

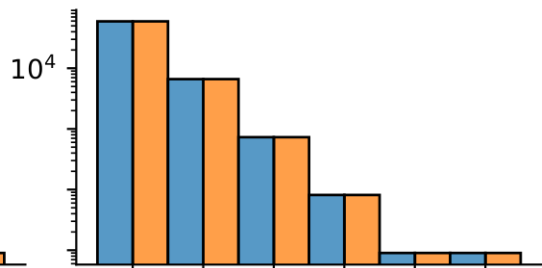
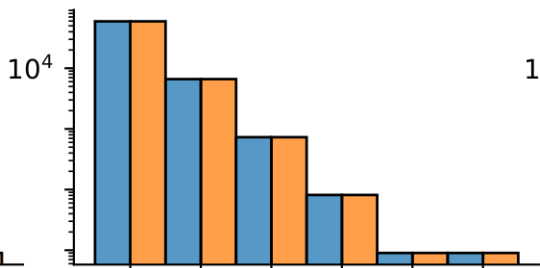
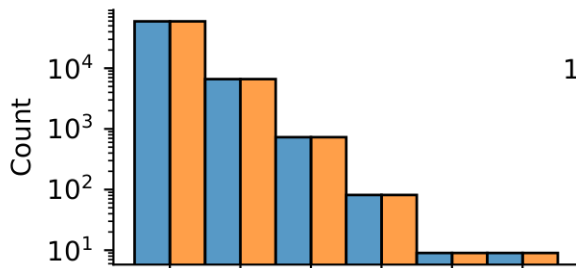
normomutator hypermutator

n=256 agents per node

available beneficial mutations = 4

available beneficial mutations = 8

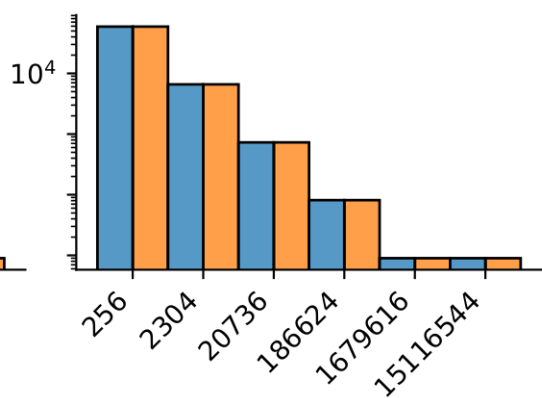
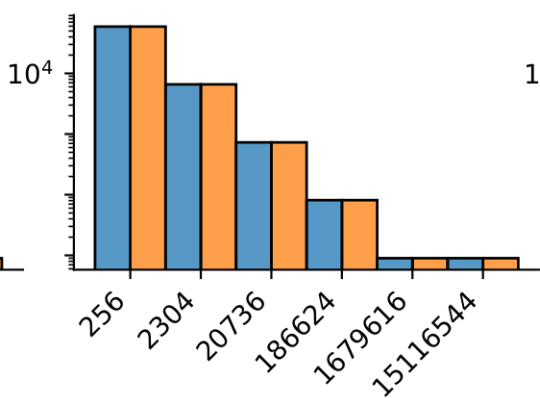
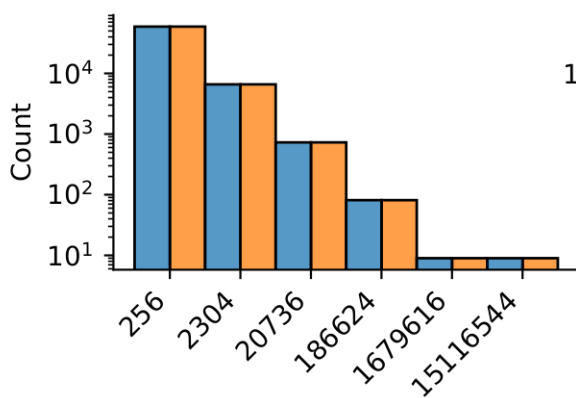
available beneficial mutations = 10



available beneficial mutations = 12

available beneficial mutations = 14

available beneficial mutations = 16



population size cat

population size cat

population size cat