

Figure S1

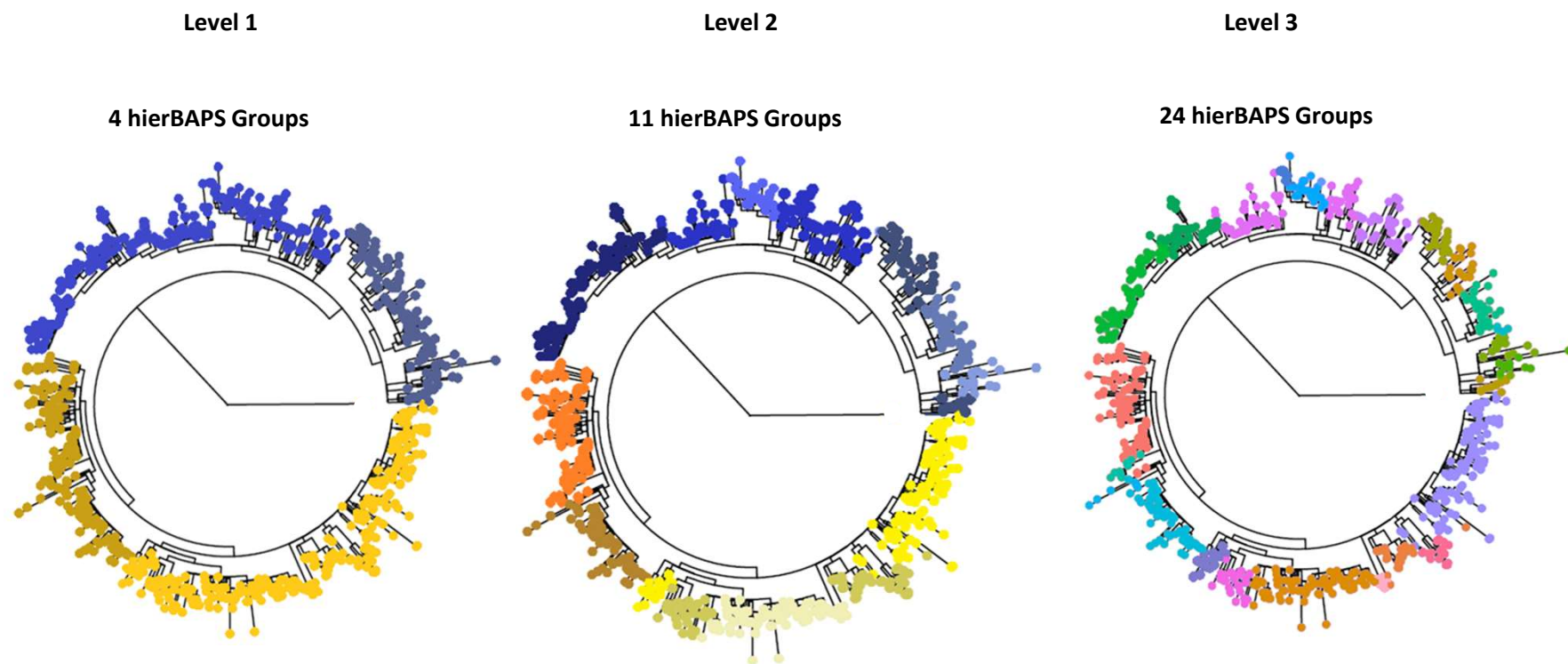


Figure S1. Three analysis levels of hierBAPS groups superimposed onto a maximum likelihood phylogenetic tree.

Figure S2

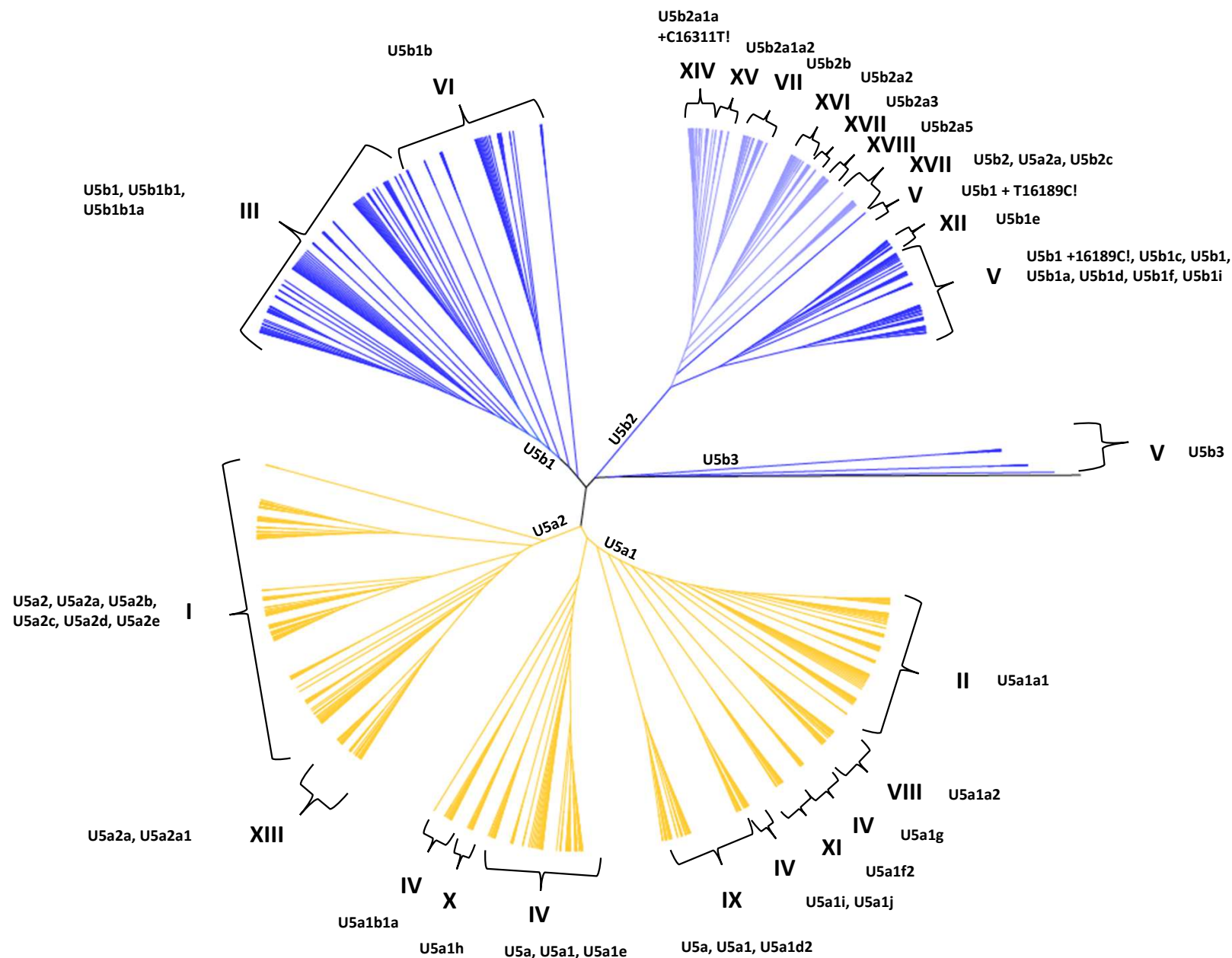


Figure S2. Coding region only analysis of hierBAPS group identification using mtDNA. The hierBAPS groups have been superimposed on a phylogenetic tree, generated using maximum likelihood analysis to view the phylogenetic relationships of each sequence.