Microbiome

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Correction to: Metagenomic analysis of isolation methods of a targeted microbe, *Campylobacter jejuni,* from chicken feces with high microbial contamination

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Following publication of the original article [1], the authors reported an error in Fig. 2. The correct figure is shown below.

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1. Kim J, Guk J-H, Mun S-H, An J-U, Song H, Kim J, Ryu S, Jeon B, Cho S. Metagenomic analysis of isolation methods of a targeted microbe, *Campylobacter jejuni,* from chicken feces with high microbial contamination. Microbiome. 2019;7:67.

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Fig. 2 Proportion of competing microbes according to isolation procedures of *C. jejuni* based on culture-dependent tools. Proportion of **a** ESBL-producing *E. coli* and **b** *P. mirabilis* among competing microbes according to the isolation procedure of *C. jejuni* (mean ± SEM). Significance was determined by *t* test and one-way ANOVA. The proportion of ESBL-producing *E. coli* was significantly different according to the isolation procedure including the type of enrichment broth, selective media, and the combination of different enrichment broths, the ratio of sample-to-enrichment broth, and selective media. The proportion of *P. mirabilis* was significantly different according to the isolation procedure including the type of selective media and combination of different enrichment broths, the ratio of sample to enrichment broth, and selective agars. *$p < 0.05$, **$p < 0.01$