- 1. Consider a sample of n = 2 DNA sequences from a population of size $N_e = 250$ diploid individuals (i.e., 2N = 500 sequences).
 - a) What is the expected coalescence time for the two sequences?
 - b) What is the probability that coalescence occurs at generation 500?
- 2. Consider a sample of n = 10 DNA sequences from a population of size $N_e = 2500$ diploid individuals.
 - a) What is the expected time to the 1^{st} , 2^{nd} , 3^{rd} ... 9^{th} coalescence event?
 - b) What is the expected coalescence time to the MRCA of all 10 sequences?
 - c) What is the expected number of polymorphisms in the sample? Assume that the sequences are 1,000 base pairs long and that the mutation rate is 10^{-8} per base-pair per generation.
 - d) What is the probability that all sequences are identical and no polymorphisms are observed in the sample?
- 3. Two 1,000 bp sequences were compared and 10 differences were observed. Assuming the mutation rate is 10^{-8} per base-pair, and the effective population size is $N_e = 10,000$, what is the probability of observing more than 3 differences? What assumptions did you make to complete the calculation?

What might explain the large number of differences between the two sequences?