



Supplementary Fig. 1. Sampling sites outside the contact region, marked by lineage. This figure provides detail for Fig. 1a. Lineage was determined by mtDNA as well as nuclear loci at some sites. N lineage sites are marked with an N and S lineage sites with an S. Sample size was 10-20 at most sites. The broad scale distribution of the two lineages was determined by Schneider *et al.* (ref. 1).