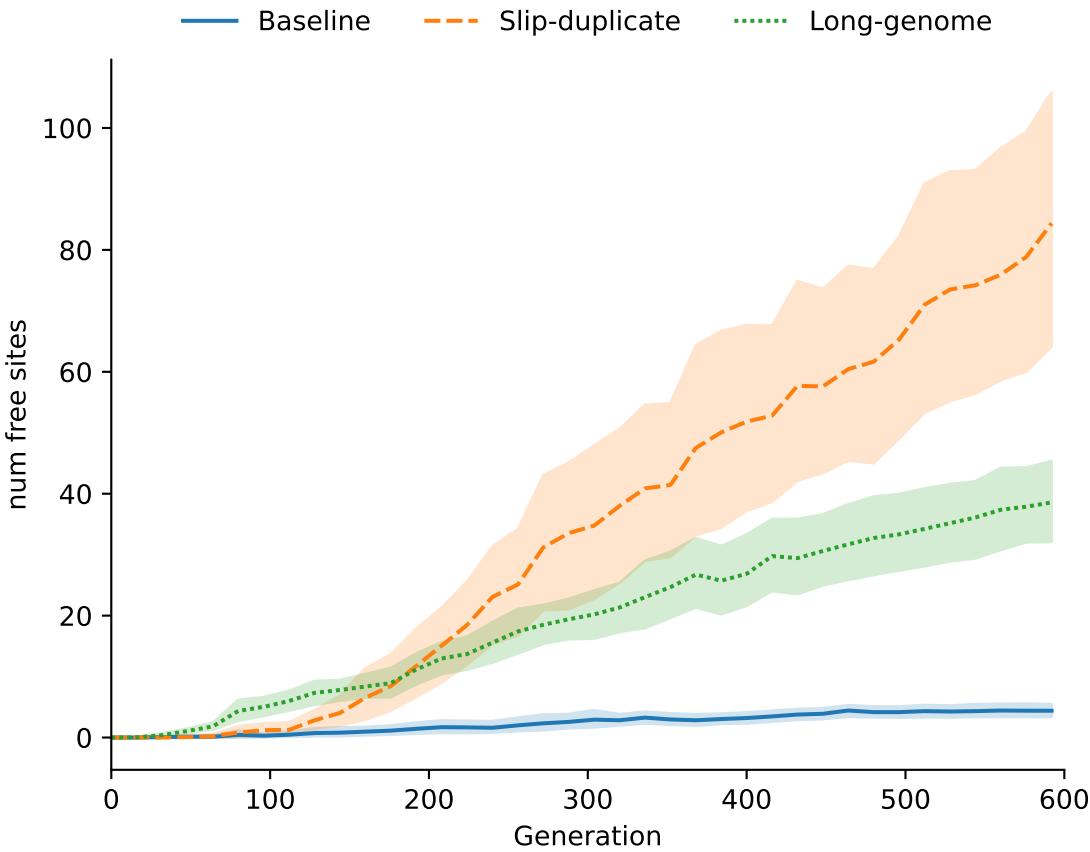
Baseline
Slip-duplicate
Long-genome
num free sites
0
20
40
60
80
100
0
100
200
300
400
500
600
Generation