

n=256 agents per node

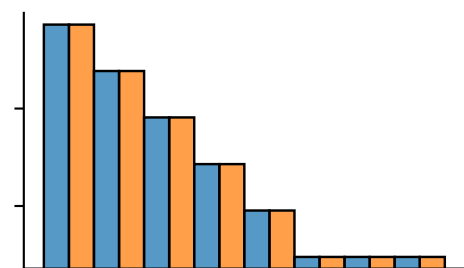
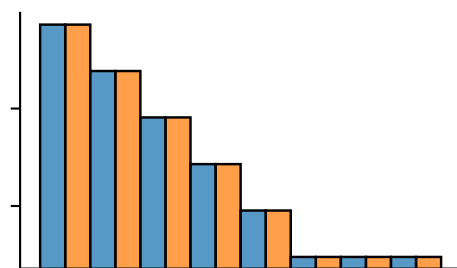
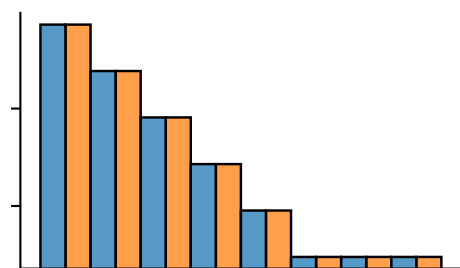
normomutator hypermutator

available beneficial mutations = 1

available beneficial mutations = 2

available beneficial mutations = 3

Count

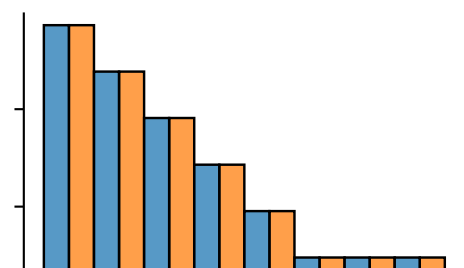
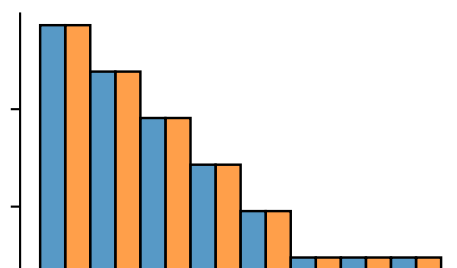
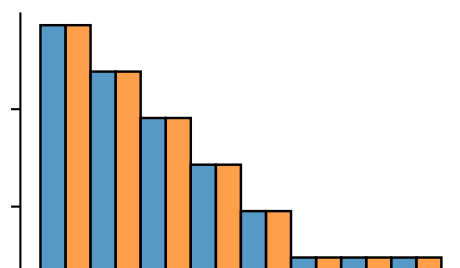


available beneficial mutations = 4

available beneficial mutations = 6

available beneficial mutations = 8

Count

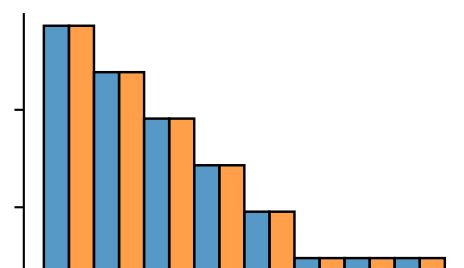
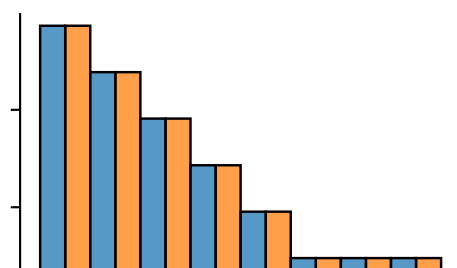
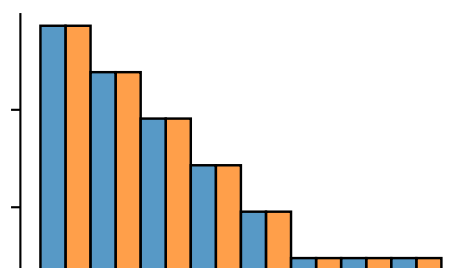


available beneficial mutations = 10

available beneficial mutations = 12

available beneficial mutations = 14

Count



available beneficial mutations = 16

population size cat

population size cat

Count

