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AUTHOR CORRECTION

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Author Correction: Assembly of hundreds of novel bacterial genomes from the chicken caecum

Laura Glendinning^{1*}, Robert D. Stewart¹, Mark J. Pallen^{2,3,4}, Kellie A. Watson¹ and Mick Watson¹

The original article can be found online at <https://doi.org/10.1186/s13059-020-1947-1>.

* Correspondence: laura.glendinning@roslin.ed.ac.uk

¹Genetics and Genomics, The Roslin Institute and Royal (Dick) School of Veterinary Studies, University of Edinburgh, Edinburgh, Midlothian, UK

Full list of author information is available at the end of the article

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Following publication of the original paper [1], the authors have reported several typographical and linguistic errors in the new *Candidatus* names presented in Additional file 5: Dataset S4. In addition, the presentation of the names without full protologues has precluded their registration in the *International Journal of Systematic and Evolutionary Microbiology*, in line with the recommendations of the *International Code of Nomenclature of Prokaryotes*. The original Additional File 5 has therefore been replaced by Additional File 5A, which presents Dataset S4 with corrected names, and by Additional File 5B, which contains protologues for the new *Candidatus* names.

Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s13059-021-02284-4>.

Additional file 5 A: Clustering of samples at 60% AAI to form genus clusters. Novel genera were defined as clusters of MAGs at 60% AAI which were not assigned a genus by GTDB-Tk. B: Protologues for the new *Candidatus* names.

Author details

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¹Genetics and Genomics, The Roslin Institute and Royal (Dick) School of Veterinary Studies, University of Edinburgh, Edinburgh, Midlothian, UK. ²Microbes in the Food Chain, Quadram Institute Bioscience, Norwich, UK. ³School of Biological Sciences, University of East Anglia, Norwich, Norfolk, UK. ⁴School of Veterinary Medicine, University of Surrey, Guildford, Surrey, UK.

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Reference

1. Glendinning L, Stewart RD, Pallen MJ, et al. Assembly of hundreds of novel bacterial genomes from the chicken caecum. *Genome Biol.* 2020;21:34 <https://doi.org/10.1186/s13059-020-1947-1>.



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