

## CORRECTION

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## Correction: The chemopreventive effects of Huangqin-tea against AOM-induced preneoplastic colonic aberrant crypt foci in rats and omics analysis

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Correction for 'The chemopreventive effects of Huangqin-tea against AOM-induced preneoplastic colonic aberrant crypt foci in rats and omics analysis' by Jie Shen *et al.*, *Food Funct.*, 2020, **11**, 9634–9650, DOI: 10.1039/D0FO01731K.

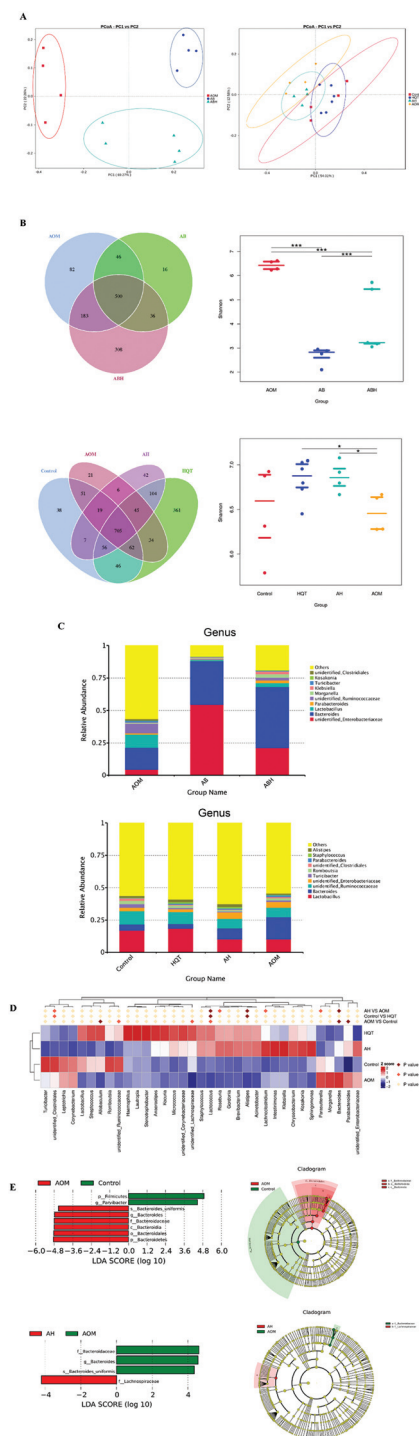
The authors regret that the panels for Fig. 3C–E are missing in the original article. The correct version of Fig. 3 is as shown below.

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**Fig. 3** HQT modulated the gut microbiota community in rats with AOM-induced precancerous colorectal lesions. (A) Principal co-ordinates analysis. (B) Venn diagram showing shared and exclusive bacterial OTUs in the various treatment groups, and alpha diversity was presented by a box plot of the Shannon. (C) The top 10 relative abundances of bacteria at the genus level in different groups. (D) Heatmap: MetaStat analysis of differences in the gut microbiota at the genus level among the groups (control, HQT, AOM, and AH groups). (E) LefSe analysis histogram (left) and biomarker taxa (right).

Consequently, a section of the text in the manuscript should be adjusted as detailed below.

The sentence on page 9640 beginning “In order to find biomarkers between the AOM and AH...” should be correctly given as “In order to find biomarkers between the AOM and AH groups using relative abundances, the LefSe method was used to identify the gut microbiota that was different in abundance in the control, model, and treatment groups (Fig. 3E and Fig. S5†).”

The Royal Society of Chemistry apologises for these errors and any consequent inconvenience to authors and readers.

