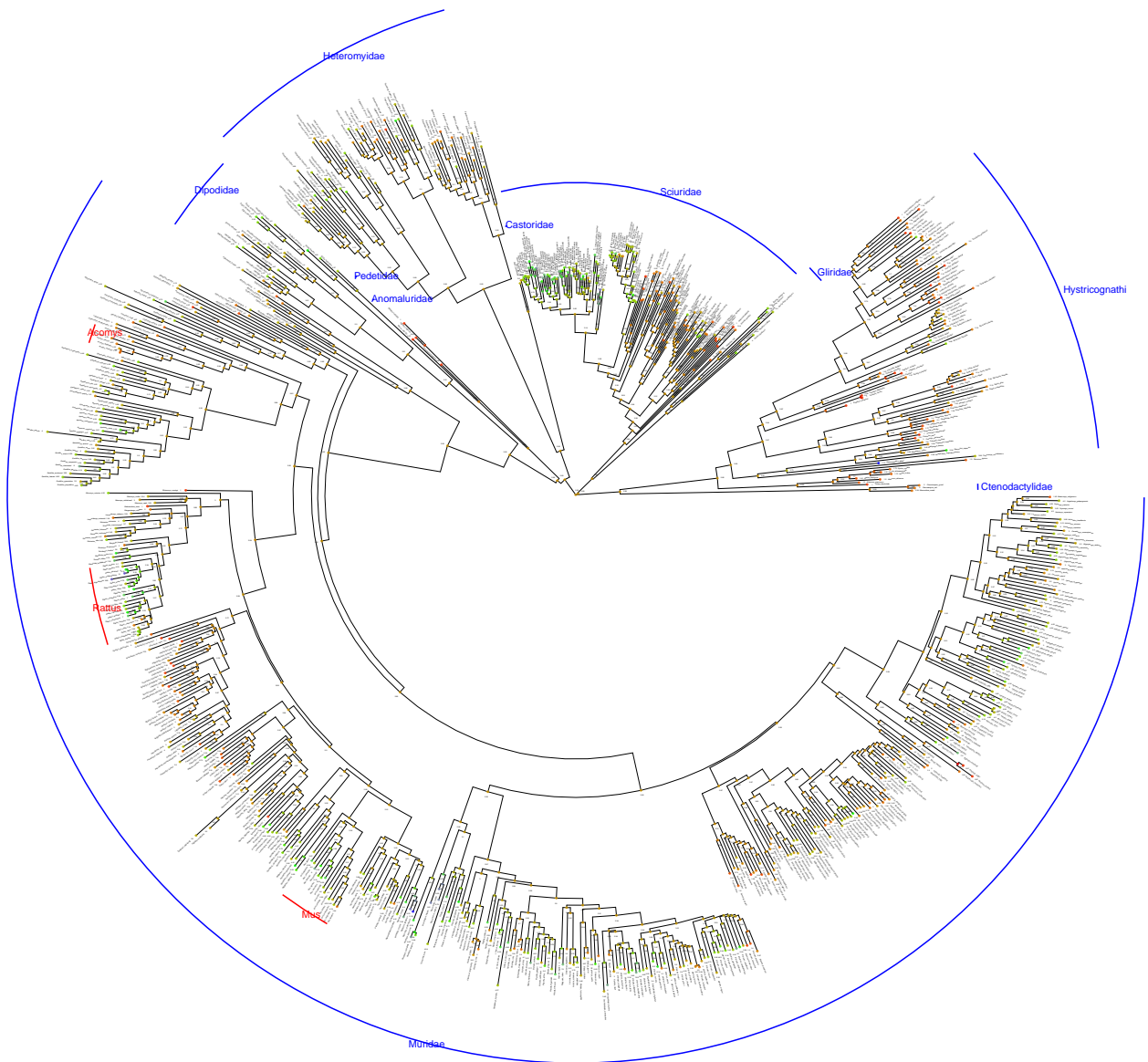
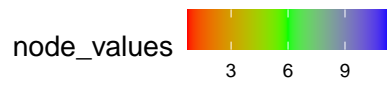


