ML ancestral state reconstruction using rayDISC (R:corHMM) pollination, ER model

Node ML state Prob Angiospermae biotic 0.9993

abioticbiotic

Angiospermae biotic 0.9993

Mesangiospermae biotic 1

Magnoliidae biotic 1

Monocotyledoneae biotic 1

Eudicotyledoneae biotic 1

Model LogL Npar AIC AICc DeltaAICc w q01 q10 ARD -71.86 2 147.72 147.76 1.85 0.16 0.0022 0.0013 ARDeq -71.17 2 146.34 146.38 0.47 0.31 0.0021 0.0013 ER* -71.95 1 145.9 145.92 0 0.39 0.0013 0.0013 UNI01 -78.61 1 159.22 159.23 13.32 0 0.0145 UNI10 -73.01 1 148.02 148.03 2.12 0.14 0.0013

