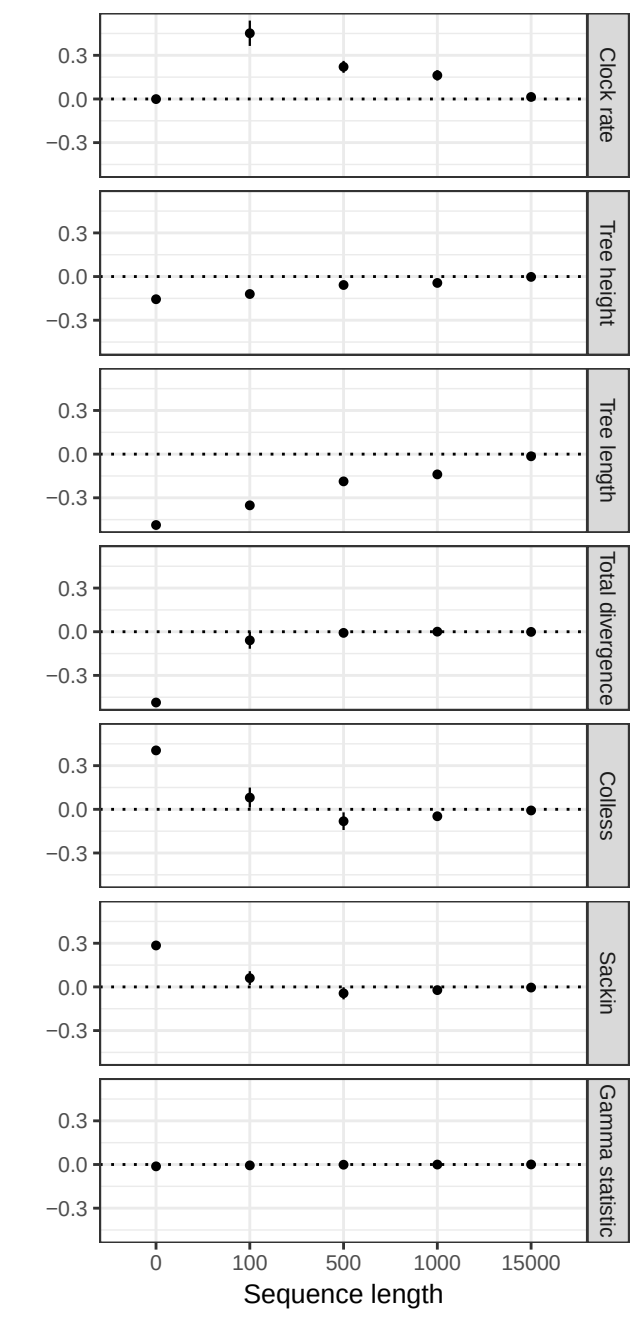
Clock rate
Tree height
Tree length
Total divergence
Colless
Sackin
Gamma statistic
0.3
0.0
−0.3
0
100
500
1000
15000
Sequence length