

Figure S1. Median-joining network inferred from sequences from C lineage (Q), with the C.lineage.Europe sequences correspond to individuals sampled in Europe with no data reported on the sub-species name

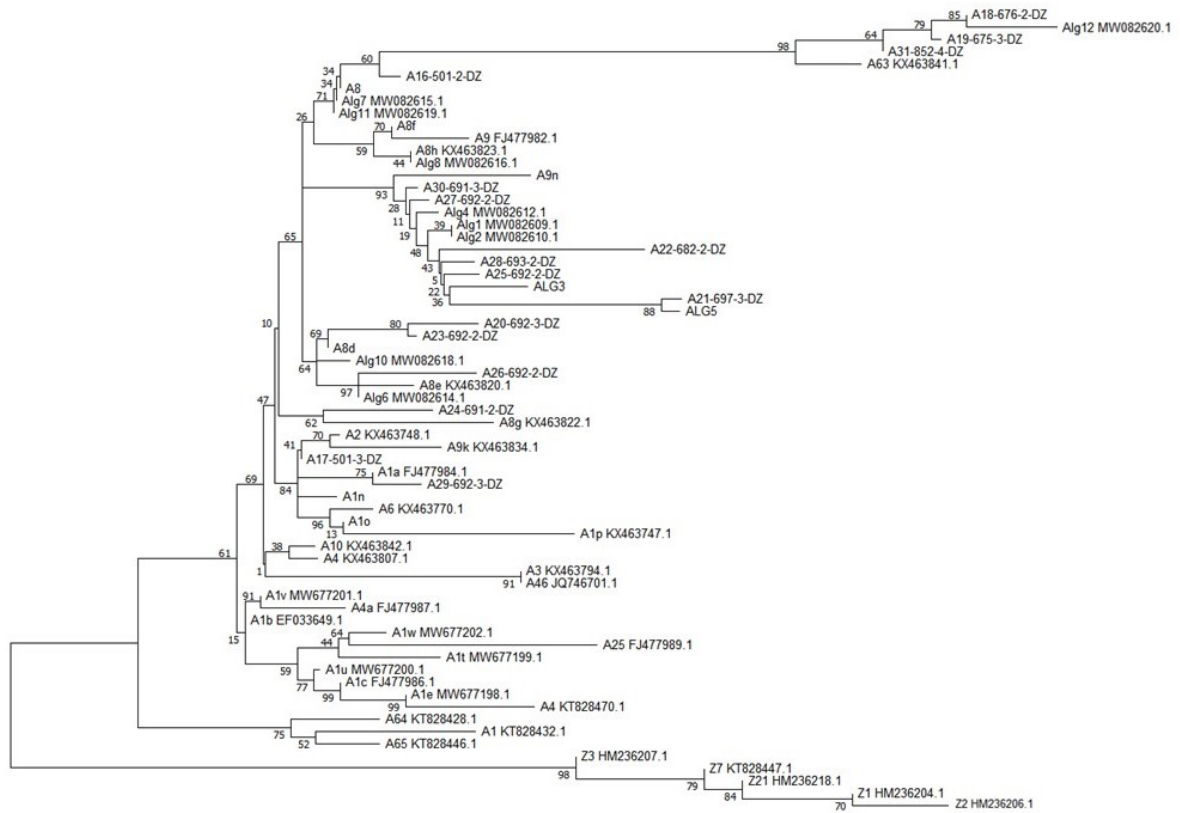


Figure S2. Phylogenetic tree of A lineage haplotypes. The tree is neighbor-joining. Nodes are supported by 1000 bootstraps