

Figure S4. SAMOVA (Spatial Analysis of Molecular Variance) design and results. A: Geographic location of mitochondrial clades estimated as the centroid between member localities (see also Fig.1). B: Localities and sequences included in each mitochondrial clade as defined by previous phylogenetic analyses. C: SAMOVA results showing relative fixation indices F_{CT} and F_{SC} ($P < 0.001$) for pre-defined value of K from 2 to 10 (after $K=10$ structures are not informative as one population at a time is removed from the groups structure).

