- 1. Simulate the following with paper and pencil and a dice (or the R command sample (6, 1)) as source of random.
 - (a) A haploid Wright-Fisher population of 6 haploid individuals
 - (b) Three individuals are sampled from the present generation. Identify their most recent common ancestors.
- 2. A number $k \in \{2, 3, 4, 5, 10, 20, 50, 100, 200, 1000\}$ of haploid individuals are sampled from a large Hardy-Weinberg population. Calculate for their genealogy...
 - (a) ... the expected time τ to their most recent common ancestor and ...
 - (b) ... the expected total length ℓ of all branches and ...
 - (c) ... the expected number of mutations on these trees for $\theta = 10$.
 - (d) Visualize how τ and ℓ depend on k.
 - (e) Find a function that can easily be expressed without sum signs and approximates how ℓ depends on k.
- 3. Use the R package scrm to simulate trees under three different scenarios:
 - (a) The standard case of a panmictic population of constant size
 - (b) An exponentially growing population
 - (c) A population that consists of two subpopulations with migration.

Visualize the genealogies. For which growth rates and migration rates do the trees begin to look different than the standard coalescent?

- 4. FOR BIOINFORMATICIANS AND OTHER PROGRAMMING ENTHUSIASTS: Develop a program that reads a sample size and outputs a random standard coalescent for this sample size in Newick format.
- 5. Assume a haploid Wright-Fisher population of N individuals. Thus, we assume discrete generations and each individual in generation 0 selects purely randomly and independent of all other individuals one parent in generation -1.
 - (a) Let a be an individual in generation -1 and let X be the number of his or her kids in generation 0. Compute Pr(X = 2).
 - (b) Compute Pr(X = k) for general $k \in \mathbb{N}$.
 - (c) Which fraction the individuals of a generation do not have offspring?
 - (d) Assume that the population size N is very large. Then, $\Pr(X = k)$ can be approximated by a term of the form

$$b^c \cdot e^d / f!.$$

Which are the appropriate values for b, c, d and f?

(e) Make a rough calculation: Which fraction of the individuals of a generation will have offspring

- i. after 10 generations?
- ii. after 100 generations?
- iii. after $10 \cdot N$ generations?
- (f) in a generation -g, that is g generations before the present generation 0, let 0.2 be the fraction of individuals who still have offspring in the present population 0. Which fraction of individuals from generation -(g + 1) do still have offspring in generation 0?
- (g) Does your answer to question 5f shed new light on part 5e?