

Author Correction: Human origins in a southern African palaeo-wetland and first migrations

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 Check for updates

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It has come to our attention that 14 (out of 198) of the mitogenomes identified in this Article with relatively poorer sequence quality resulted in artificial recombinants. This has been corrected by modifying the set of parameterizations that was used in the original Article to deduce the consensus sequences. Each of the corrected sequences differs from the original sequence by 9 to 14 nucleotides, but the haplogroup assignments or estimates of divergence time were not affected. The Supplementary Information to this Amendment contains the minor changes made to the parameters used for re-deducing consensus mitochondrial DNA (mtDNA) from ion torrent short-read sequencing data, including Table S1 showing the corrected sequences and lack of change in haplogroup assignment, and Table S2 showing the repeat divergence time estimates using the corrected mtDNA sequences. Only one parallel mutation (A16129G, a common mutation found in many lineages) was found in one sample after correction. Three non-indel mutations remain missing from three samples: a L0d3b-defining mutation (C152T) missing from BM25 (MK248280), a L0d3-defining mutation (A16300G) missing from DIV489 (MK248380), and a L0d2c-defining mutation (A13827G) missing from WB202 (MK248420). Corrected sequences for the 14 mitogenomes (with GenBank accession numbers MK248280, MK248281, MK248285, MK248370, MK248376, MK248377, MK248380, MK248381, MK248382, MK248385, MK248387, MK248393, MK248420, MK248422) have been updated with GenBank. The original Article has not been corrected.

Supplementary Information is available in the online version of this Amendment.