. tuberculosis* F11
- CDC abs = Number of single amino acid polymorphisms (SAPs) observed in *M. tuberculosis* Beijing, but absent from *M. tuberculosis* CDC1551