Erratum

This erratum refers to:

Dietmar Zinner, Jenny Wertheimer, Rasmus Liedigk,
Linn Fenna Groeneveld, Christian Roos

Baboon phylogeny as inferred from complete mitochondrial genomes.

American Journal of Physical Anthropology Volume
150, Issue 1, 133–140, published online 26 November
2012 in Wiley Online Library, doi:10.1002/ajpa.22185

Erratum Text

The designation of haplotype ranges and sampling location with capital letters D and F in Figure 1a have been exchanged. Here we present the corrected version of the figure that includes the correct designations for range D of haplogroup D (*P. anubis* West 1) and for range F of haplogroup F (*P. anubis* West 2).

Fig. 1. (a) Distribution of the six baboon species and the seven major mitochondrial haplogroups (A–G) (map based on Kingdon, 1997; Jolly, 2007; Zinner et al., 2009, in press), and (b) simplified phylogenetic relationships among the haplogroups (adapted from Zinner et al., 2009). In (a), dashed lines and colored circles indicate the distribution of the seven major mitochondrial haplogroups and the geographic provenance of samples used in this study (see also Table 1), respectively. In (b), numbers on branches represent support values from Bayesian inference and maximum-likelihood analysis, respectively.

DOI: 10.1002/ajpa.22973

Published online 22 March 2016 in Wiley Online Library
(wileyonlinelibrary.com).