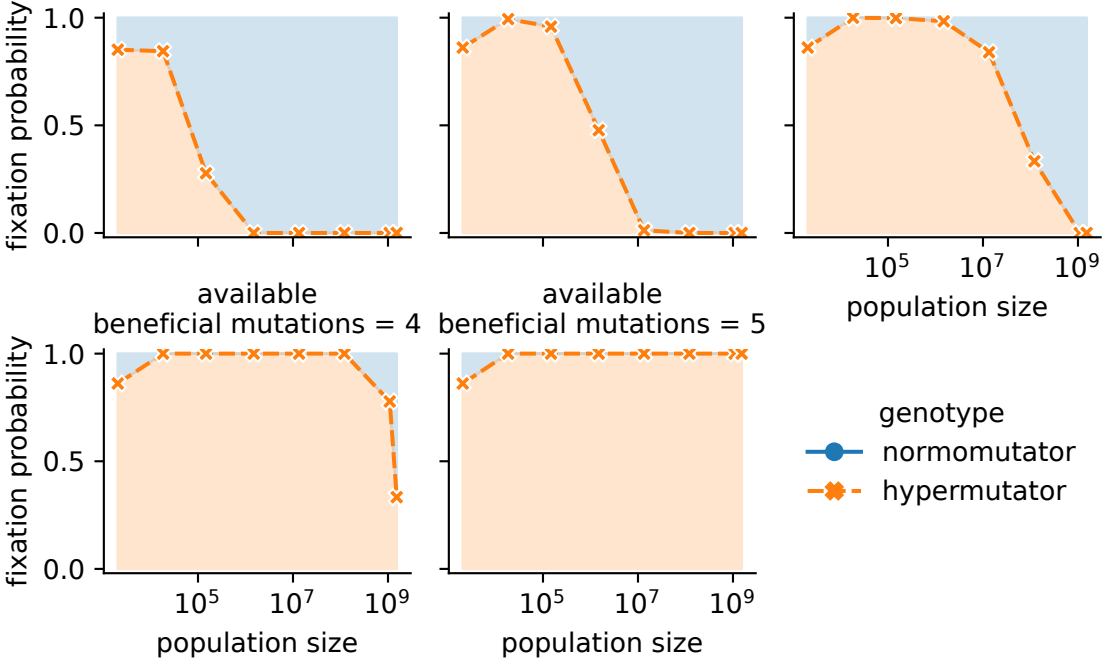
available
beneficial mutations = 1
available
beneficial mutations = 2
available
beneficial mutations = 3
available
beneficial mutations = 4
available
beneficial mutations = 5
fixation probability
1.0
0.5
0.0
10^5
10^7
10^9
population size
genotype
normomutator
hypermutator