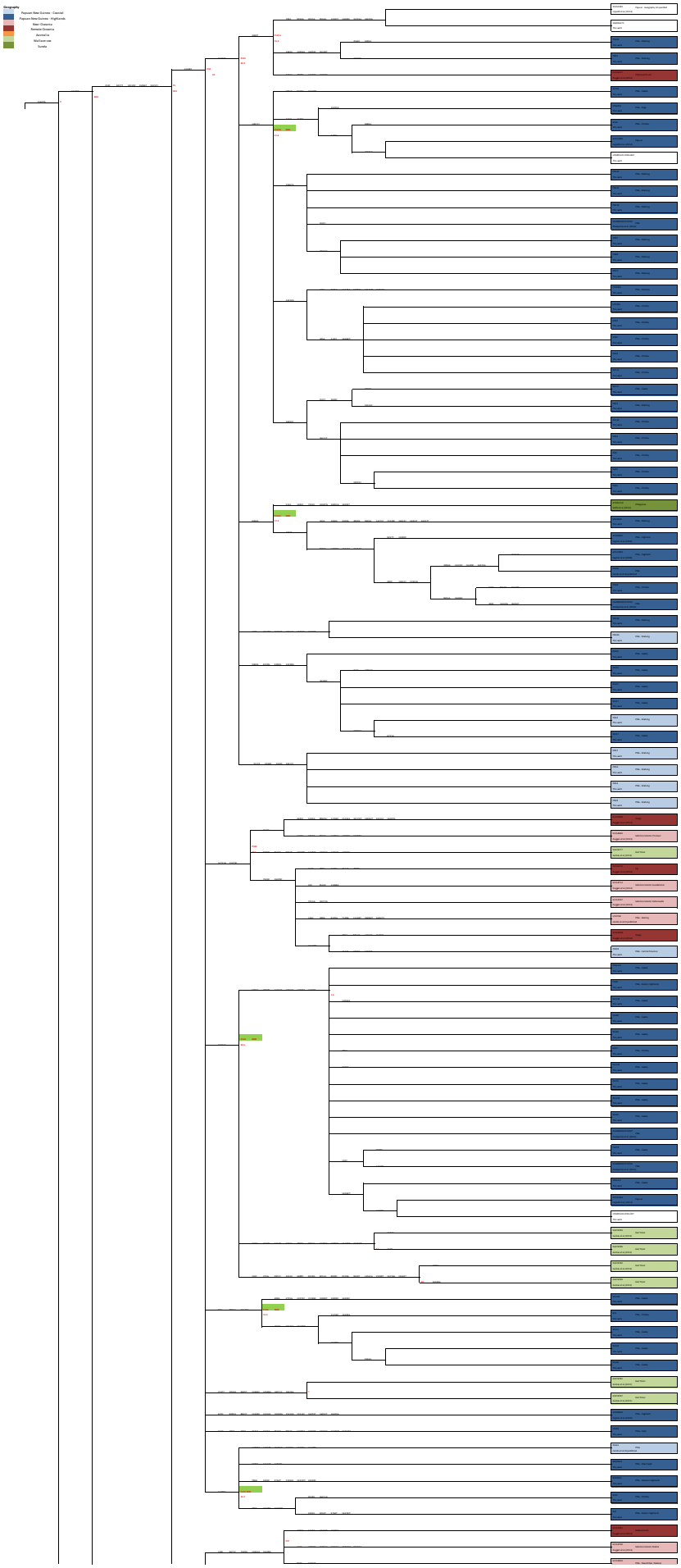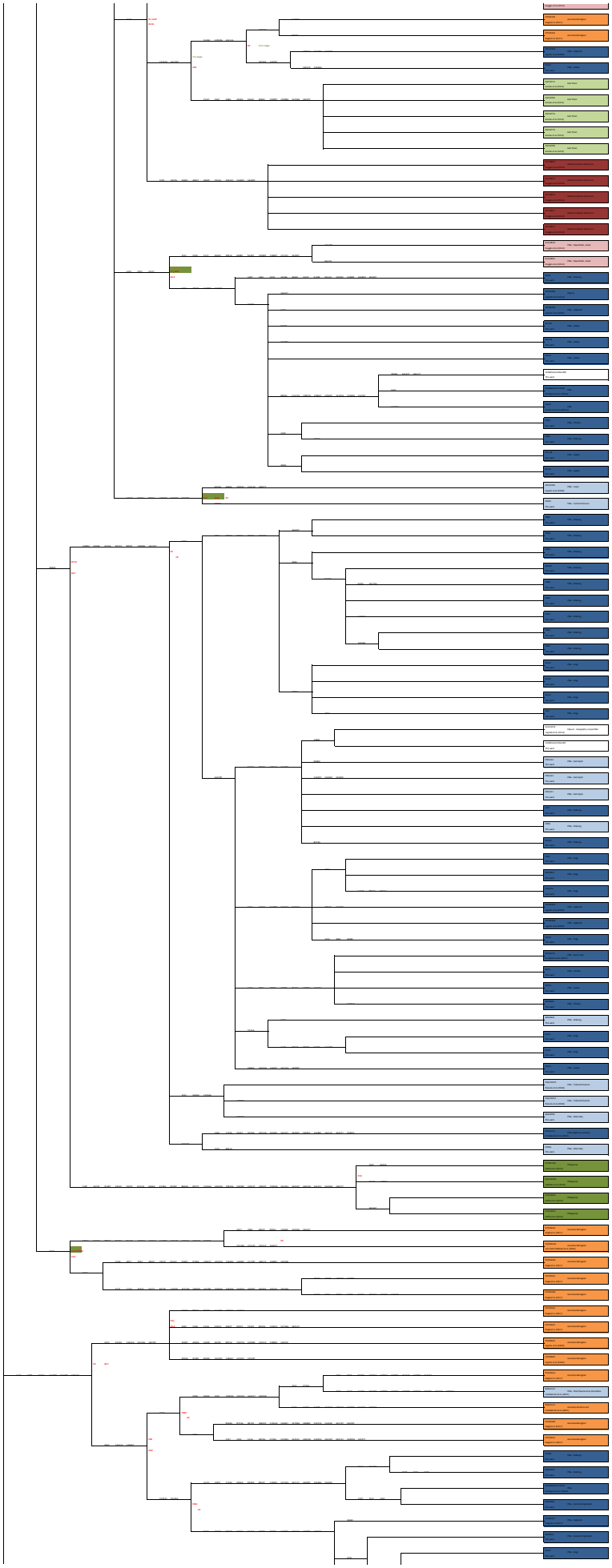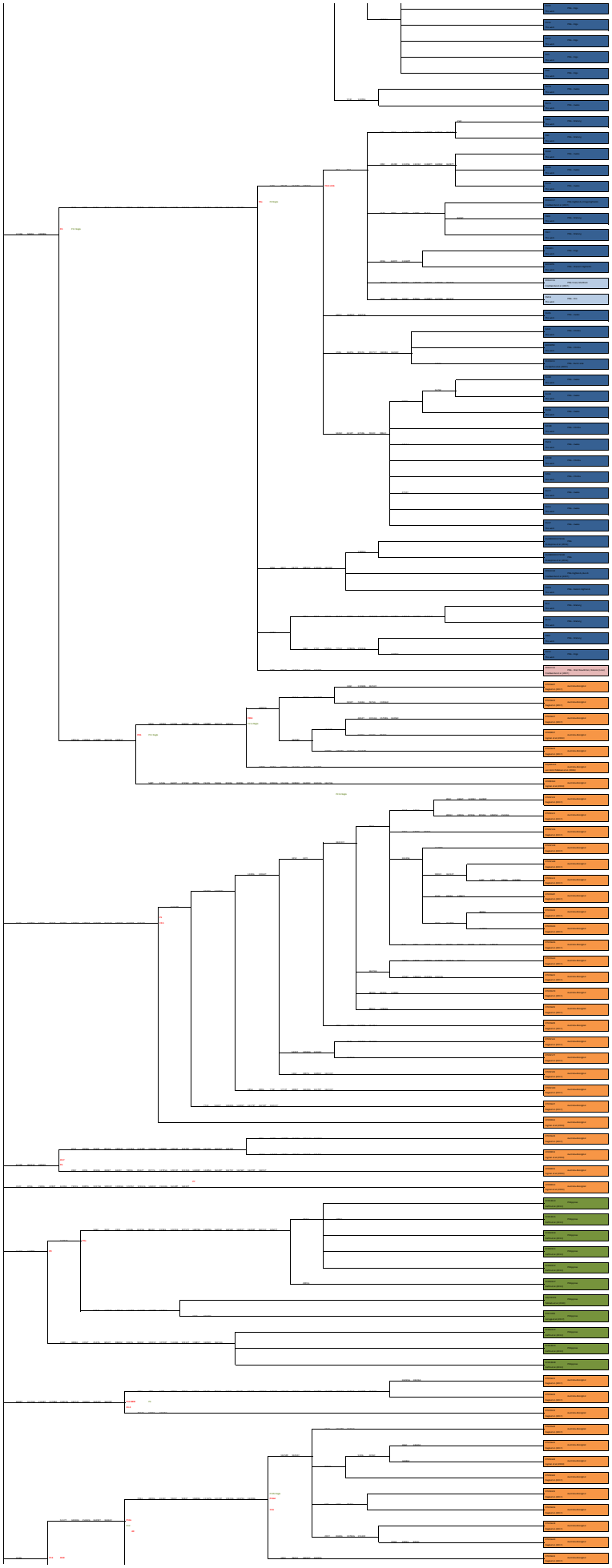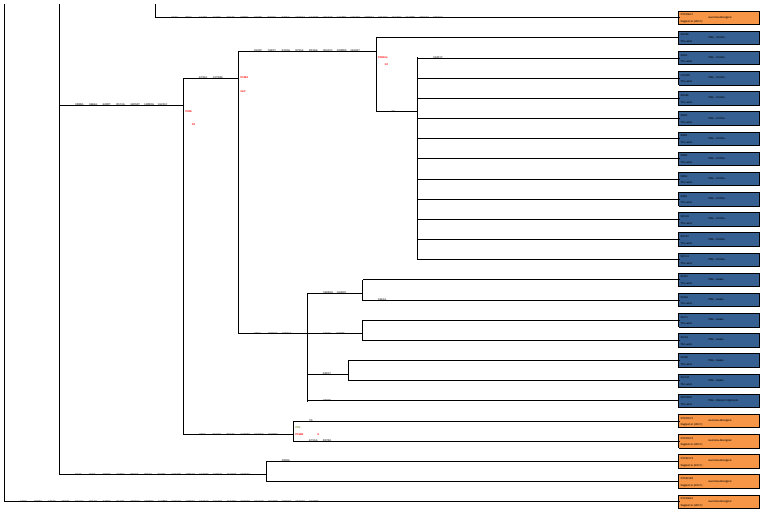Figure S6. Maximum Parsimony Phylogenetic Tree of P. falciparum mtDNA.
Substitutions associated with length variation in the poly-C tracts of the hypervariable regions as well as the ambiable 16189 position were excluded from phylogenetic analysis.