

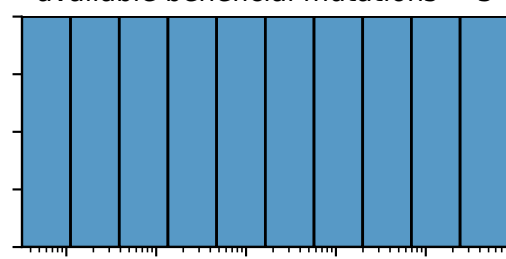
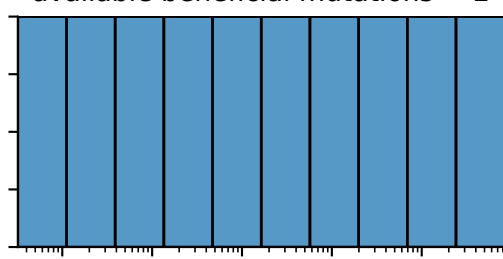
False

n=32 agents per node

available beneficial mutations = 1

available beneficial mutations = 2

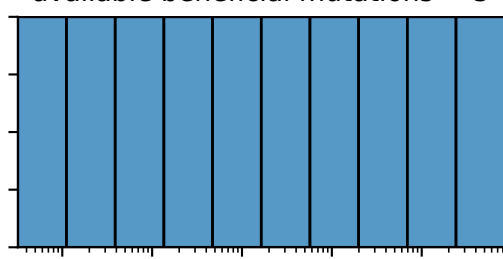
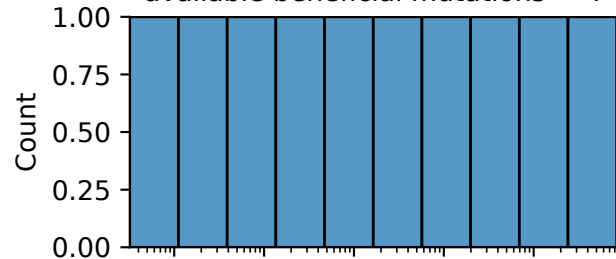
available beneficial mutations = 3



population size

available beneficial mutations = 4

available beneficial mutations = 5



population size

population size