

n=256 agents per node

normomutator hypermutator

available beneficial mutations = 2

available beneficial mutations = 4

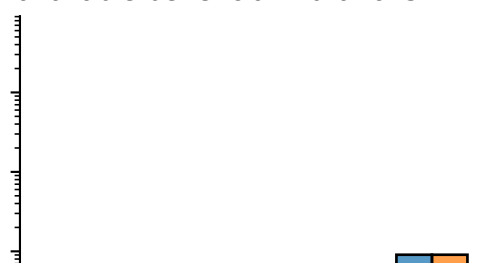
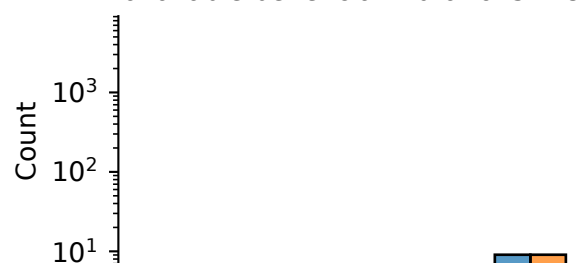
available beneficial mutations = 6



available beneficial mutations = 8

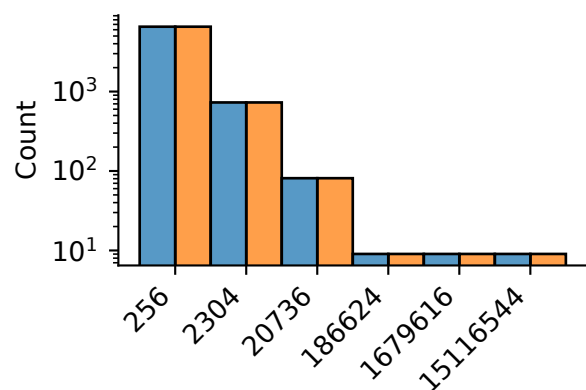
available beneficial mutations = 10

available beneficial mutations = 12



available beneficial mutations = 14

available beneficial mutations = 16



population size cat