


- ▶ Scale: rateAC.s:seqs Scale GTR A-C substitution parameter of partition s:seqs 0.1
- ▶ Scale: rateAG.s:seqs Scale GTR A-G substitution parameter of partition s:seqs 0.1
- ▶ Scale: rateAT.s:seqs Scale GTR A-T substitution parameter of partition s:seqs 0.1
- ▶ Scale: rateCG.s:seqs Scale GTR C-G substitution parameter of partition s:seqs 0.1
- ▶ Scale: rateGT.s:seqs Scale GTR G-T substitution parameter of partition s:seqs 0.1
- ▼ Scale: clockRate.c:seqs Scale substitution rate of partition c:seqs 3.0

☒ estimate 

Parameter 0.1

Scale Factor 0.75

☐ Scale All☐ Scale All Independently

Degrees Of Freedom 0

☐ Root Only☒ Optimise

Upper 0.99999999

Lower 1.0E-8

Weight 0.1

- ▶ Up Down: clockRate.c:seqs Tree.t:seqs Scale up substitution rate c:seqs and scale down tree t:(\$n) 0.0
- ▶ Scale: Tree.t:seqs Scales all internal nodes for tree t:seqs 3.0
- ▶ Scale: Tree.t:seqs Scales root node for tree t:seqs 3.0
- ▶ Uniform: Tree.t:seqs Draws new internal node heights uniformly for tree t:seqs 30.0
- ▶ Subtree Slide: Tree.t:seqs Performs subtree slide rearrangement of tree t:seqs 15.0
- ▶ Exchange: Tree.t:seqs Narrow exchange performs local rearrangement of tree t:seqs 15.0
- ▶ Exchange: Tree.t:seqs Wide exchange performs global rearrangement of tree t:seqs 3.0
- ▶ Wilson Balding: Tree.t:seqs Performs Wilson-Balding global rearrangement of tree t:seqs 3.0
- ▶ Scale: popSize.t:seqs Scale population size of Coalescent prior of tree t:seqs 3.0