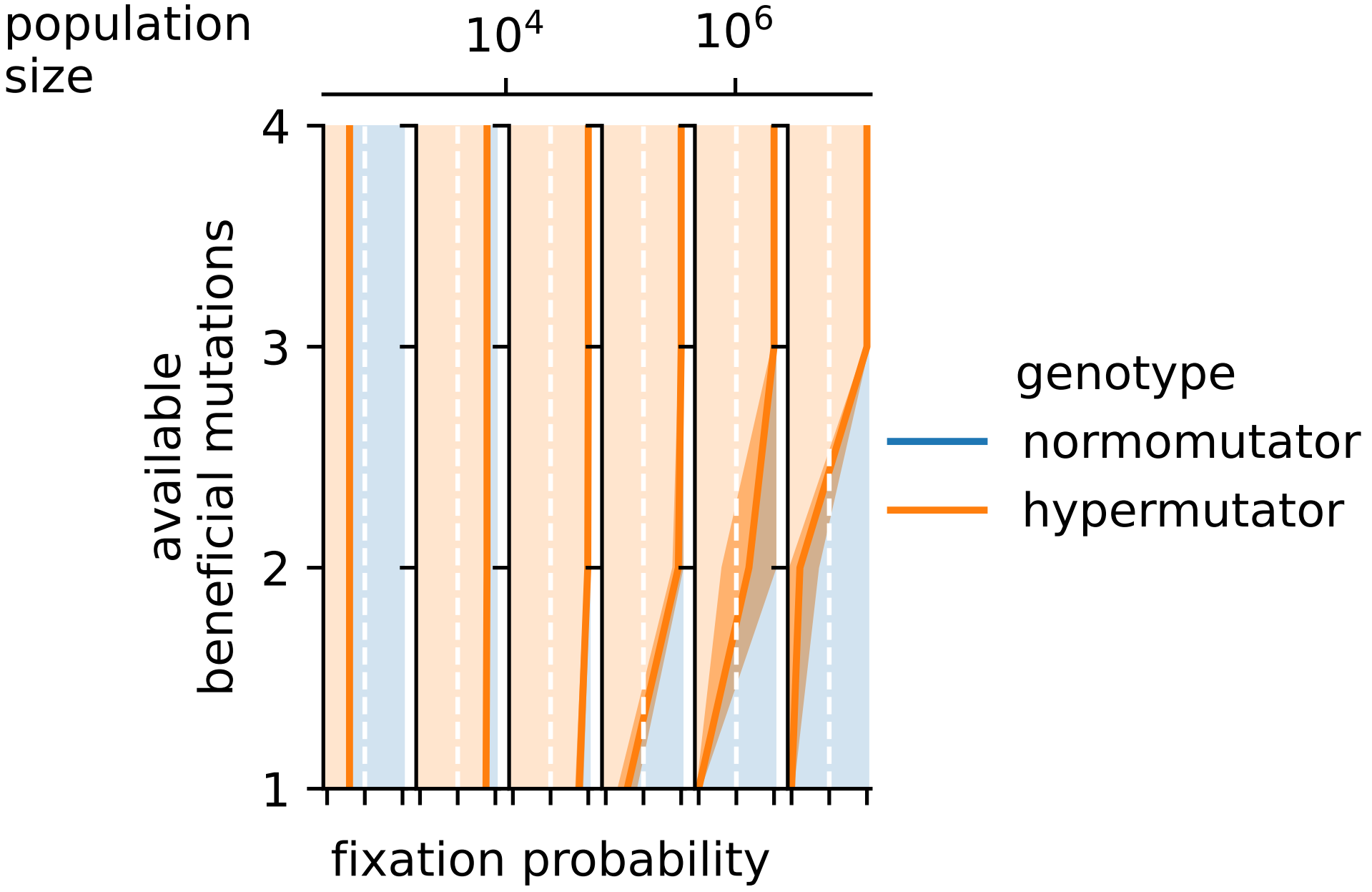
population size
10^4
10^6
4
3
2
1
available beneficial mutations
fixation probability
genotype
normomutator
hypermutator