

Figure 1
Ancestral recombination graph.

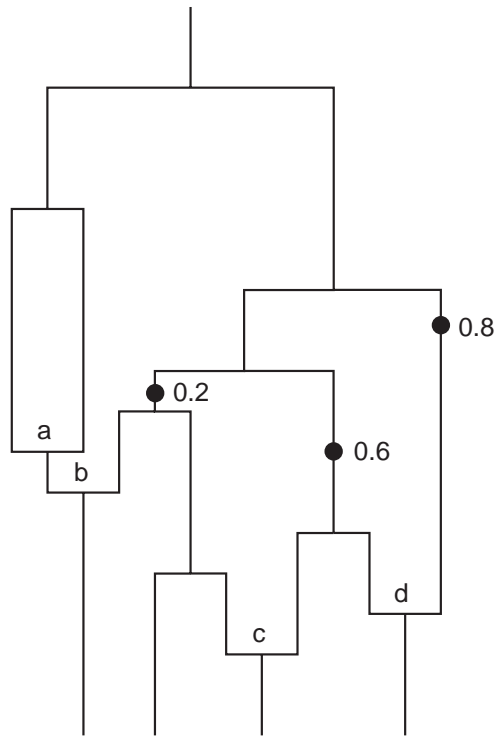


Figure 2
Marginal ancestral tree.

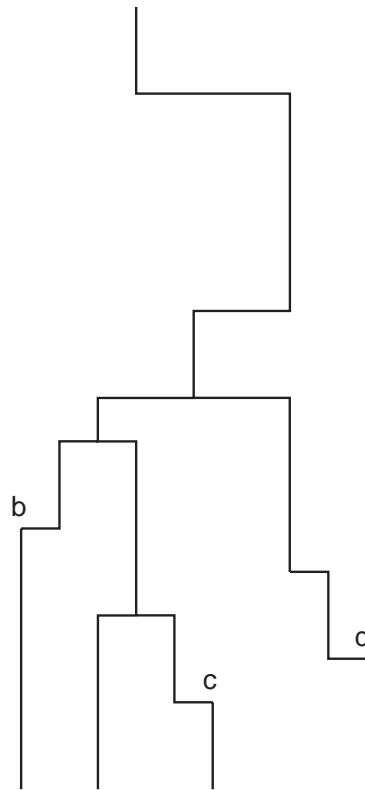


Figure 3

A sample of sequences, showing mutations.

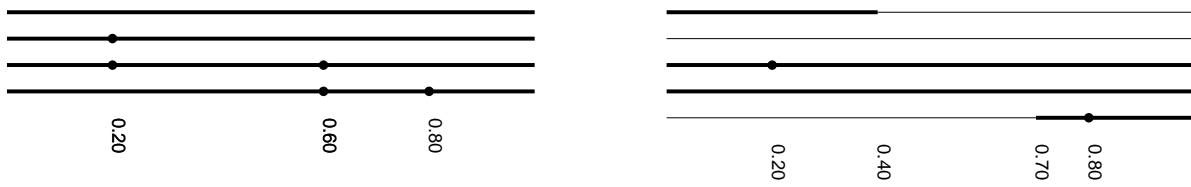


Figure 4
TMRCA along the sequence.

