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*CORRESPONDENCE Jie Yu ⊠ yujie1@dongeejiao.com

[†]These authors have contributed equally to this work and share first authorship

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Corrigendum: Metabolic changes before and after weaning in Dezhou donkey foals in relation to gut microbiota

Qiwen Yang^{1,2†}, Haibing Liu^{1†}, Halima Jafari², Bing Liu¹, Zhaofei Wang², Jiangtian Su², Fuwen Wang², Ge Yang², Minhao Sun², Jie Cheng¹, Boying Dong¹, Min Li¹, Mingjian Gen¹ and Jie Yu^{1*}

¹National Engineering Research Center for Gelatin-Based Traditional Chinese Medicine, Dong-E-E-Jiao Co. Ltd., Dong'e County, Shandong, China, ²Key Laboratory of Animal Genetics, Breeding and Reproduction of Shaanxi Province, College of Animal Science and Technology, Northwest A&F University, Xianyang, Shaanxi, China

KEYWORDS

donkey foal, weaning, gut microbes, serum, metabolome

A Corrigendum on

Metabolic changes before and after weaning in Dezhou donkey foals in relation to gut microbiota

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In the published article, there was an error in Figure 8 as published. Figure 8C and Figure 8B were erroneously duplicated. The corrected Figure 8 and its caption appear below:

In the published article, there was an error in section 3 "**Results**", subsection 3.2 " β *diversity of donkey manure*", paragraph 2. This sentence previously stated: "The other three groups M.F.1, M.F.3, and M.F.6 contain 2,402,1,671 and 7,519 ASVs, respectively."

The corrected sentence appears below:

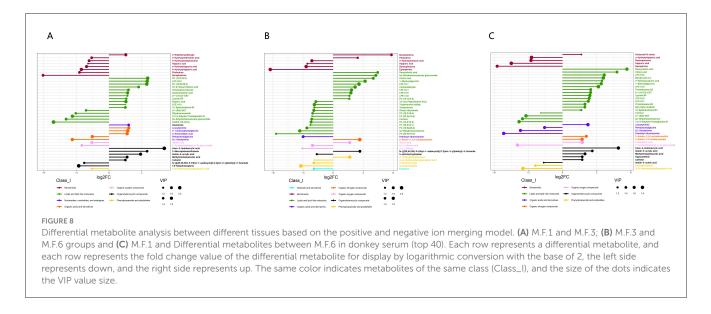
"The other three groups M.F.1, M.F.3, and M.F.6 contain 2,402,1,671 and 7,529 ASVs, respectively."

In the published article, there was an error in section 3 "**Results**", subsection 3.5 "*Identification and evaluation of differential metabolites*", paragraph 3. This sentence previously stated:

"The top 10 differential metabolites were norephedrine (log2FC = 5.82), deoxycholic acid (log2FC = 4.45), 5 α -dihydrotestosterone glucuronide (log2FC = 4.26), cholic acid (log2FC = 4.00), 1-stearoylglycerol (log2FC = 3.20), N-{((2R,4S,5R)-5-ethyl-1-azabicyclo [2.2.2]oct-2-yl]methyl}-2-furamide (aldehyde (log2FC = 2.68), LPA 16:1 (log2FC = 2.58), pholedrine (log2FC = 2.58) and cuminaldehyde (log2FC = 2.20)."

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The authors apologize for these errors 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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