

CORRIGENDUM

Correction to: High-coverage genomes to elucidate the evolution of penguins

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In the original version of the article “High-coverage genomes to elucidate the evolution of penguins” by Hailin Pan *et al.* [1], the authors received a request to make some clarifications and changes in the acknowledgements, the following is the modified version:

“We thank the following: John Cockrem, Scott Flemming, Helen McConnell, Chris Rickard, Sarah Fraser, Otto Whitehead, Kyle Morrison, and Amy Van Buren for help collecting samples; Jonathan Banks, Kirsten Rodgers, and Jo Hiscock for sample information; Manuel Paredes Oyarzún and Hernán Rivera Meléndez for facilitating permits and sample collection; Lauren Tworowski, Richard O’Rorke, and Joanna Sumner for facilitating sample collection; Adrian Smith for providing laboratory support to extract 2 DNA samples; Peter Dearden, Neil Fowke,

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References

1. Pan H, Cole T L, Bi X *et al.* High-coverage genomes to elucidate the evolution of penguins. *GigaScience* 2019;8(9):1–17.