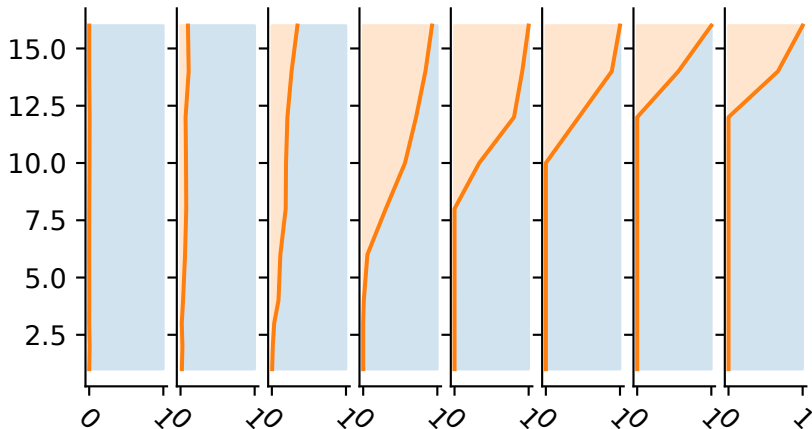


population
size

10^3 10^4 10^5 10^6 10^7 10^8

available
beneficial mutations



fixation probability



normomutator



hypermutator