Print out and fill in the last page.

- 1. likelihood calculations
  - (a) As data, use a character in which C is the state for taxa 1 and 2 , taxon 3 has state G, and taxon 4 has a T
  - (b) Fill in the circle with character state (either the observed data for tips or the inferred states)
  - (c) There are 16 trees on each page, use them to show all 16 possible ancestral character state combinations.
  - (d) Calculate the probability of each reconstruction under the Kimura model.
  - (e) Use  $\kappa = 2$
  - (f) For branch lengths, use (1:0.03,3:0.4, (2:0.03,4:0.4):0.03)
- 2. Which branch length has the higher ratio of the transition probability to transversion probability?
- 3. What is the likelihood for this character?

I recommend, using a spreadsheet (or some other software) to:

- 1. calculate all three change probabilities (no change, transition, transversion) for the 0.03 branch length.
- 2. calculate all three change probabilities (no change, transition, transversion) for the 0.4 branch length
- 3. multiply the appropriate numbers.

If you label the cells of the spreadsheet logically, I'll be able to figure out your system.

