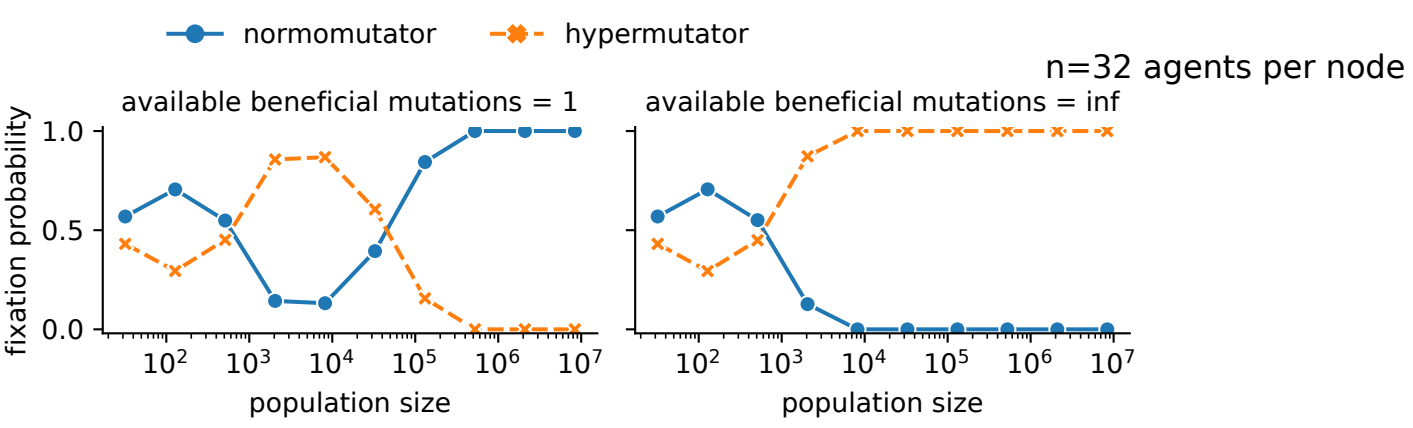
normomutator
hypermutator
n=32 agents per node
available beneficial mutations = 1
available beneficial mutations = inf
fixation probability
1.0
0.5
0.0
10^2
10^3
10^4
10^5
10^6
10^7
population size