

CORRECTION

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# Correction to: Aging progression of human gut microbiota



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**Correction to: BMC Microbiol 19, 236 (2019)**  
<https://doi.org/10.1186/s12866-019-1616-2>

Following the publication of the original article [1], we were notified that the supplementary material contains the incorrect links.

The corrected Additional files can be found below:

## Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s12866-021-02200-7>.

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| <p><b>Additional file 1: Supplementary Table 1.</b> Significant genera from PERMANOVA analysis</p> <p><b>Additional file 2: Supplementary Table 2.</b> Genera correlated with aging with Spearman correlation</p> <p><b>Additional file 3: Supplementary Table 3.</b> Critical genera revealed by SPD</p> <p><b>Additional file 4: Supplementary Figure 1.</b> The relative abundance of all the 35 critical genera across different age groups.</p> |
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## Reference

1. Xu, et al. Aging progression of human gut microbiota. *BMC Microbiol.* 2019; 19:236. <https://doi.org/10.1186/s12866-019-1616-2>.

The original article can be found online at <https://doi.org/10.1186/s12866-019-1616-2>.

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