

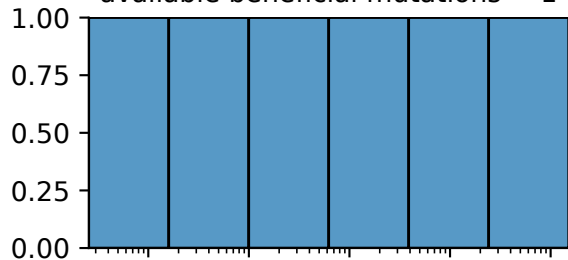
False

n=256 agents per node

available beneficial mutations = 1

available beneficial mutations = 2

available beneficial mutations = 3



available beneficial mutations = 4

