

Figure S1 BAYESIAN ANALYSIS OF POPULATION STRUCTURE (BAPS) of 83 mitochondrial sequences (803 bp). The best clustering solution of five (PP = 0.81) is shown.

Cluster 1 MDC 191

- Cluster 2 MDC31, MDC44, MDC66, MDC80, MDC144, MDC146, DC62, DC64, DEC65, DC81, DC82, DC14, DC157
- Cluster 3 MDC2, MDC4, MDC41, MDC63, MDC120, MDC159
- Cluster 4 MDC25, MDC27, MDC49, MDC83, MDC88, MDC99, MDC104, MDC106, MDC131, MDC133, MDC138, MDC138, MDC150, MDC152, MDC155, DC36, DC53, DC54, DC56, DC59, DC60, DC64, DC66, DC68, DC69, DC73, DC74, DC75, DC76, DC77, DC87, DC88, DC89, DC90, DC92, DC93, DC94, DC175, DC176, DC178, DC186, DC188, DC198, DC199, DC201, DC203, DC205
- Cluster 5 MDC 1, MDC 37, MDC 70, MDC 73, DC 70, DC 83, DC 96, DC 98, DC 204, DC 250, DC 251, DC 253, DC 254