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

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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.

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