



Corrigendum: Frequent somatic transfer of mitochondrial DNA into the nuclear genome of human cancer cells

Young Seok Ju, Jose M.C. Tubio, William Mifsud, et al.

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Corrigenda

Genome Research 26: 291–300 (2016)

Corrigendum: Model-based analyses of whole-genome data reveal a complex evolutionary history involving archaic introgression in Central African Pygmies

PingHsun Hsieh, August E. Woerner, Jeffrey D. Wall, Joseph Lachance, Sarah A. Tishkoff, Ryan N. Gutenkunst, and Michael F. Hammer

The authors would like to correct the omission of the term “model-based” in the Abstract. The corrected sentence should read as follows:

“our inference method rejects the hypothesis that the ancestors of AMH were genetically isolated in Africa, thus providing model-based whole genome-level evidence of African archaic admixture.”

In addition, they also would like to make a change in the last sentence in the Introduction, and the corrected sentence should read as follows:

“Together, our results provide a model-based whole-genome perspective on archaic introgression in Africa.”

The article has already been corrected in both the PDF and full-text HTML files online.

doi: 10.1101/gr.206524.116

Genome Research 25: 814–824 (2015)

Corrigendum: Frequent somatic transfer of mitochondrial DNA into the nuclear genome of human cancer cells

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The names of two co-authors, Ola Myklebost and G. Steven Bova, were omitted from the authorship list of this article. Please note the corrected list above. The author and affiliation lists have already been corrected in both the PDF and full-text HTML files online.

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