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Corrigendum: *Edwardsiella piscicida* infection reshapes the intestinal microbiome and metabolome of big-belly seahorses: mechanistic insights of synergistic actions of virulence factors

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Edwardsiella piscicida, metagenome, metabolome, virulence factor, big-belly seahorse, pathogenesis

A Corrigendum on

Edwardsiella piscicida infection reshapes the intestinal microbiome and metabolome of big-belly seahorses: mechanistic insights of synergistic actions of virulence factors

by Zhang L, Wang F, Jia L, Yan H, Gao L, Tian Y, Su X, Zhang X, Lv C, Ma Z, Xue Y, Lin Q and Wang K (2023) *Front. Immunol.* 14:1135588. doi: 10.3389/fimmu.2023.1135588

In the published article, there was an error in Figure 2A as published. E9 1-4 and E21 1-4 should be changed to E9D 1-4 and E21D 1-4, respectively. The corrected Figure 2 and its caption appear below.

Figure 2. Effects of *Edwardsiella piscicida* infection on kingdom-level composition **(A)**, structure **(B)**, and diversity **(C)** of intestinal microbiota in big-belly seahorses. (B, C) represent bacterial intestinal microbiota at the species level. Con represents healthy controls (4.1–4.5 g, PSS); E9D and E21D represent the samples collected on days 9 and 21 of *E. piscicida*- challenged group (4.1–4.5 g, 1×10^5 cfu/mL), respectively (similarly hereinafter). ** $P < 0.01$.

In the published article, there was an error. A repeated and wrong sentence was found in the last paragraph of the **Results** section.

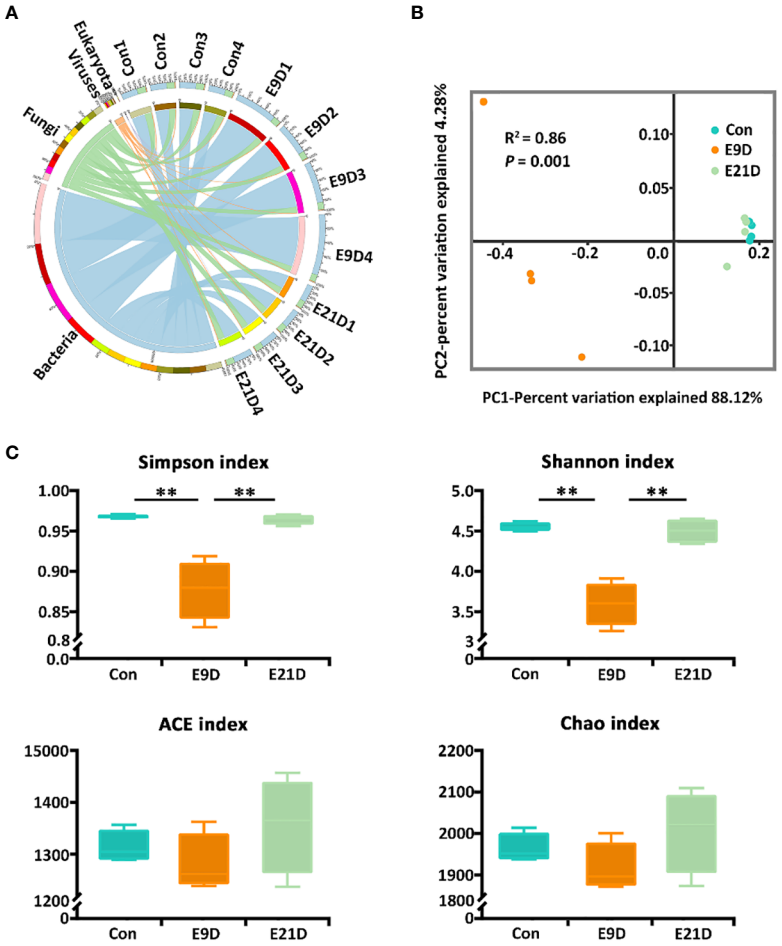

FIGURE 2
Effects of *Edwardsiella piscicida* infection on kingdom-level composition **(A)**, structure **(B)**, and diversity **(C)** of intestinal microbiota in big-belly seahorses. **(B, C)** represent bacterial intestinal microbiota at the species level. Con represents healthy controls (4.1–4.5 g, PSS); E9D and E21D represent the samples collected on days 9 and 21 of *E. piscicida*- challenged group (4.1–4.5 g, 1×10^5 cfu/mL), respectively (similarly hereinafter). $^{**}P < 0.01$.

A correction has been made to **3 Results**, *3.6 Molecular pathogenesis of E. piscicida-induced enteritis*. This sentence previously stated: "In addition, seven of the eight KMBs L-Malic acid and L-Glutamate significantly decreased ($P < 0.05$) (Figures 7A; 4D left panel; 5C). In addition, seven of the eight KMBs with central roles were enriched and significantly negatively correlated with two key intestinal microbiota functions, TCS and ABC transporters ($P < 0.05$)."

The corrected sentence appears below:

"In addition, seven of the eight KMBs with central roles were enriched and significantly negatively correlated with two key intestinal microbiota functions, TCS and ABC transporters ($P < 0.05$)."

The authors apologize for these errors and state that they do not change the scientific conclusions of the article in any way. The original article has been updated.

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