

## Correction



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Correction to: 'TiDeTree: a Bayesian  
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# Correction to: 'TiDeTree: a Bayesian phylogenetic framework to estimate single-cell trees and population dynamic parameters from genetic lineage tracing data' (2022) by Seidel & Stadler

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We would like to submit the following correction.

In figure 5*a*, the axis labels have to be exchanged, meaning that 'mean triplet distance' should show on the *y*-axis and 'mean RF distance' should show on the *x*-axis.

This has now been corrected on the publisher's website.