

CORRECTION

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Correction: Bayesian kinetic modeling for tracer-based metabolomic data

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The original article can be found online at <https://doi.org/10.1186/s12859-023-05211-5>.

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Following the publication of the original article [1], the authors identified that the funding note was incorrect. The correct funding note is given below.

The incorrect funding is:

This work was supported by National Institutes of Health [1R03CA211835, 5P20GM103436-15, 1P01CA163223-01A1], the Biostatistics and Bioinformatics and Redox Metabolism Shared Resource Facilities of the University of Kentucky Markey Cancer Center [P30CA177558] and University of Kentucky Center for Cancer and Metabolism [1P20GM121327-01].

The correct funding is:

This work was supported by National Institutes of Health [1R03CA211835, 5P20GM103436-15, 1P01CA163223-01A1], the Biostatistics and Bioinformatics and Redox Metabolism Shared Resource Facilities of the University of Kentucky Markey Cancer Center [P30CA177558] and University of Kentucky Center for Cancer and Metabolism [P20GM121327].

The original article [1] has been corrected.

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Reference

1. Zhang et al. *BMC Bioinformatics* (2023) Bayesian kinetic modeling for tracer-based metabolomic data (2023) 24:108 DOI: <https://doi.org/10.1186/s12859-023-05211-5>

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