

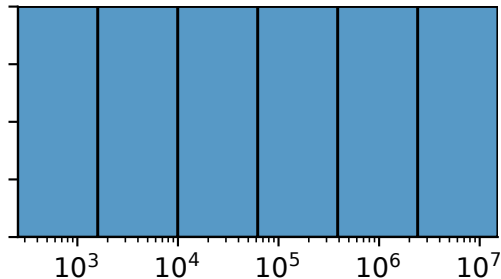
False

n=256 agents per node

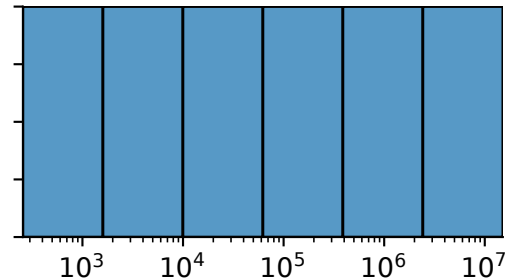
available beneficial mutations = 1



available beneficial mutations = 2



available beneficial mutations = 3



available beneficial mutations = 4

