

CORRECTION

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# Correction: Automatic disease prediction from human gut metagenomic data using boosting GraphSAGE

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The original article can be found online at <https://doi.org/10.1186/s12859-023-05251-x>.

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**Correction: BMC Bioinformatics (2023) 24:126**

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Following the publication of the original article [1], the authors would like to correct affiliation 2.

The incorrect affiliation is: Nuclera Nucleics Ltd, Cambridge, UK

The correct affiliation is: Department of Biotechnology, Birla Institute of Technology and Science Pilani Dubai Campus, Dubai International Academic City, Dubai, UAE.

The original article [1] has been corrected.

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## Reference

1. Syama K, et al. Automatic disease prediction from human gut metagenomic data using boosting GraphSAGE. *BMC Bioinform.* 2023;24:126. <https://doi.org/10.1186/s12859-023-05251-x>.

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