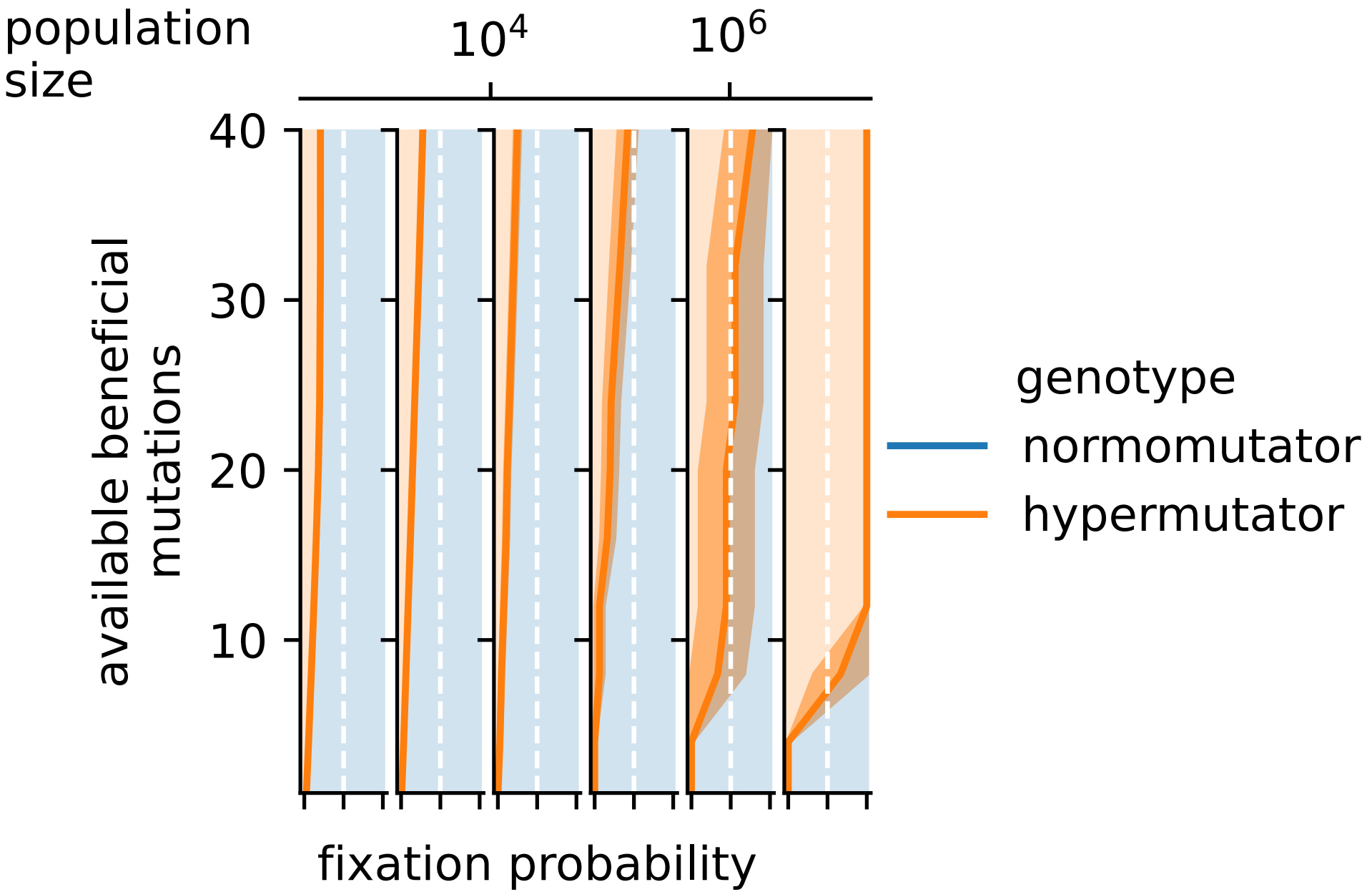
population size
10^4
10^6
40
30
20
10
available beneficial mutations
fixation probability
genotype
normomutator
hypermutator