

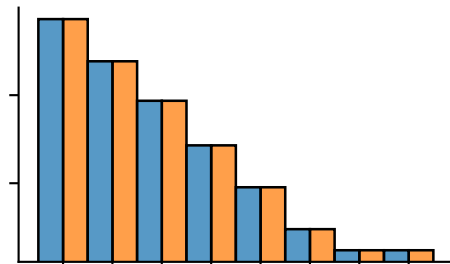
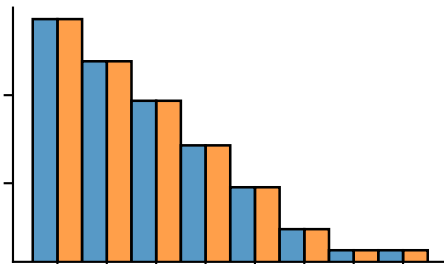
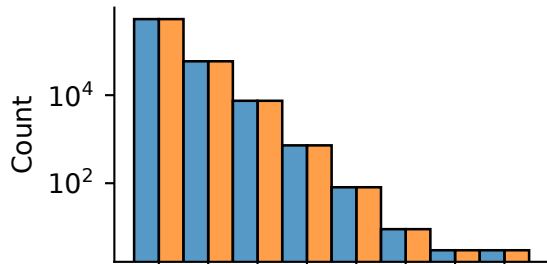
normomutator hypermutator

n=2048 agents per node

available beneficial mutations = 1

available beneficial mutations = 2

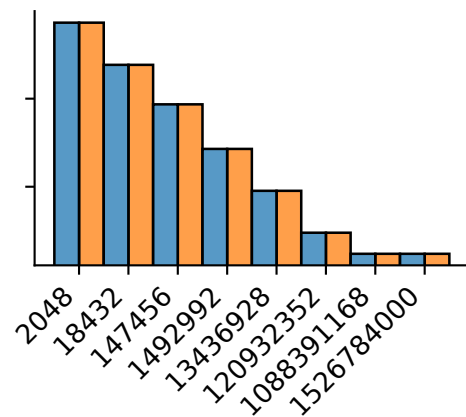
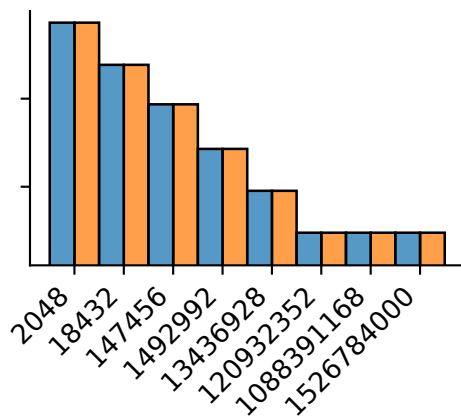
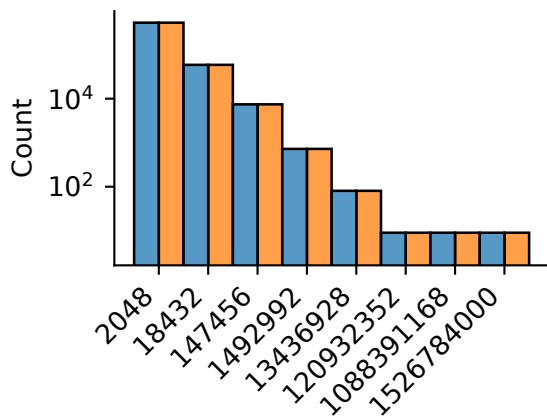
available beneficial mutations = 3



available beneficial mutations = 4

available beneficial mutations = 5

available beneficial mutations = 6



population size cat

population size cat

population size cat