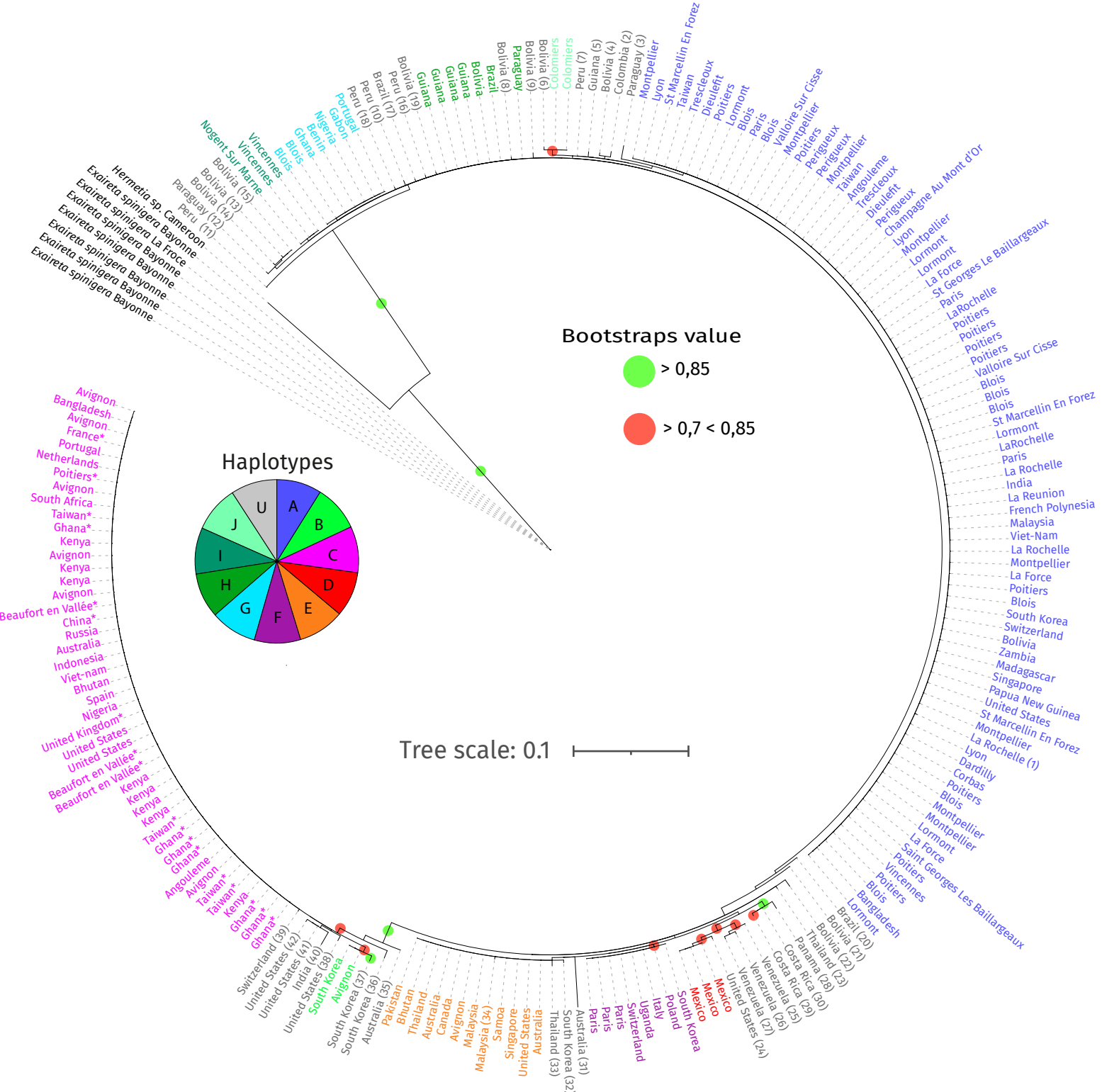

Phylogenetic tree of the CO1 gene.

This tree was made by keeping only one sequence per haplotype except for France.

The visualization is done using ITOL. An asterisk is placed for individuals from breeding.

The statistical support was estimated with 500 Bootstraps with MegaX software using the Tamura 92 G+I model.