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:

†Shakeel Ahmad and Xuexin Zhai have contributed equally to this work.

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The updated Fig. 1 is as follows:

![Fig. 1](image)

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 ≤ 0.01. V = Variety, B = Biochar, V x B = Variety x Biochar

The original article [1] has been updated.

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