

#### Exercises, part 4:

1. Simulate a birth-death tree with 200 tips using  $b=0.1$  and  $d=0.05$ . Plot the tree and make an LTT plot.
2. Simulate a character on your tree using a brownian motion model with a rate of 1.0.
3. Test your simulated character on the tree for phylogenetic signal.
4. Estimate ancestral character states for your character. The root state in the tree was actually 0 - how close did you get?

#### Exercises, part 5

1. Fit a birth-death model to the phelsuma tree. Also calculate the gamma statistic. What do these results suggest?
2. Use `fitContinuous` to compare BM, EB, and OU to the phelsuma data. Which model is best supported by the data?