

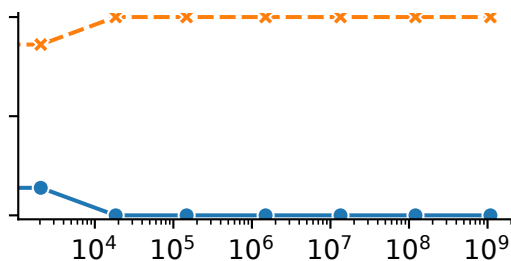
—●— normomutator    -✕- hypermutator

n=0 agents per node

available beneficial mutations = 1

available beneficial mutations = inf

fixation probability



population size

population size