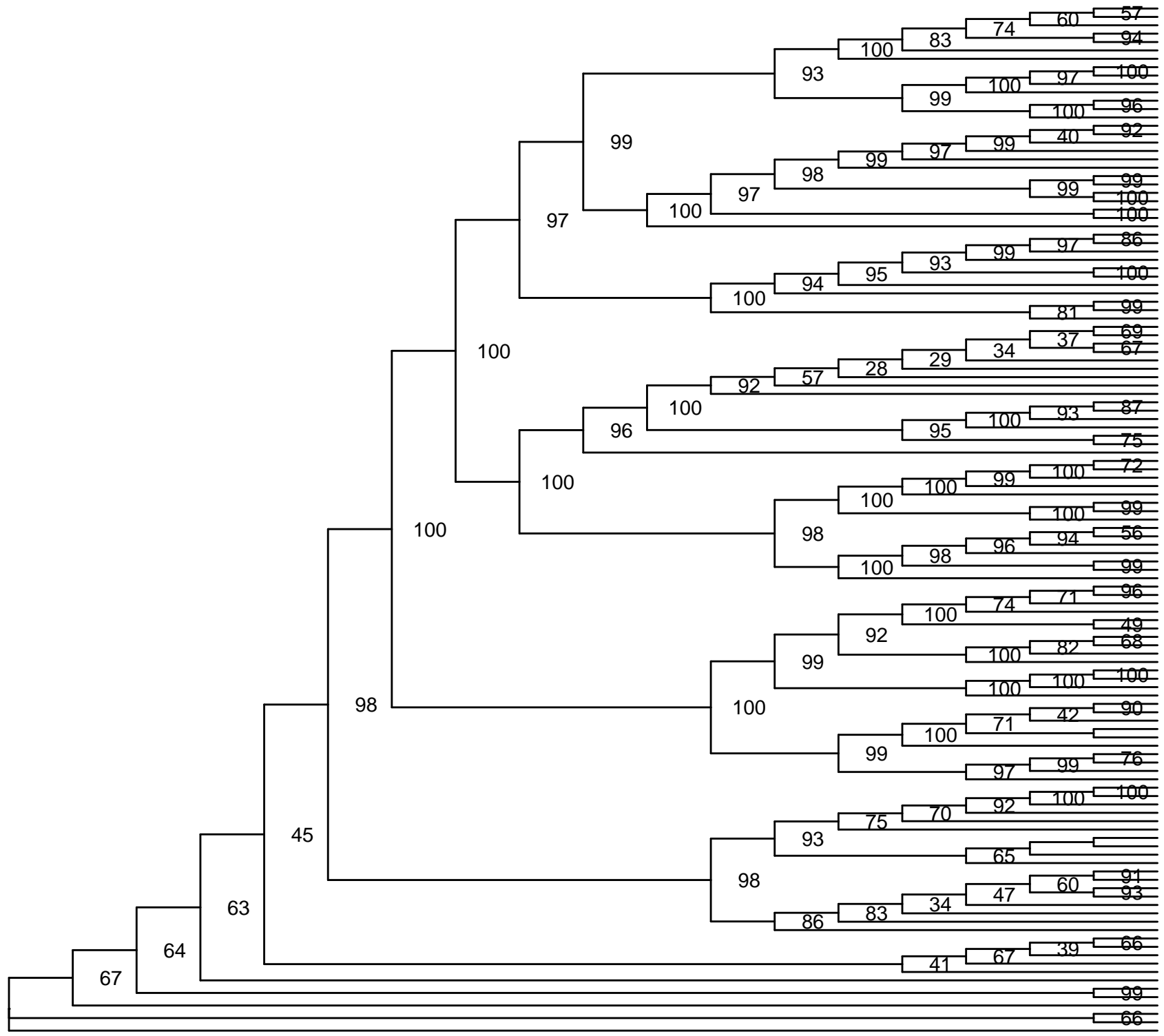


Tree



Sequence alignment data for the taxa shown in the tree. The sequences are represented by a series of vertical bars, each corresponding to a taxon. The bars are color-coded to represent different nucleotide bases: A (Adenine), C (Cytosine), G (Guanine), and T (Thymine). The alignment shows the relative positions of the bases across the different taxa, highlighting conserved regions and areas of divergence. The sequences are labeled with their respective taxon numbers on the left side of the alignment.

MSA

