

normomutator hypermutator

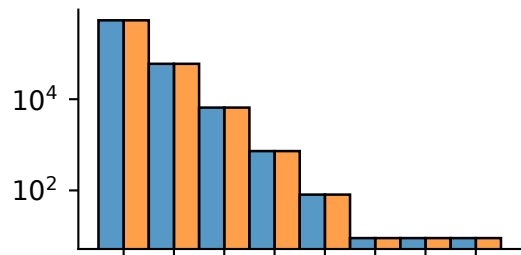
n=256 agents per node

available beneficial mutations = 1

available beneficial mutations = 2

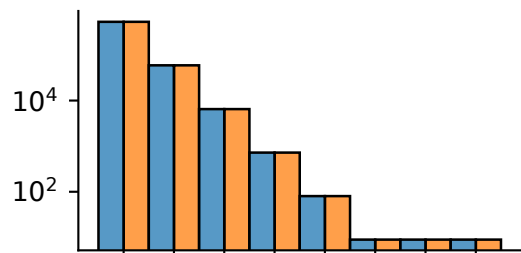
available beneficial mutations = 3

Count



available beneficial mutations = 4

Count



population size cat

population size cat