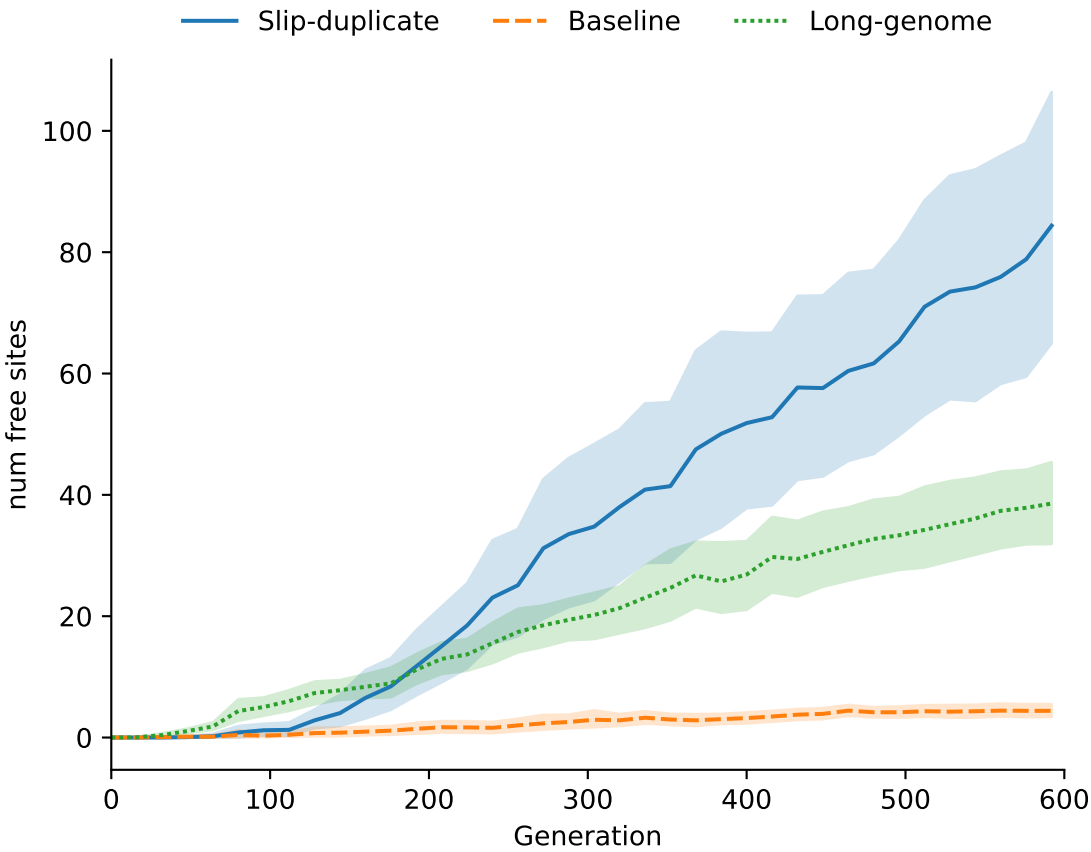
Slip-duplicate
Baseline
Long-genome
num free sites
Generation
0
20
40
60
80
100
0
100
200
300
400
500
600