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Different population histories of the Mundari- and Mon-Khmer-speaking Austro-Asiatic tribes inferred from the mtDNA 9-bp deletion/insertion polymorphism in Indian populations

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In the May 2005 issue of the Journal, in the article entitled “Different population histories of the Mundari- and Mon-Khmer-speaking Austro-Asiatic tribes inferred from the mtDNA 9-bp deletion/insertion polymorphism in Indian populations,” by Thangaraj et al. (116:507–517), there were three typographical errors, one each in text, table 2 and figure 3:

1. In the ‘Electronic database information’ heading in the third line on page 515, “for mtDNA HVS-I,

accession numbers AY255133–AY255180” should be “for mtDNA HVS-I, accession numbers AY368704–AY368863”.

2. In Table 2 on page 511, the HVS-I motif of haplotype H3 “223-156-274-320-360” should be “223-256-274-320-360”.
3. The 22 Nicobarese of B5a1 haplogroup in Figure 3 on page 515 should have the additional mutation 13145 along with 11881 and 13395 (fig. 1). Our reanalysis did not exhibit any further discrepancies with our original sequence data. The authors regret these errors.

The original article can be found at <http://dx.doi.org/10.1007/s00439-005-1271-6>

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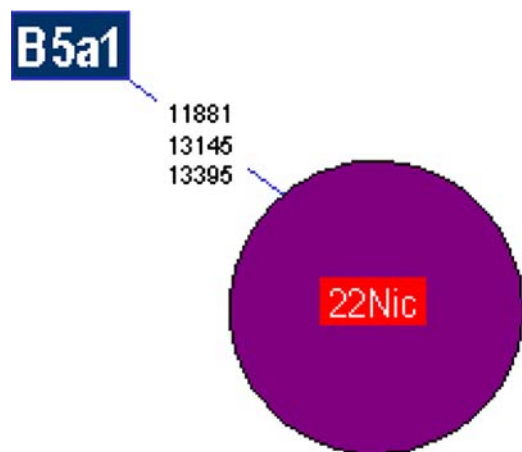


Fig. 1 The phylogenetic position of 22 Nicobarese samples in B5a1 haplogroup with an additional mutation 13145