

ERRATA

M. A. Bermisheva, I. A. Kutuev, T. Yu. Korshunova, N. A. Dubova, R. VILLEMS, and E. K. Khusnutdinova regret the errors that occurred in their paper “Phylogeographic Analysis of Mitochondrial DNA in the Nogays: A Strong Mixture of Maternal Lineages from Eastern and Western Eurasia” published in *Molecular Biology*, 2004, vol. 38, no. 4, pp. 516–523.

Below are the corrected Fig. 2 and legend to it.

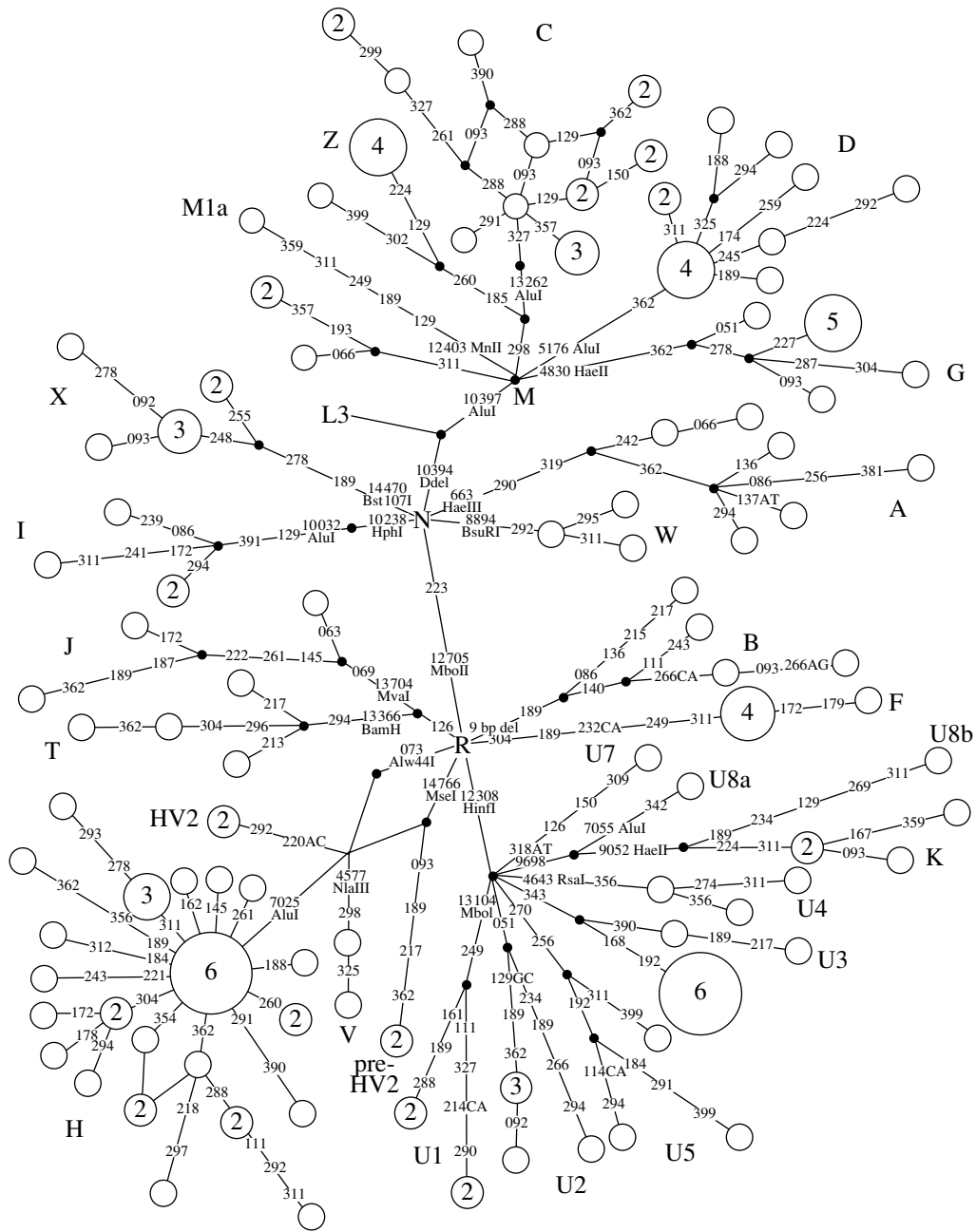


Fig. 2. The phylogenetic tree of mtDNA in the Nogay population. The size of each circle is proportional to the haplotype frequency. The mutations are identified relative to the Cambridge Reference Sequence of the hypervariable segment I; the positions are indicated without 16,000, i.e., 16,223 is written as 223. Mutations are transversions; in the case of transversions, base changes are specified. For some clusters, restriction sites of the mtDNA coding region used for haplogroup identification are indicated.