

Additional Figure S1. Results of statistical parsimony network analyses based on mtDNA data: Sub-networks recovered under the 95% limit for a parsimony connection. Haplotypes are represented by circles with size proportional to their frequencies, coloured according to the main mtDNA lineages, and labelled according to the sublineages recovered in the phylogenetic analysis (see text for more details). Small white circles represent unsampled or extinct haplotypes.

