

Erratum to: Evaluating length heteroplasmy in the human mitochondrial DNA control region

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After the proofs stage, the following typographical errors were introduced in the published version.

"we will quantify the two mutation rates in Table 5 as separate sections 1 and 2" has incorrectly been published as "we will quantify the two mutation rates in Table 5 as separate sections Introduction and Subjects and methods".

The sentence "Assuming in the high-radiation area a mutation rate of 1/600 transmissions for loss of four cytosines, 1/600 for loss of three cytosines, and 4/600 for a loss of two cytosines (Table 5), the probability to observe the child's sequence, given the putative mother is the true mother, is approximately 1/600=0.17%" has been incorrectly published as "Assuming in the high-radiation area a mutation rate of one out of 1/600 transmissions for loss of four cytosines, one out of 1/600 for loss of three cytosines, and four out of 4/600 for a loss of two cytosines (Table 5), the probability to observe the child's sequence, given the putative mother is the true mother, is approximately 1/600=0.17%".

The online version of the original article can be found under <http://dx.doi.org/10.1007/s00414-009-0385-0>.

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