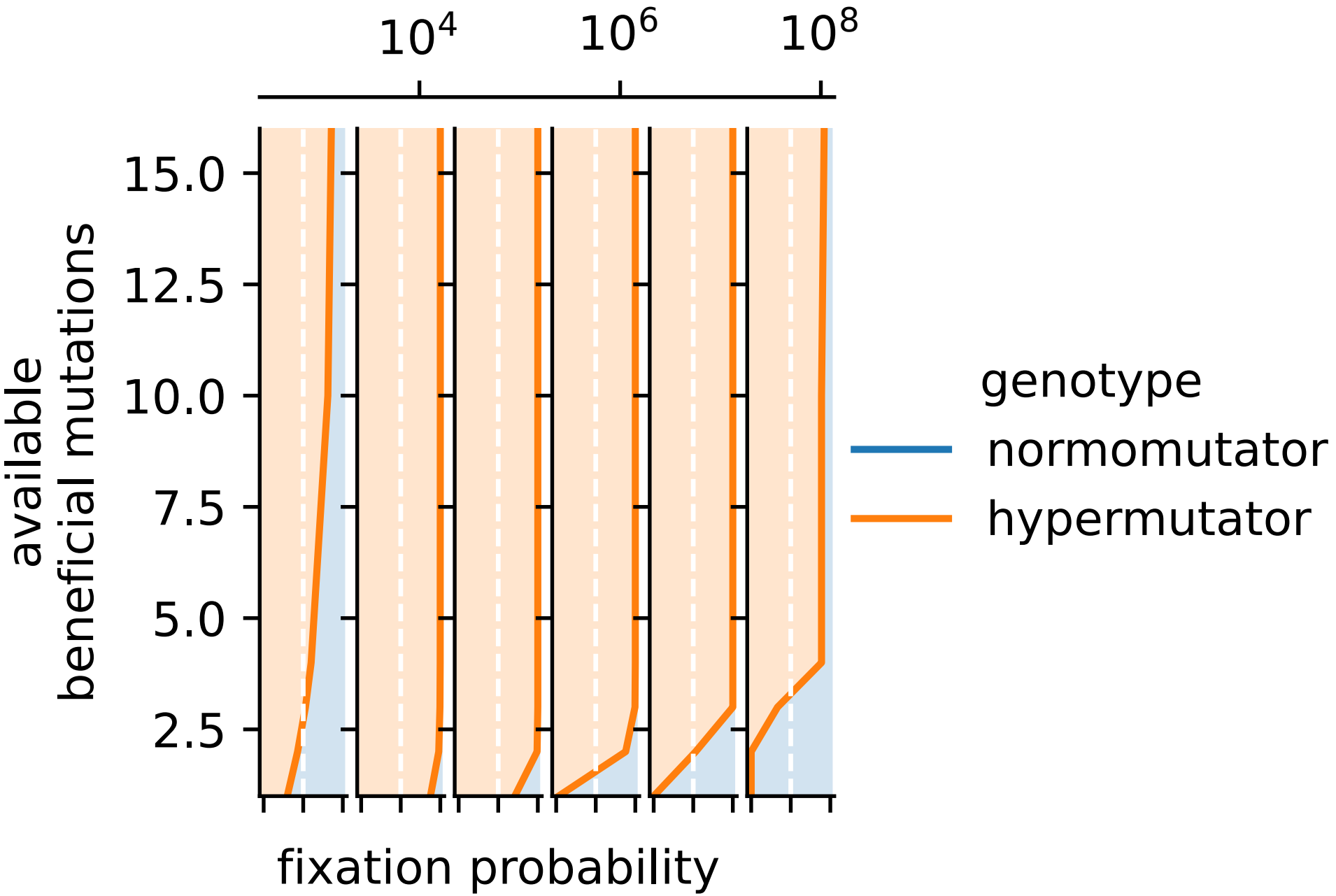
10^4
10^6
10^8
15.0
12.5
10.0
7.5
5.0
2.5
available beneficial mutations
fixation probability
genotype
normomutator
hypermutator