Correction to: Neolithic phylogenetic continuity inferred from complete mitochondrial DNA sequences in a tribal population of Southern India

Charles Sylvester^{1,2} · Mysore Siddaiah Krishna¹ · Jaya Sankar Rao² · Adimoolam Chandrasekar²

Published online: 10 September 2018
© Springer Nature Switzerland AG 2018

Correction to: Genetica

<https://doi.org/10.1007/s10709-018-0030-2>

Unfortunately, the original version of this article was published with an error in the second sentence of the ‘**Molecular dating**’ section: “The first mutation rate based on substitution rate of the entire coding region..... was mentioned as **5.39** years for each mutation” (Mishmar et al. 2003).

However, the correct mutation rate is 5,139 years for each mutation (Mishmar et al. 2003).

Reference

Mishmar D, Ruiz-Pesini E, Golik P, Macaulay V, Clark AG, Hosseini S, Brandon M, Easley K, Chen E, Brown MD, Sukernik RI (2003) Natural selection shaped regional mtDNA variation in humans. *Proc Natl Acad Sci* 100(1):171–176

The original article can be found online at <https://doi.org/10.1007/s10709-018-0030-2>.

✉ Mysore Siddaiah Krishna
drosokrish@gmail.com

¹ Department of Studies in Zoology, University of Mysore, Mysore, India

² Anthropological Survey of India, Southern Regional Center, Mysore, India