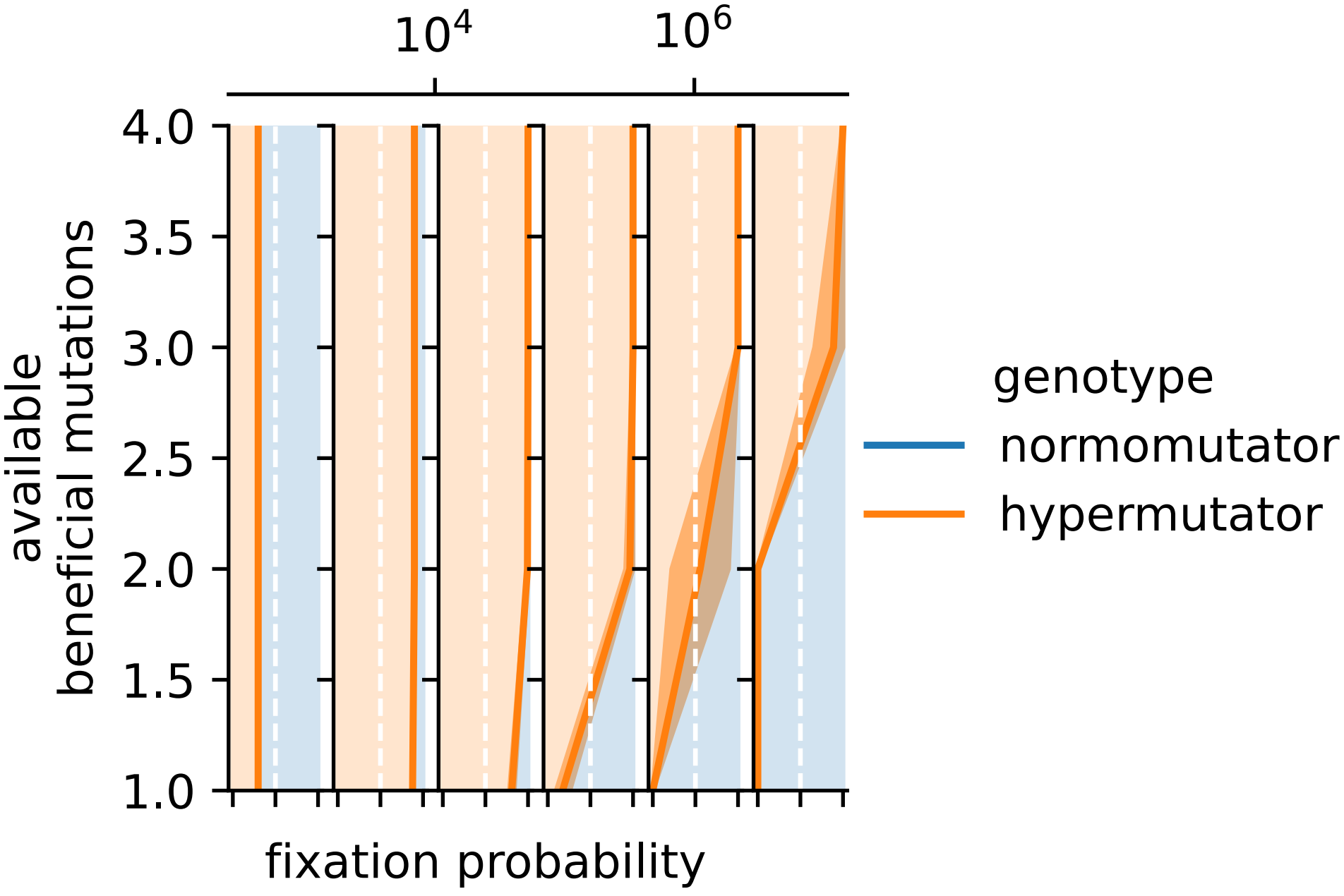
10^4
10^6
4.0
3.5
3.0
2.5
2.0
1.5
1.0
available beneficial mutations
fixation probability
genotype
normomutator
hypermutator