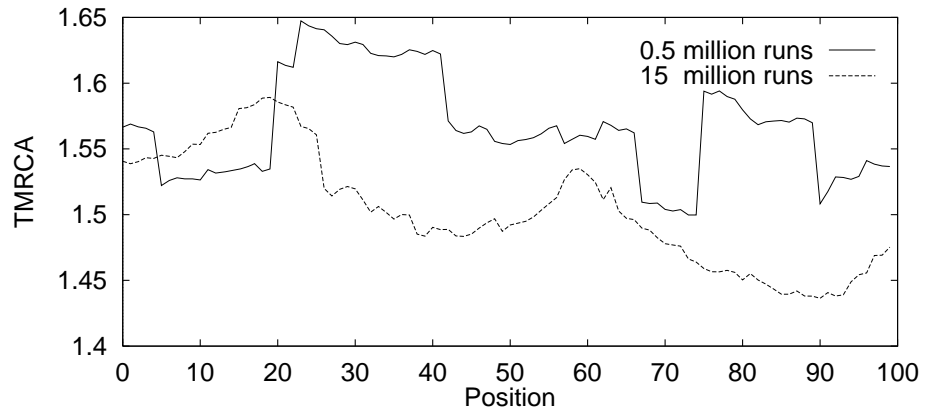


Figure 5
TMRCA distribution.

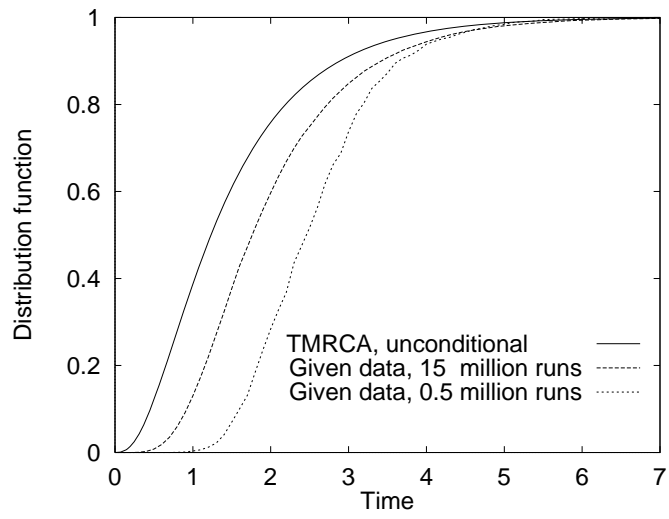


Figure 6

TMRCAs in two sequences with no mutations.

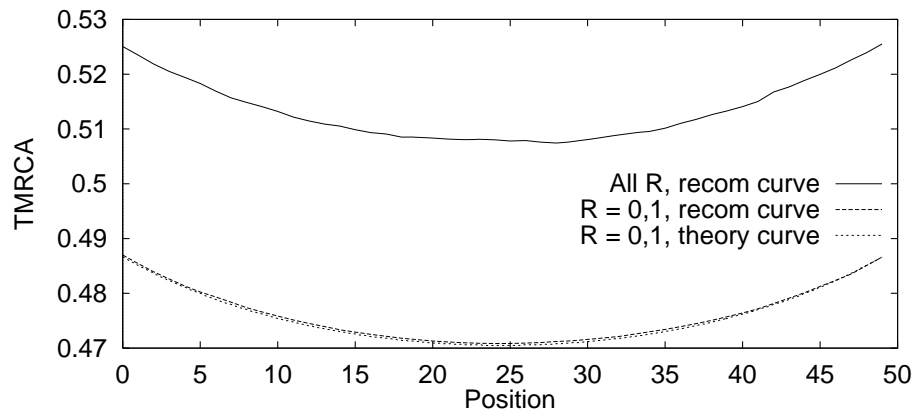


Figure 7

TMRCAs in two sequences with mutations.

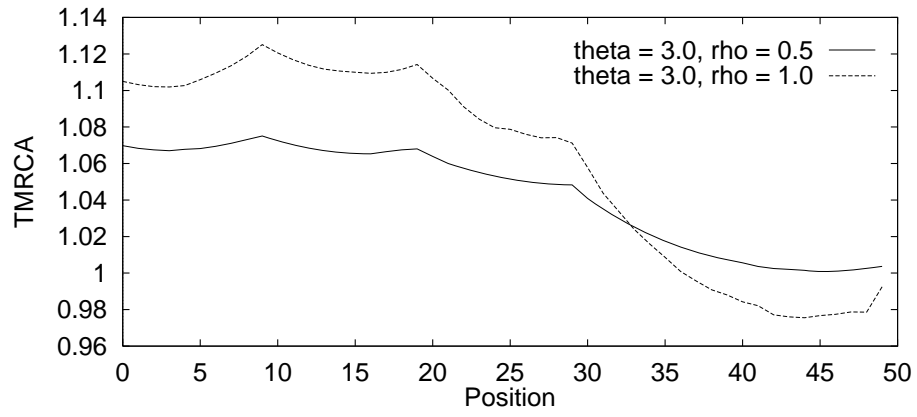


Figure 8

Sample sequences, showing recombination.

