

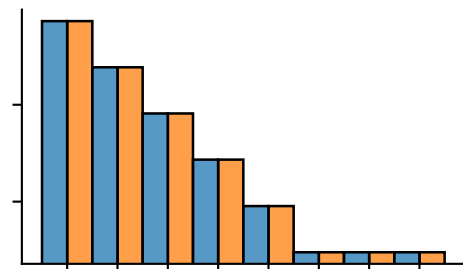
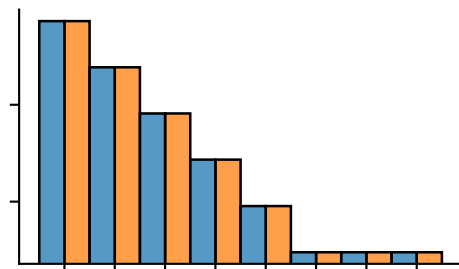
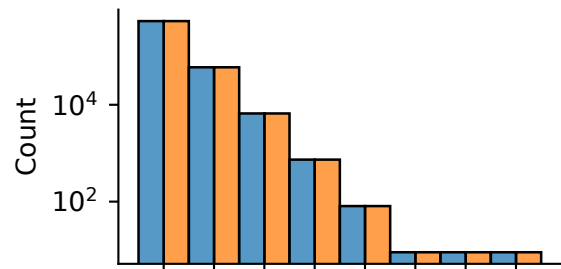
normomutator hypermutator

n=256 agents per node

available beneficial mutations = 1

available beneficial mutations = 2

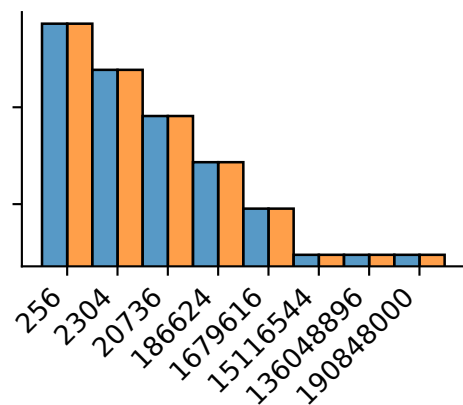
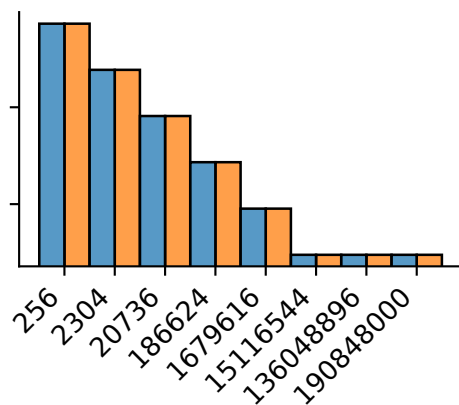
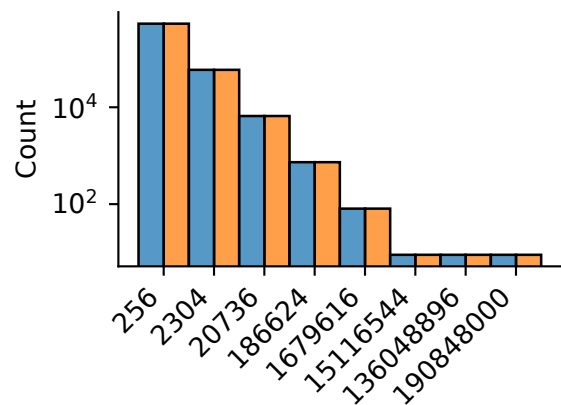
available beneficial mutations = 3



available beneficial mutations = 4

available beneficial mutations = 10

available beneficial mutations = 30



population size cat

population size cat

population size cat