

CORRECTION

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Correction: Human impact on the recent population history of the elusive European wildcat inferred from whole genome data

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Correction: *BMC Genomics* 23, 709 (2022)
<https://doi.org/10.1186/s12864-022-08930-w>

Following publication of the original article [1], the authors reported an error in the "Availability of data and materials" section, as they did not provide the European Nucleotide Archive (ENA) project number of the newly sequenced samples.

Therefore, the section should read as follows:

Availability of data and materials

Newly generated whole genome individual sequencing data is available at European Nucleotide Archive (ENA) project number: PRJEB40421 (<https://www.ebi.ac.uk/ena/browser/view/PRJEB40421>). fastq files for 21

domestic cat individuals can be found under ENA project PRJNA343389 (<https://www.ebi.ac.uk/ena/browser/view/PRJNA343389>).

The original article [1] has been updated.

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Reference

1. Nieto-Blázquez ME, et al. Human impact on the recent population history of the elusive European wildcat inferred from whole genome data. *BMC Genomics*. 2022;23:709. <https://doi.org/10.1186/s12864-022-08930-w>.

The original article can be found online at <https://doi.org/10.1186/s12864-022-08930-w>.

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