

Allman Ch 4.4 Prob 3,5,8,9,10, 15 Allman Ch 4.5 Ex 1,3,5,7,13

4.4.3. a. About 27 steps to be within .05; about 67 steps to be within .01.

4.4.5. Because mutation is rare, the conditional probabilities describing no change should be largest.

4.4.8. a. The first theorem applies to M , but the second does not since M has some zero entries. (However, since M^2 has all non-zero entries, you can apply the second theorem to it.)

b. (.1849, .3946, .2819, .1386)

4.4.9. a. $\mathbf{p}_0 = (.3, .225, .25, .225)$, $M = \begin{pmatrix} .833 & 0 & 0 & .111 \\ .083 & .889 & 0 & 0 \\ 0 & .111 & 1 & .111 \\ .083 & 0 & 0 & .778 \end{pmatrix}$

b. \mathbf{p}_0 is reasonable close to (.25, .25, .25, .25). M may seem less close to a Jukes-Cantor matrix than you might expect, because of the variation in the off-diagonal entries. One way to estimate α is to average the off-diagonal entries to estimate $\alpha/3$. This gives $\alpha/3 = .0416$, so $\alpha = .1248$.

4.4.10. a. The Jukes-Cantor model is more appropriate for the pair S'_0, S'_1 , since a particular base seems to mutate to any of the other three bases with roughly the same frequency. Note also that the bases in S'_0 are in roughly equal numbers.

b. The Kimura 2-parameter model is more appropriate for the pair S_0, S_1 , since the data shows that transitions are more likely than transversions. Note also that the bases in S_0 are in roughly equal numbers.

4.4.15. a. $\mathbf{p}_0 = (.15, .25, .35, .25)$ is not an equilibrium base distribution for the Jukes-

Cantor matrix $M = \begin{pmatrix} .7 & .1 & .1 & .1 \\ .1 & .7 & .1 & .1 \\ .1 & .1 & .7 & .1 \\ .1 & .1 & .1 & .7 \end{pmatrix}$

b. $\mathbf{p}_0 = (.19, .25, .31, .25)$ is not an equilibrium base for the transition matrix

$M = \begin{pmatrix} .5526 & .06 & .0484 & .06 \\ .1316 & .7 & .0806 & .1 \\ .1842 & .14 & .7903 & .14 \\ .1316 & .1 & .0806 & .7 \end{pmatrix}$, which is not Jukes-Cantor.

c. \mathbf{p}_0 is unchanged by multiplying by M ; it is an equilibrium. Notice that \mathbf{p}_0 is an eigenvector of M with eigenvalue 1.

d. The initial vector is drawn towards the stable equilibrium (.25, .25, .25, .25). This \mathbf{p}_0 corresponds to an initial sequence comprised entirely of G 's.

4.5.1. .1367

- 4.5.3. a. .2224580274
 b. .2308224444
 c. The Kimura 2-parameter distance is probably a better choice (assuming we did not already know that the sequences were created with the Kimura 2-parameter model). The frequency table shows a definite pattern of more transitions than transversions. Notice too that the distances differ in the second decimal position.
- 4.5.5. Graph the Jukes-Cantor distance on a graphing calculator or computer.
 a. If the sequences are identical, then $p = 0$. This means the Jukes-Cantor distance is $-.75 \log(1) = 0$.
 b., c. Mathematically, if two sequences differ in more than $3/4$ of the sites, then $p > 3/4$. Then the Jukes-Cantor distance formula requires taking the logarithm of a negative number, which is impossible. This is not a limitation with real data. If we took two sequences that were in no way related, we would expect that about $1/4$ of the sites agree and about $3/4$ of the sites disagree, since with a uniform distribution of bases about 25% of the time the two sequences should agree if everything is chosen at random. For related sequences the formulas for the Jukes-Cantor model derived in the last section show p is at most $3/4$, and in practice p is usually much less than $3/4$. Notice that the Jukes-Cantor distance gets huge as the values of p get close to .75. This is desirable, since distances should be large when comparing sequences that appear almost unrelated.
- 4.5.7. Substituting $(1 - q)$ for p yields $d_{JC} = -\frac{3}{4} \ln(1 - \frac{4}{3}p) = -\frac{3}{4} \ln(1 - \frac{4}{3}(1 - q)) = -\frac{3}{4} \ln(\frac{4}{3}q - \frac{1}{3}) = -\frac{3}{4} \ln(\frac{4q-1}{3})$.
- 4.5.13. The Kimura 3-parameter distance is given by $d_{K3} = -\frac{1}{4} (\ln(1 - 2\beta - 2\gamma) - \ln(1 - 2\beta - 2\delta) - \ln(1 - 2\beta - 2\gamma))$. Substituting $\alpha/3$ for β , γ , and δ gives

$$\begin{aligned}
 d &= -\frac{1}{4} (\ln(1 - 2\alpha/3 - 2\alpha/3) - \ln(1 - 2\alpha/3 - 2\alpha/3) - \ln(1 - 2\alpha/3 - 2\alpha/3)) \\
 &= \frac{1}{4} (3 \ln(1 - 4\alpha/3)) = d_{JC}.
 \end{aligned}$$