

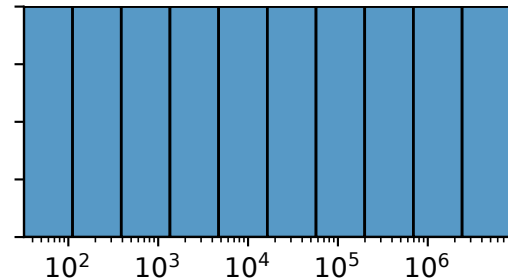
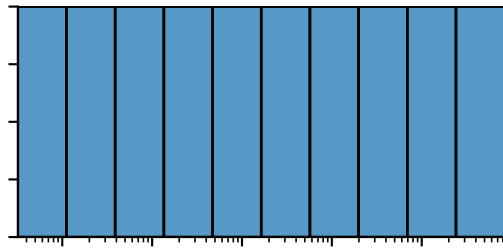
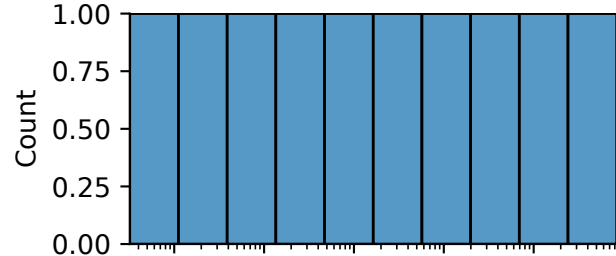
False

n=32 agents per node

available beneficial mutations = 1

available beneficial mutations = 2

available beneficial mutations = 3



available beneficial mutations = 4

available beneficial mutations = 5

