

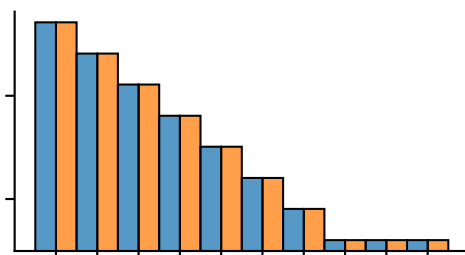
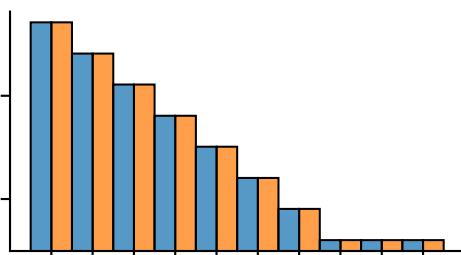
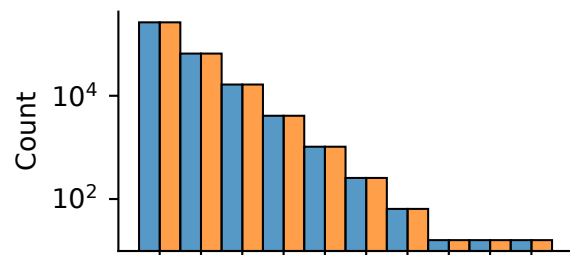
normomutator hypermutator

n=32 agents per node

available beneficial mutations = 1

available beneficial mutations = 2

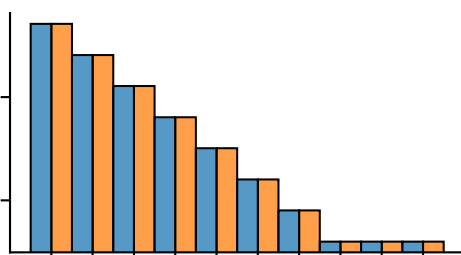
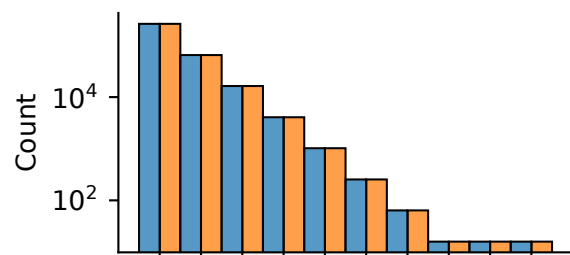
available beneficial mutations = 3



32 128 512 2048 8192 32768 131072 524288 2097152 8388608

available beneficial mutations = 4

available beneficial mutations = 5



population size cat

32 128 512 2048 8192 32768 131072 524288 2097152 8388608

32 128 512 2048 8192 32768 131072 524288 2097152 8388608

population size cat

population size cat