

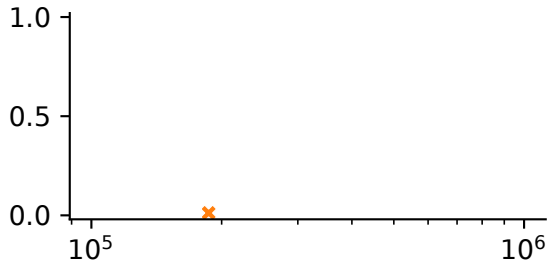
● normomutator      ✕ hypermutator

n=256 agents per node

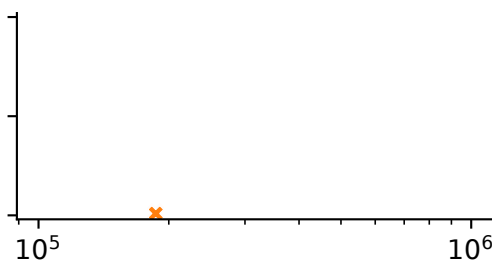
available beneficial mutations = 1

available beneficial mutations = 2

fixation probability



population size



population size