Shared Homology with Reference Alignment

SP-Score

Default Guide Tree   Chained Guide Tree

Number of Sequences
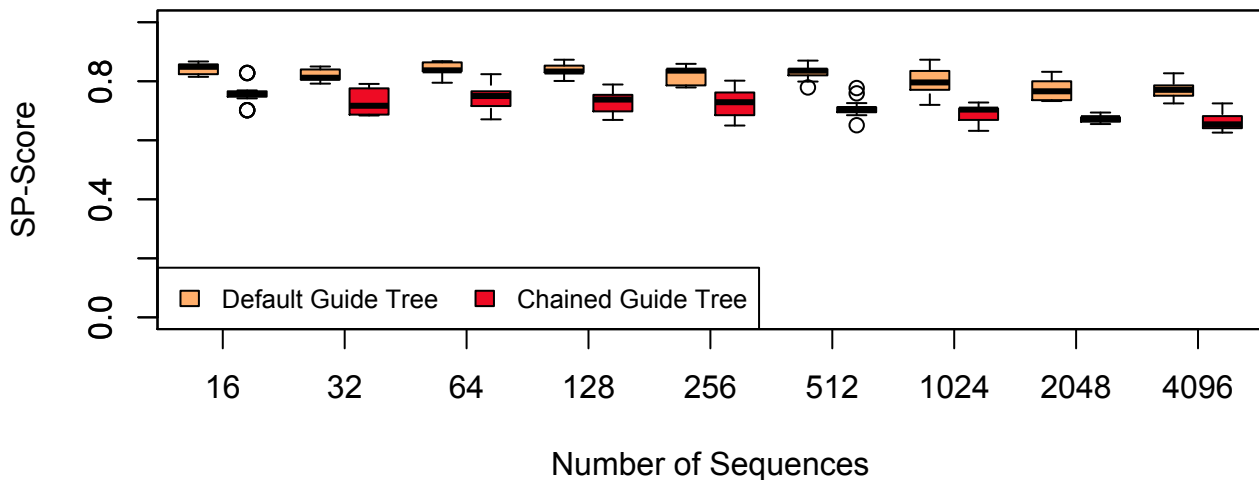
Total Reference Columns Preserved

TC-Score

Default Guide Tree   Chained Guide Tree

Number of Sequences