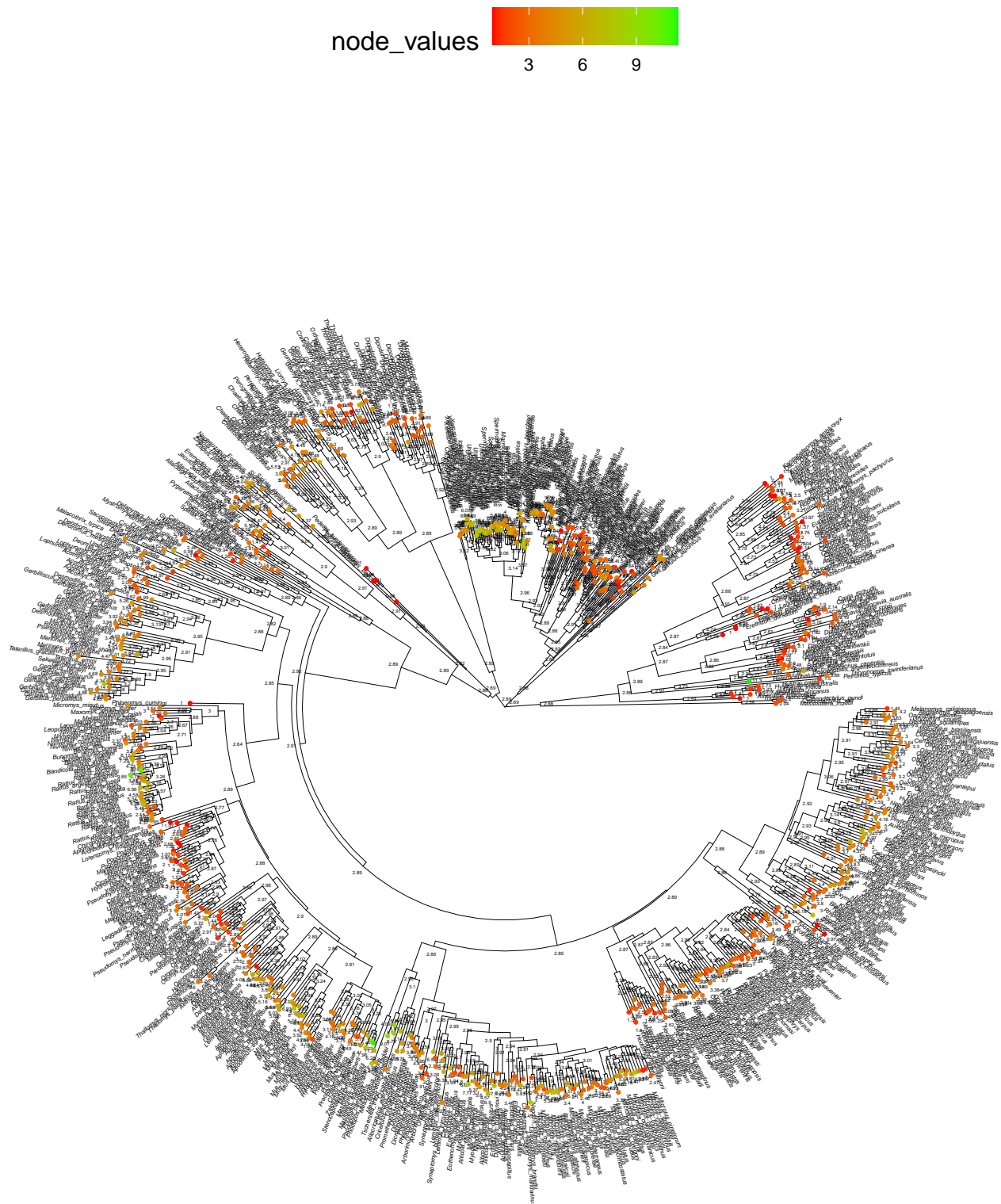


# Litter Size Evolution

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**



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

- `$ace` contains the reconstructed ancestral character states. There is one value per internal node, and they are in the order that the internal nodes are numbered in.
- `$sigma2` is the estimate of the Brownian Motion parameter.
- `$CI95` are the confidence intervals on the ancestral character state reconstructions.