CORRECTION



Correction: Association between twocomponent systems gene mutation and *Mycobacterium tuberculosis* transmission revealed by whole genome sequencing



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Following publication of the original article [1], it was reported that there was an error in affiliation 1 and that the Background section of the Abstract had to be revised.

The incorrect version of affiliation 1 was: Department of Chinese Medicine Integrated with Western Medicine, College of Traditional Chinese Medicine, Shandong Uni-

The online version of the original article can be found at https://doi.org/10.1186/s12864-023-09788-2.

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The corrected version of affiliation 1 is: Shandong University of Traditional Chinese Medicine, Jinan, Shandong, 250014, People's Republic of China.

The incorrect background text was: Two-component systems (TCSs) assume a pivotal function in *Mycobacte-rium tuberculosis* (*M.tuberculosis*) growth. However, the exact regulatory mechanism of this system needs to be elucidated, and only a few studies have investigated the effect of gene mutations within TCSs on *M.tuberculosis* transmission. This research explored the relationship between TCSs gene mutation and the global transmission of (*M.tuberculosis*).

The corrected background text is: Two-component systems (TCSs) play a crucial role in the growth of *Mycobacterium tuberculosis* (*M. tuberculosis*). However, the precise regulatory mechanism of their contribution remains to be elucidated, and only a limited number of studies have investigated the impact of gene mutations within TCSs on the transmission of *M. tuberculosis*. Therefore, this study aims to explore the relationship between TCSs gene mutation and the global transmission of *M. tuberculosis*.

The original article has been updated.



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References

1. Li Y, Kong X, Li Y, et al. Association between two-component systems gene mutation and Mycobacterium tuberculosis transmission revealed by whole

genome sequencing. BMC Genomics. 2023;24:718. https://doi.org/10.1186/s12864-023-09788-2.

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