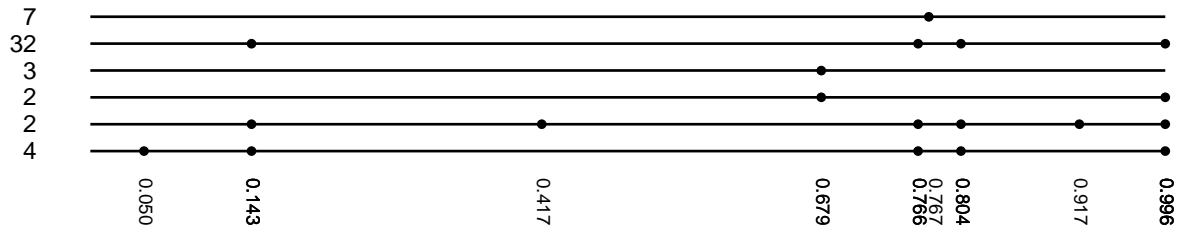


Figure 9
Genealogical tree.

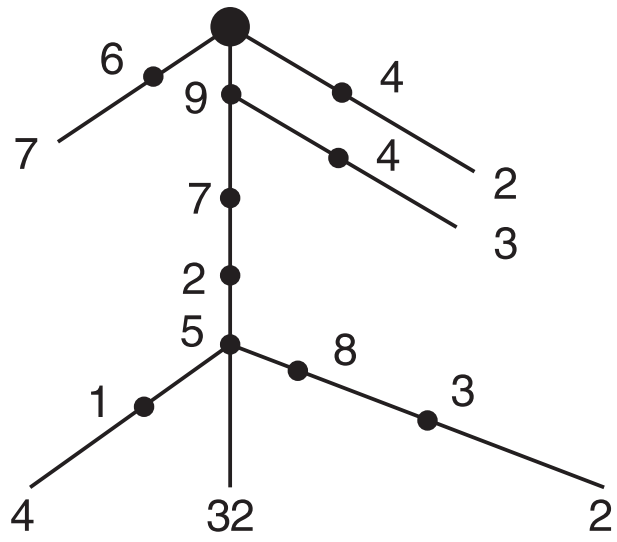


Figure 10
Recombination graph.

