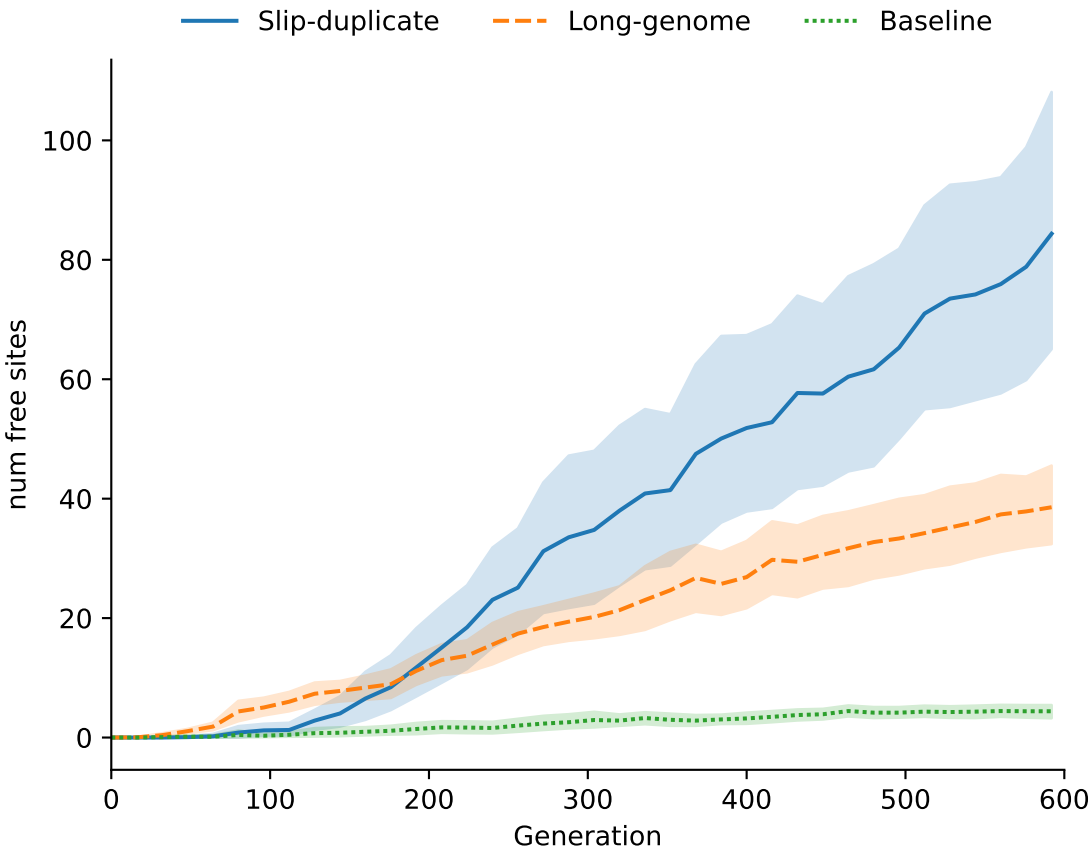
Slip-duplicate
Long-genome
Baseline
num free sites
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20
40
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100
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200
300
400
500
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Generation