

CORRECTION

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Correction to: Archaic mitochondrial DNA inserts in modern day nuclear genomes



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Correction to: BMC Genomics

<https://doi.org/s12864-019-6392-8>

Following the publication of this article [1], the authors reported that the captions of Figs. 3 and 4 were published in the incorrect order, whereby they mismatch with their corresponding images. The figures are reproduced in the correct sequence with the correct captions in this Correction article. The original article has been corrected.

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Published online: 17 January 2020

Reference

1. Bücking, et al. Archaic mitochondrial DNA inserts in modern day nuclear genomes. *BMC Genomics*. 2019;20:1017 <https://doi.org/s12864-019-6392-8>.

The original article can be found online at <https://doi.org/10.1186/s12864-019-6392-8>

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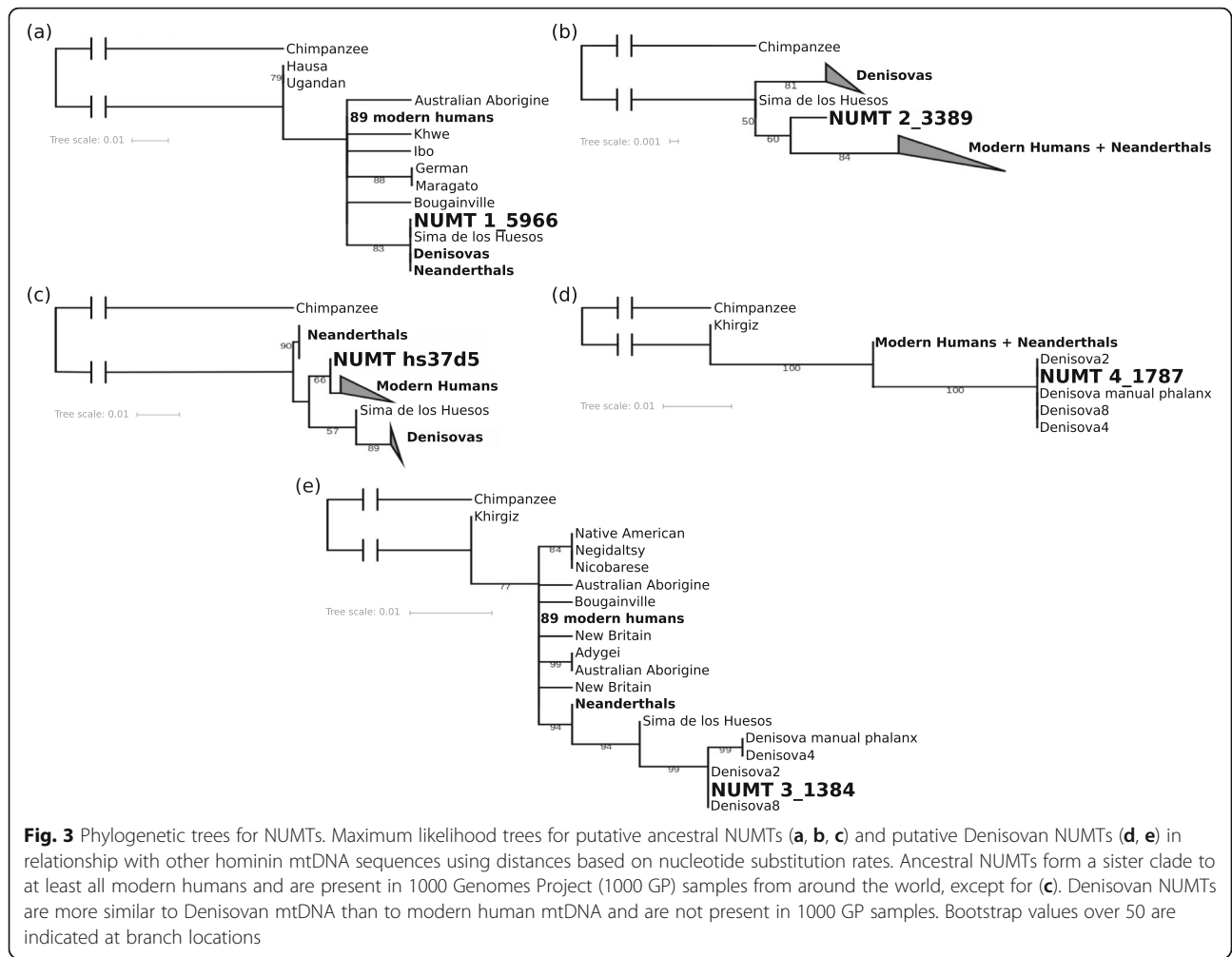


Fig. 3 Phylogenetic trees for NUMTs. Maximum likelihood trees for putative ancestral NUMTs (**a, b, c**) and putative Denisoan NUMTs (**d, e**) in relationship with other hominin mtDNA sequences using distances based on nucleotide substitution rates. Ancestral NUMTs form a sister clade to at least all modern humans and are present in 1000 Genomes Project (1000 GP) samples from around the world, except for (**c**). Denisoan NUMTs are more similar to Denisoan mtDNA than to modern human mtDNA and are not present in 1000 GP samples. Bootstrap values over 50 are indicated at branch locations

