**Core \_NS5B**

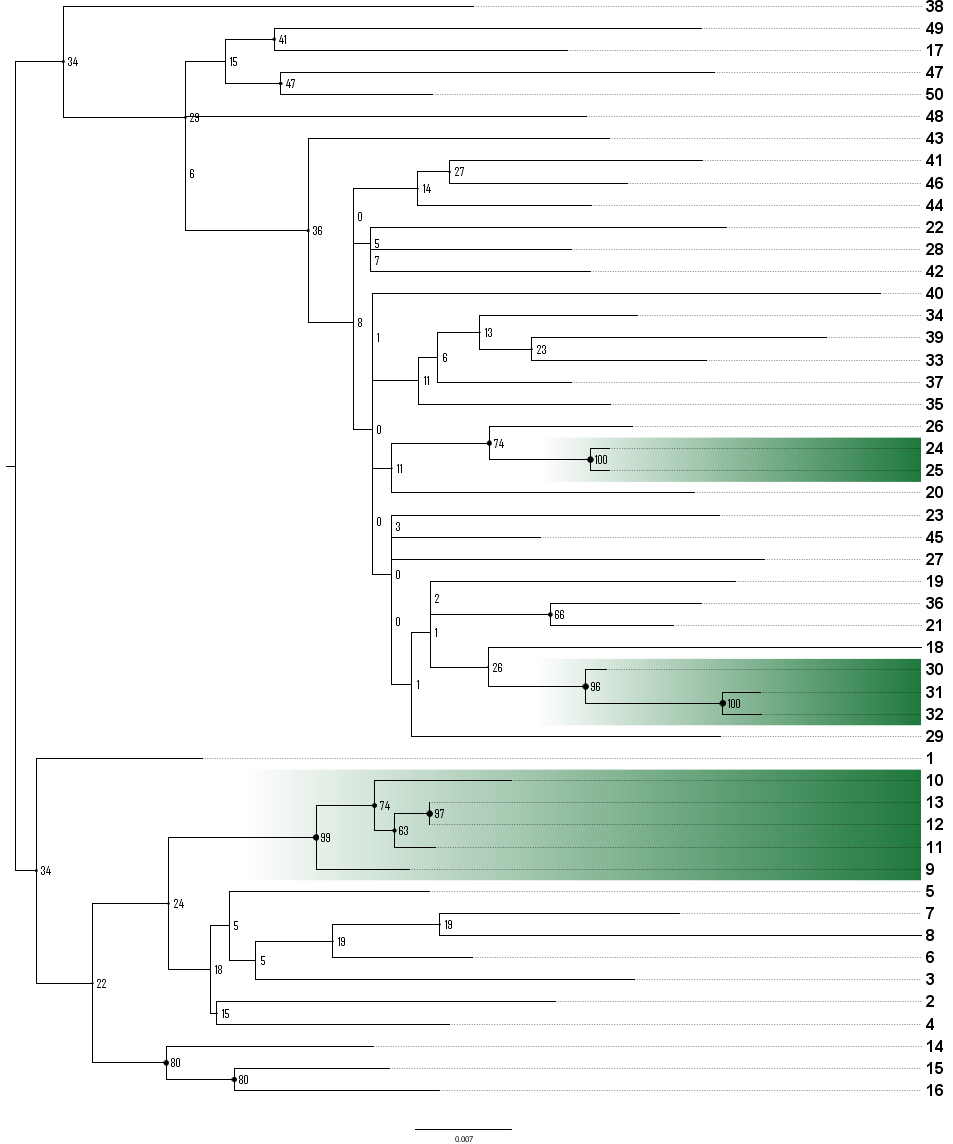
Genetic distance



Percentage of sequence clustering and average cluster size



Phylogenetic tree

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**S5 Figure: Clustering results among 50 GT1a ATAHC sequences with genetic distance, percentage of sequences, tree, patristic distance and bootstrap values.**