Author’s response to reviews

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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The abstract was formatted according to your requirement (Background, Methods, Results, Conclusions, Keywords).

The contributions of Nurgul Sikhayeva, Maka Akhalaia, Arike Alenova and Pavel Tarlykov was clarified in the appropriate section ("authors contribution").

The funding information was removed from the Acknowledgements and include in the Funding section.

The maps depicted in figure 2 is our own. All details were already given in Methods (lane 205).