



Corrigendum: Metagenome Analysis of Intestinal Bacteria in Healthy People, Patients With Inflammatory Bowel Disease and Colorectal Cancer

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A Corrigendum on

Metagenome Analysis of Intestinal Bacteria in Healthy People, Patients With Inflammatory Bowel Disease and Colorectal Cancer

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FIGURE/TABLE LEGEND

In the original article, there was a mistake in the legend for **Figure 5** as published. There is a duplicate statement. The correct legend appears below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

CORRECT LEGEND

Figure 5 Stratified analyses of metabolic pathways by bacterial species. (A) The circle shows the pathways in which 16 differential bacteria species are involved. The black font represents the metabolic pathways. Red represents the bacterium that belongs to the phylum Bacteroidetes, green represents the bacterium that belongs to the phylum Firmicutes, brown represents the bacterium that belongs to the phylum Actinobacteria. (B) Log abundance for 10 major bacteria species

compared between Healthy, IBD, and CRC cases. P values were determined by Kruskal–Wallis tests. The midline is the median, and the red diamond shape represents the mean values.

INCORRECT AUTHOR NAME

An author name was incorrectly spelled as Jianghou Xiang. The correct spelling is Houxiang Jiang.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

An author name was incorrectly spelled as Zhongmin Yang. The correct spelling is Zhongming Yang.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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