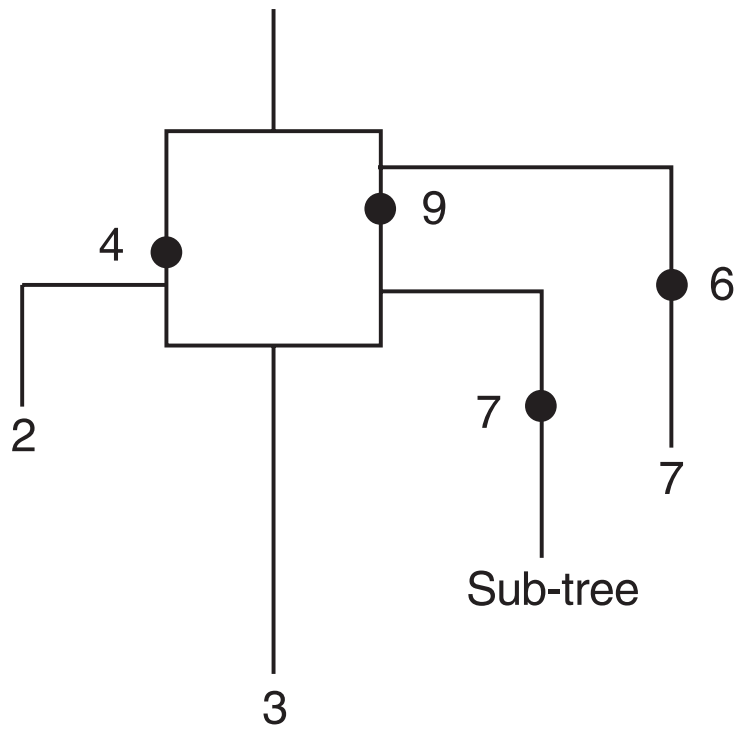


Figure 11

Likelihood surface, generated with $\theta = 2.0$, $\rho = 0.5$.

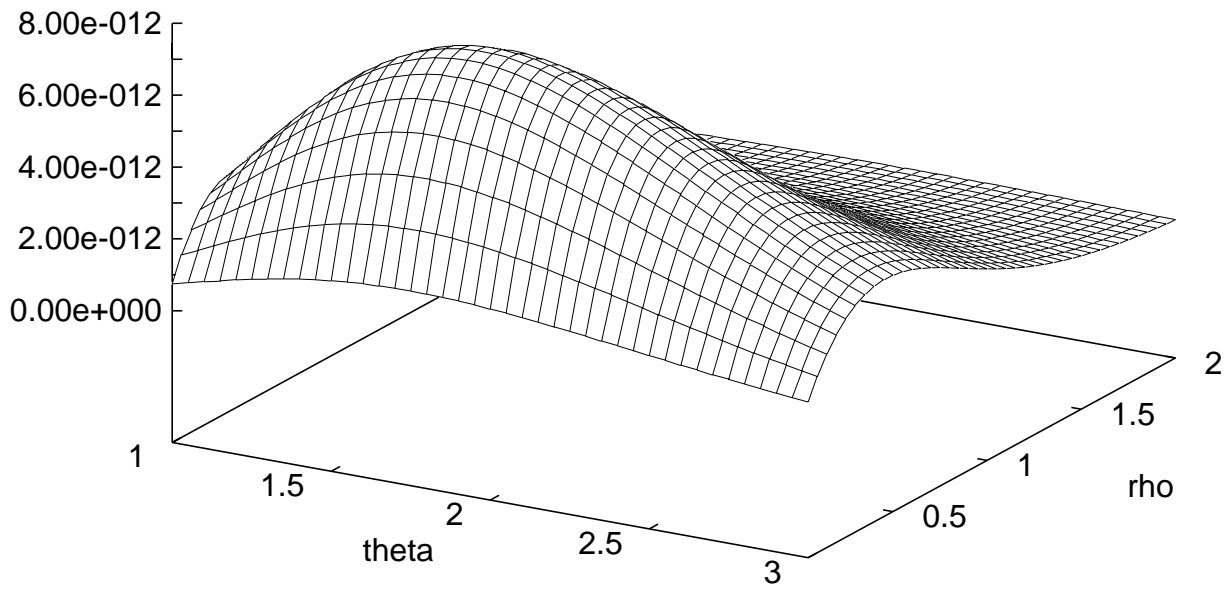


Figure 12
Recombination hits along sequences.

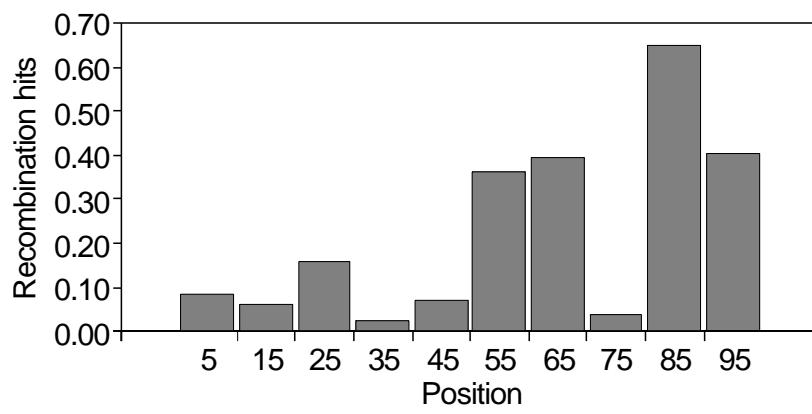


Figure 13
TMRCAs along sequences.

