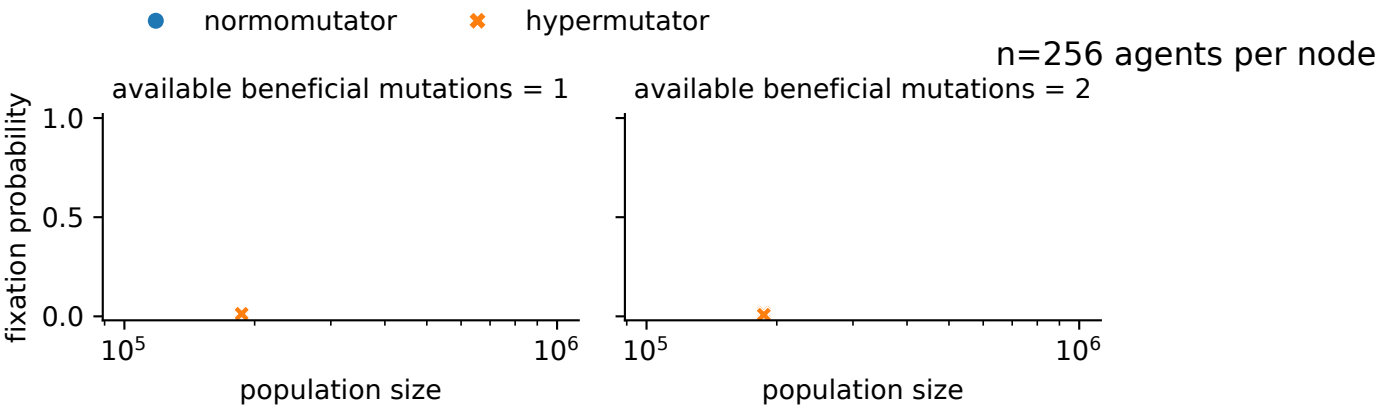
normomutator
hypermutator
n=256 agents per node
available beneficial mutations = 1
available beneficial mutations = 2
fixation probability
1.0
0.5
0.0
10^5
10^6
population size