

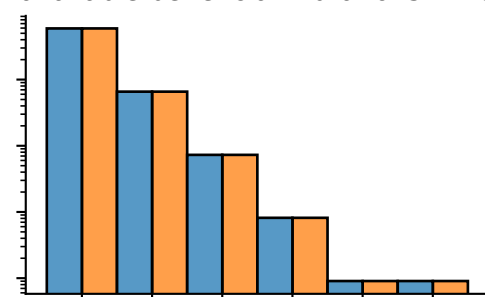
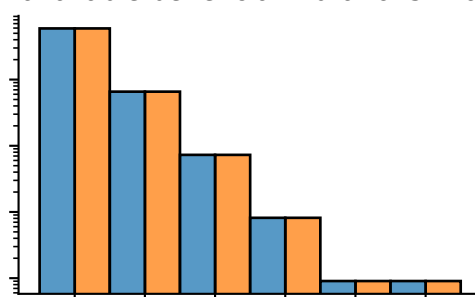
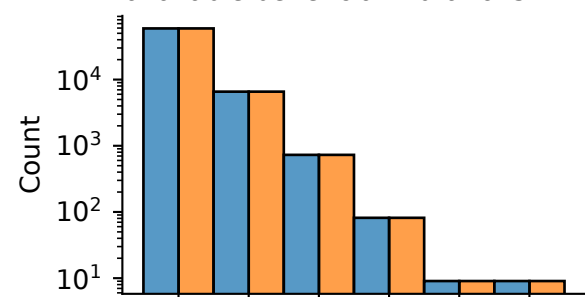
n=256 agents per node

normomutator hypermutator

available beneficial mutations = 2

available beneficial mutations = 6

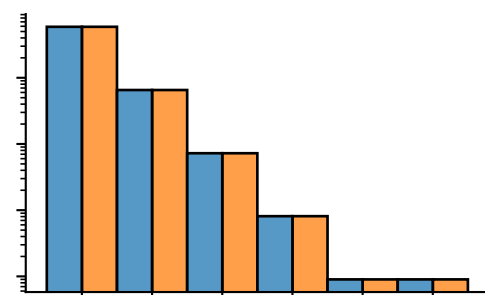
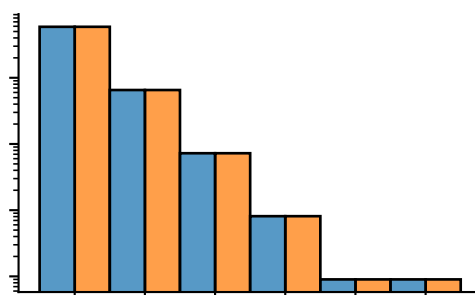
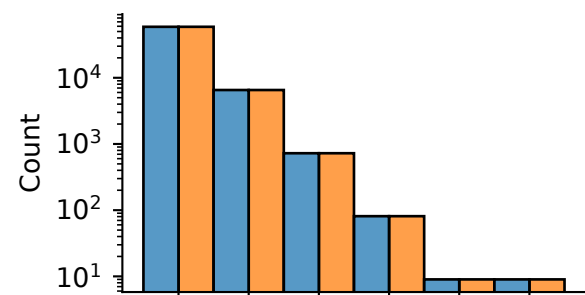
available beneficial mutations = 10



available beneficial mutations = 12

available beneficial mutations = 14

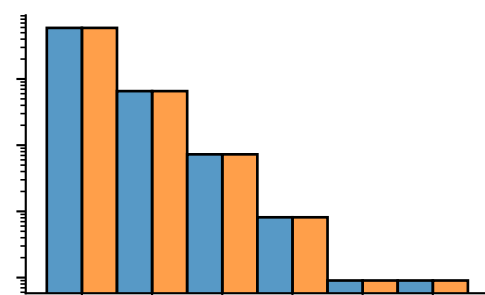
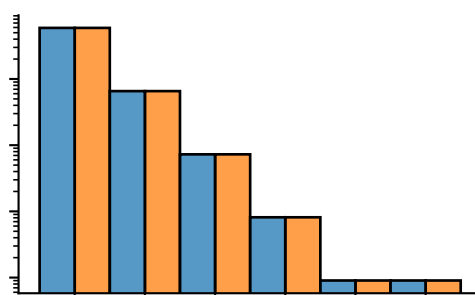
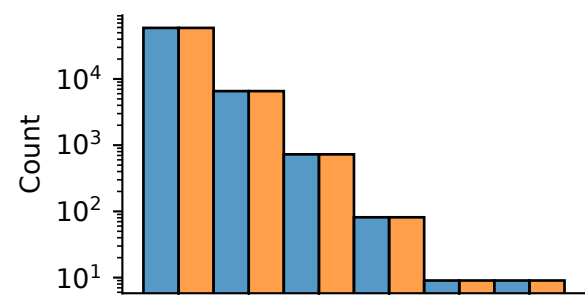
available beneficial mutations = 16



available beneficial mutations = 18

available beneficial mutations = 20

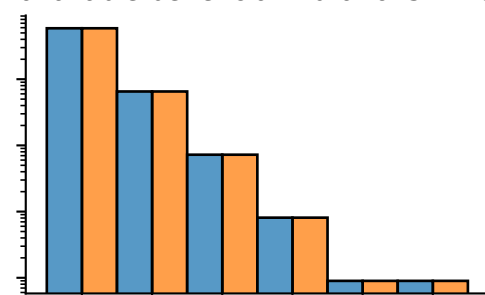
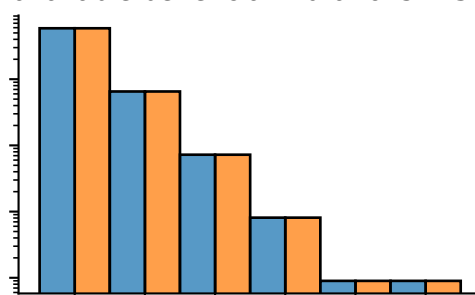
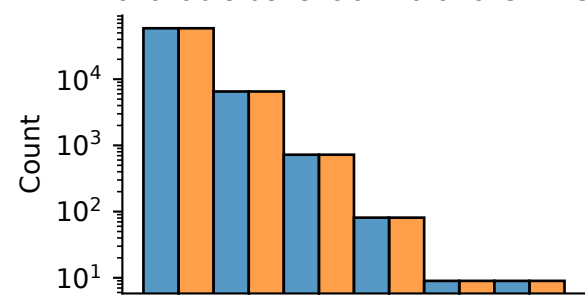
available beneficial mutations = 24



available beneficial mutations = 28

available beneficial mutations = 32

available beneficial mutations = 40



population size cat

population size cat

population size cat