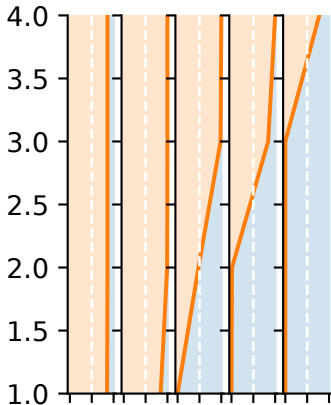


population
size

10^5 10^7 10^9

available
beneficial mutations



fixation probability

genotype

nonmutator

mutator