Shared Homology with Reference Alignment

Q-Score

Default Guide Tree   Chained Guide Tree

16   32   64   128   256   512   1024   2048   4096   8192

Number of Sequences
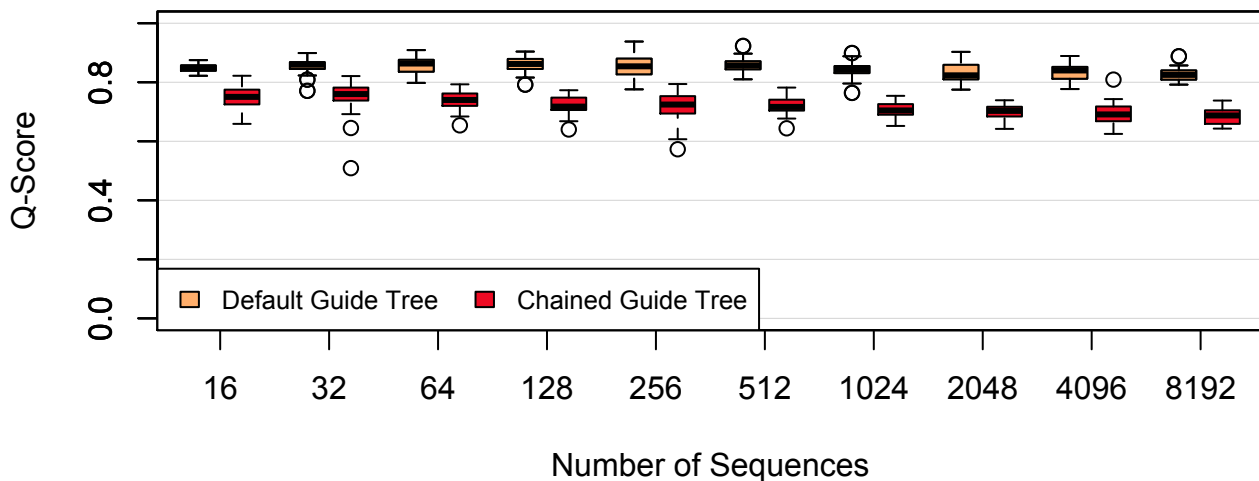
Total Reference Columns Preserved

TC-Score

Default Guide Tree   Chained Guide Tree

16   32   64   128   256   512   1024   2048   4096   8192

Number of Sequences