Recent introgression; speciation rate = 1

Tree age (Million years) 5 50 500 1.0 1.0 1.0 0.9 proportion 8.0 8.0 8.0 0.7 0.6 0.6 0.6 1.00 1.00 1.00 Cunningham 0.98 0.98 0.96 metric 0.96 0.96 0.92 0.94 0.94 1.0 1.0 8.0 TIGER value 8.0 8.0 0.6 0.6 0.6 0.4 0.4 0.4 0.6 8.0 0.4 0.6 0.4 Number of 0.4 0.2 0.2 taxa 0.2 5 0.0 0.0 0.0 10 0.05 0.15 0.05 20 Q-Residual value 0.04 0.04 50 0.10 0.03 0.03 100 0.02 0.02 0.05 0.01 0.01 0.00 0.00 0.00 1.0 1.0 1.0 sCF value 8.0 0.8 8.0 0.6 0.6 0.6 0.4 0.4 0.4 1.00 1.0 1.0 resolved quartets 0.9 0.75 Proportion 0.9 8.0 0.50 8.0 0.7 0.25 0.6 0.00 1.0treelike alignments 0.9 0.9 0.9 8.0 0.8 0.7 8.0 0.7 0.6 0.7 0.6 0.5 0.5 0.3 0.5 0.3 0.4 0.2 0.2 0.2 0.4 0.1

Median test statistic value

Proportion of recombinant DNA