

CORRECTION

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# Correction to: Phenotype and multi-omics comparison of *Staphylococcus* and *Streptococcus* uncovers pathogenic traits and predicts zoonotic potential

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**Correction to: *BMC Genomics* 22, 102 (2021)**

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Following publication of the original article [1], it was reported that there was a missing reference in the Methods section and should have been reference [2] at the end of the following sentence:

The GNU “parallel” package was used to perform all of the above steps in parallel [2].

This reference has been inserted in the original article, the subsequent references have been renumbered, and the original article has been corrected.

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## References

1. Zondervan NA, Martins dos Santos VAP, Suarez-Diez M, et al. Phenotype and multi-omics comparison of *Staphylococcus* and *Streptococcus* uncovers pathogenic traits and predicts zoonotic potential. *BMC Genomics*. 2021;22:102. <https://doi.org/10.1186/s12864-021-07388-6>.
2. Tange O. Gnu parallel-the command-line power tool. *USENIX Mag.* 2011; 36(1):42–7.

The original article can be found online at <https://doi.org/10.1186/s12864-021-07388-6>.

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