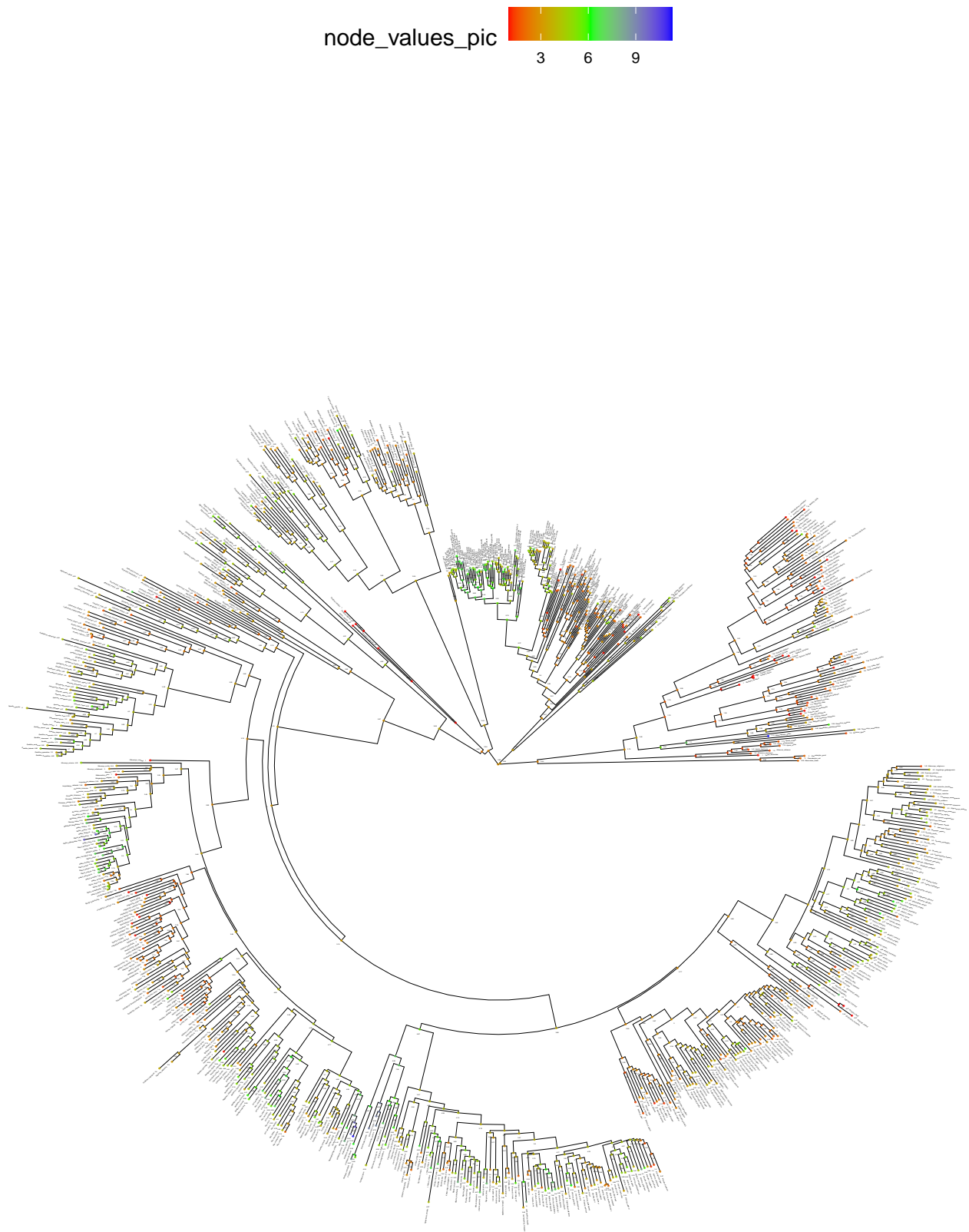
Litter Size Evolution

Consensus tree extracted from an Upham et al. paper (Jetz lab)
Data: 1006 out of 2278 species

Warning in sqrt(1/out\$hessian): NaNs produced



Ancestral Recon with pic



This gives us a variety of information about the ancestral character reconstructions:

- `$ace` contains the reconstructed ancestral character states. There is one value per internal node, and

- they are in the order that the internal nodes are numbered in.
- σ^2 is the estimate of the Brownian Motion parameter.
 - CI_{95} are the confidence intervals on the ancestral character state reconstructions.