



Figure S1. Outputs of the STRUCTURE analyses. On the left side—plots of the ΔK statistic of Evanno et al. (2005) for detecting the number of K groups that best fit the data. On the right side—the plot of mean likelihood $L(K)$ and variance per K value from STRUCTURE. (A) Analyses performed including all the individuals. (B) Analyses performed including the population assigned to the genetic cluster A. (C) Analyses performed on the populations assigned to the genetic cluster B.