Title: Genomic characterization of MDR/XDR-TB in Kazakhstan by a combination of high-throughput methods predominantly shows the ongoing transmission of the L2/Beijing 94-32 Central Asian/Russian clusters.

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Author’s response to reviews:
The paper was revised according to BMC infectious Diseases guidelines

The age and gender content was suppressed in the Supplementary tables

The Ethical statement was added

Unfortunately the files could not be put in their right order due to a bug in the electronic system, we would be grateful if you could reorder the files.

In response to the Editors, the corresponding author assumes that the list of authors change was made only for technical reasons to provide full concordance between the printed form and the electronic version. In other words, not enough care was put initially when introducing the names in the e-submission system. All authors accepted the final order of authors.