## Recent introgression; speciation rate = 0.1

Tree age (Million years) 5 50 500 1.0 1.0 1.0 0.9 0.9 proportion 8.0 8.0 0.8 0.7 0.6 0.7 0.6 0.5 0.6 1.000 1.00-1.00 0.975 Cunningham 0.98 0.99 metric 0.950 0.96 0.98 0.925 0.94 0.900 0.92 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 0.6 0.4 0.4 0.4 Number of 0.2 0.2 0.2 taxa 5 0.0 0.0 0.0 10 0.08 0.08 20 Q-Residual value 0.006 0.06 0.06 50 0.004 100 0.04 0.04 0.002 0.02 0.02 0.00 0.00 0.000 1.0 1.0 1.0 0.9 0.9 sCF value 0.8 8.0 8.0 0.7 0.7 0.6 0.6 0.6 0.5 0.5 1.00 1.0 1.00 resolved quartets 0.9 Proportion 0.95 0.95 8.0 0.90 0.90 0.7-0.6 0.85 0.850.5 1.0-1.00 1.00 treelike alignments 0.75 0.75 0.9 0.50 0.50 8.0 0.25 0.25 0.7 0.00 0.00 0.5 0.5 0.5 0.3 0.3 0.2 0.4 0.1 0.1 0.1

Median test statistic value

Proportion of recombinant DNA