



<https://doi.org/10.1038/s42003-022-04266-z>

OPEN

Author Correction: Metagenome-assembled genomes and gene catalog from the chicken gut microbiome aid in deciphering antibiotic resistomes

Yuqing Feng , Yanan Wang , Baoli Zhu, George Fu Gao , Yuming Guo & Yongfei Hu 

Correction to: *Communications Biology* <https://doi.org/10.1038/s42003-021-02827-2>, published online 18 November 2021.

In the original version of this article, several references were incorrectly numbered in the text. These have now been corrected as follows:

On Page 2, 10 changes have been made to correctly cite references 7, 11, 12, 13, 14, 15, 16, 17 and 18–21.

On Page 3, 2 changes have made to correctly cite references 7, 18 and 19.

On Page 4, 1 change has been made to correctly cite reference 7.

On Page 6, 3 changes have been made to correctly cite reference 7, 11 and 13.

In addition, 4 references were listed out of order in the references list. These have now been corrected to:

8. Broom, L. J. & Kogut, M. H. The role of the gut microbiome in shaping the immune system of chickens. *Vet. Immunol. Immunopathol.* **204**, 44–51 (2018).

9. Gordon, J. I., Raoult, D. & Henrissat, B. The abundance and variety of carbohydrate-active enzymes in the human gut microbiota. *Nat. Rev. Microbiol.* **11**, 497–504 (2013).

10. Segura-Wang, M., Grabner, N., Koestelbauer, A., Klose, V. & Ghanbari, M. Genome-resolved metagenomics of the chicken gut microbiome. *Front. Microbiol.* **12**, 726923 (2021).

11. Stewart, R. D. et al. Compendium of 4941 rumen metagenome-assembled genomes for rumen microbiome biology and enzyme discovery. *Nat. Biotechnol.* **37**, 953–961 (2019).

This has now been corrected in the PDF and HTML versions of the Article.

Published online: 24 November 2022



Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons license, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons license and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this license, visit <http://creativecommons.org/licenses/by/4.0/>.

© The Author(s) 2022