

Homework 4: Genetic algorithm functions I

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How to represent simplified genotypes

In genetic algorithms or in simulating organic evolution, it is common to represent a simplified "genotype" as a vector of numbers, characters, or bits (1s and 0s or TRUEs and FALSEs).

We will start building a system for simulating the evolution of such vectors which will represent haploid genotypes. One of the first functions we need for such a program is a function that takes two parents and produces an offspring. In our first version of this function, we will ignore mutation, but we will be concerned with recombination. The function will have the following interface:

```
# mateVectors(x, y, r):  
# Returns a child vector created by "mating" two parent  
# vectors x and y and supplying a per-adjacent-locus recombination rate.  
#  
# Args:  
# - x, y = the two parent vectors. These must be of same  
# type (numeric, character, or logical) and same  
# length.  
# - r = the per-locus recombination rate  
#  
# Returns:  
# - A vector of the same length as the parents
```

An outline of the algorithm

We are simulating a simplified haploid recombination mechanism. This is not meant to directly represent biology but to provide a way to simulate mating with limited recombination and allow spatial correlation such that adjacent alleles are more likely to stay together. Your function will use the following algorithm for mating and recombination:

- Randomly select one parent from which to begin reading values and transferring these values (alleles) to offspring
- After each vector element is read and the result copied to the child vector, check for a crossing over event (with probability r)
- On a recombination event, a crossover occurs and the reading then switches to the second parent.

