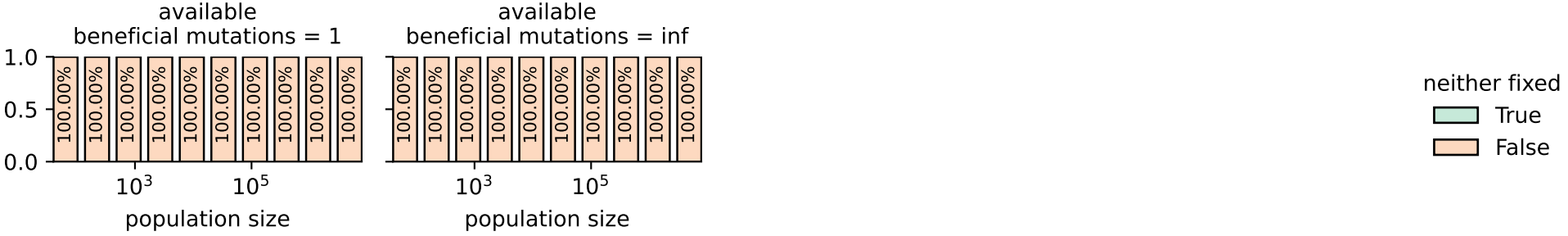
available
beneficial mutations = 1
available
beneficial mutations = inf
1.0
0.5
0.0
100.00%
10^3
10^5
population size
neither fixed
True
False