

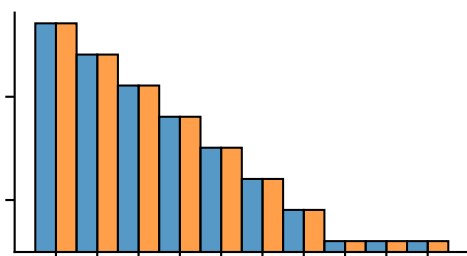
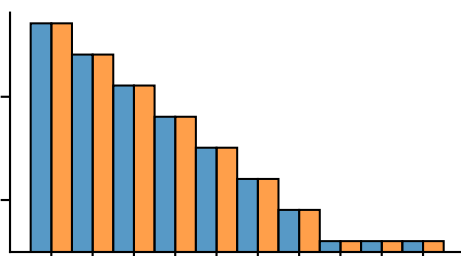
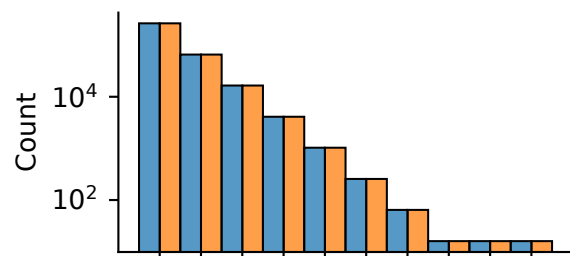
normomutator hypermutator

n=32 agents per node

available beneficial mutations = 1

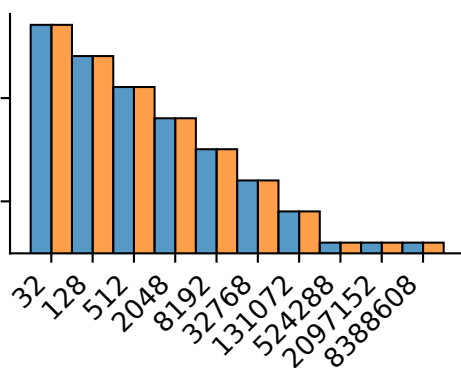
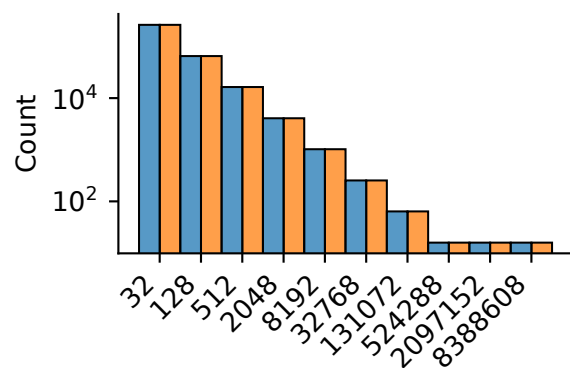
available beneficial mutations = 2

available beneficial mutations = 3



available beneficial mutations = 4

available beneficial mutations = 5



population size cat

population size cat

population size cat