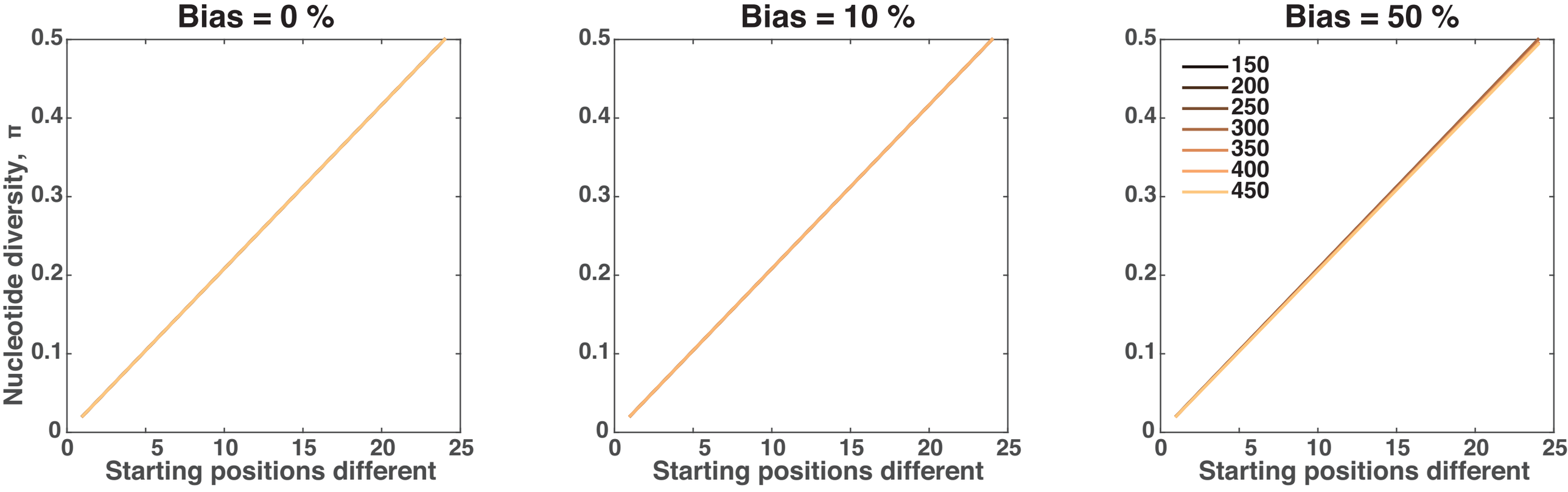
Bias = 0 %
Bias = 10 %
Bias = 50 %
Nucleotide diversity, π
Starting positions different
Starting positions different
Starting positions different
0
0.1
0.2
0.3
0.4
0.5
0
5
10
15
20
25
150
200
250
300
350
400
450