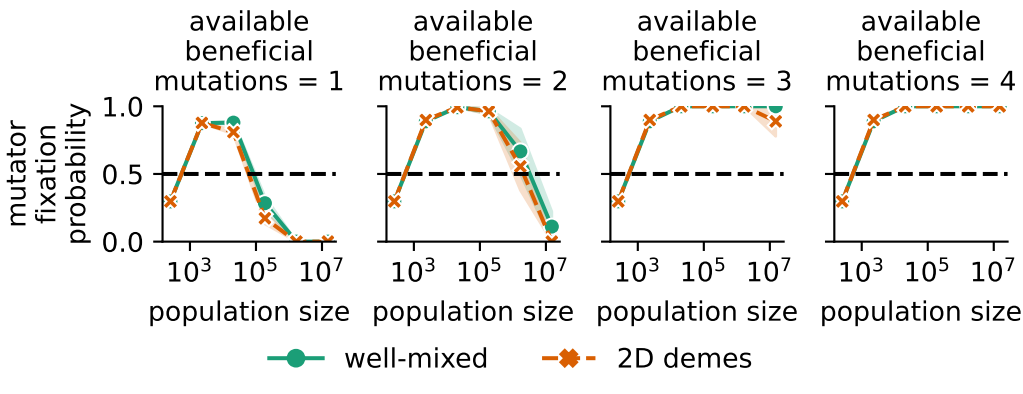
available beneficial mutations = 1
available beneficial mutations = 2
available beneficial mutations = 3
available beneficial mutations = 4
mutator fixation probability
1.0
0.5
0.0
10^3
10^5
10^7
population size
well-mixed
2D demes