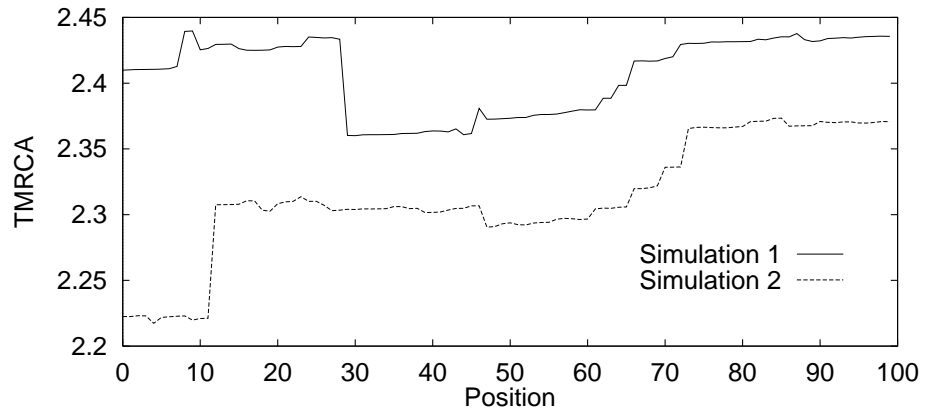


Figure 14

Likelihood surface of sample with no recombination.

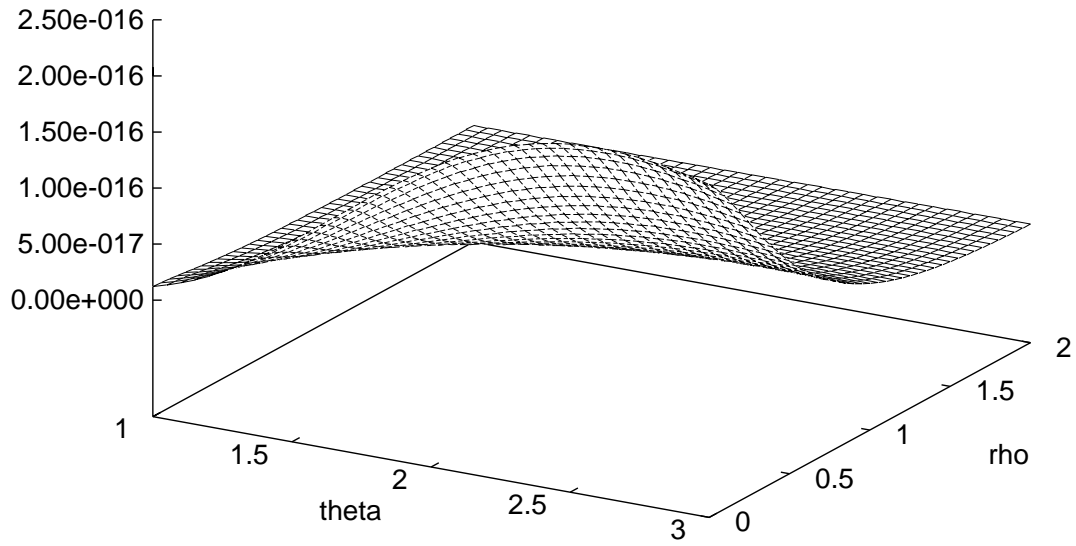


Figure 15

Sample sequences, with no recombination.

