# CORRECTION

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# **Correction: Biochar amendments improve** soil functionalities, microbial community and reduce Pokkah boeng disease of sugarcane

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Following the publication of the original article [1], the authors reported a typo in the legend of Fig. 4. It reads: LEfSe analysis of the sugarcane rhizosphere samples among different treatments on phylum level (A bacterial, **B** fungal); LEfSe analysis of the sugarcane rhizosphere samples among different treatments on genus level (A bacterial, **B** fungal).

The correct version should have read: Fig. 4 LEfSe analysis of the sugarcane rhizosphere samples among different treatments on phylum level (A bacterial, B fungal); LEfSe analysis of the sugarcane rhizosphere samples among different treatments on genus level (C bacterial, D fungal).

Furthermore, the author updated Fig. 1 to enhance clarity for the reader. The original Fig. 1 is as follows:

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The original article can be found online at https://doi.org/10.1186/s40538-024-00546-4.

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### The updated Fig. 1 is as follows:



**Fig. 1** Alpha diversity indices: Shannon, Chao1 and coverage of soil bacterial communities from different treatments (**A**); Shannon, Chao1 and coverage of fungal communities from different treatment (**B**). \* $P < 0.05 **P \le 0.01$ . V = Variety, B = Biochar,  $V \times B = \text{Variety} \times \text{Biochar}$ 

The original article [1] has been updated.

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#### Reference

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