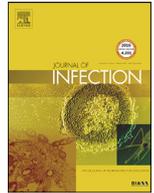




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## Corrigendum

## Corrigendum to 'The epidemic of multidrug resistant tuberculosis in China in historical and phylogenetic perspectives' [Journal of Infection, Volume 80, Issue 4 (2020) Pages 444-453]



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The authors regret that the function to calculate growth rate and results of number of clusters of RR strains were incorrect. Please see correction below:

1. In the method section, "phylogeny and population dynamics", the function used to calculate the growth rate should be as follows. This is a clerical error the results are unaffected.

$$r = n\sqrt{Pe/Ps} - 1$$

2. In the results section, "Transmission and positive selection of MDR-TB strains", this sentence: "six clusters including 7 (19.0%) RR strains had recently developed RR related mutations from rifampicin susceptible (RS) strains;" should be "four clusters including 6 (16.2%) RR strains had recently developed RR related mutations from rifampicin susceptible (RS) strains". These clusters / strains were correctly reported in Fig. 4; the four clusters which contain ten strains six of which are RR are:

- (1) CN3521(R), CN3509(S)
- (2) CN459(R), CN492(S)
- (3) CN2348(R), CN1124(R), CN2206(R), ancestor(S)
- (4) CN3218(R), CN3245(S)

The authors would like to apologise for any inconvenience caused.

### Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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