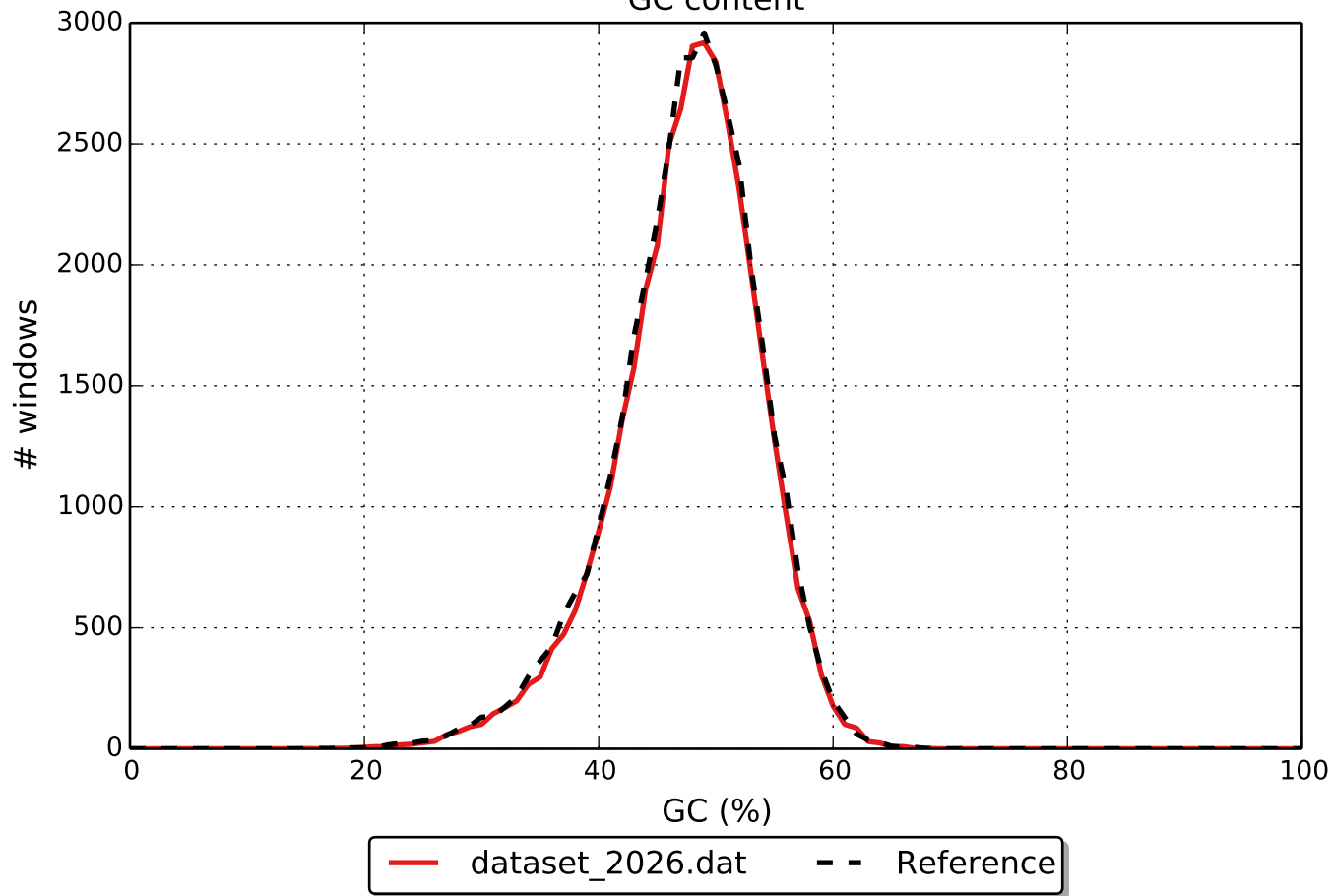
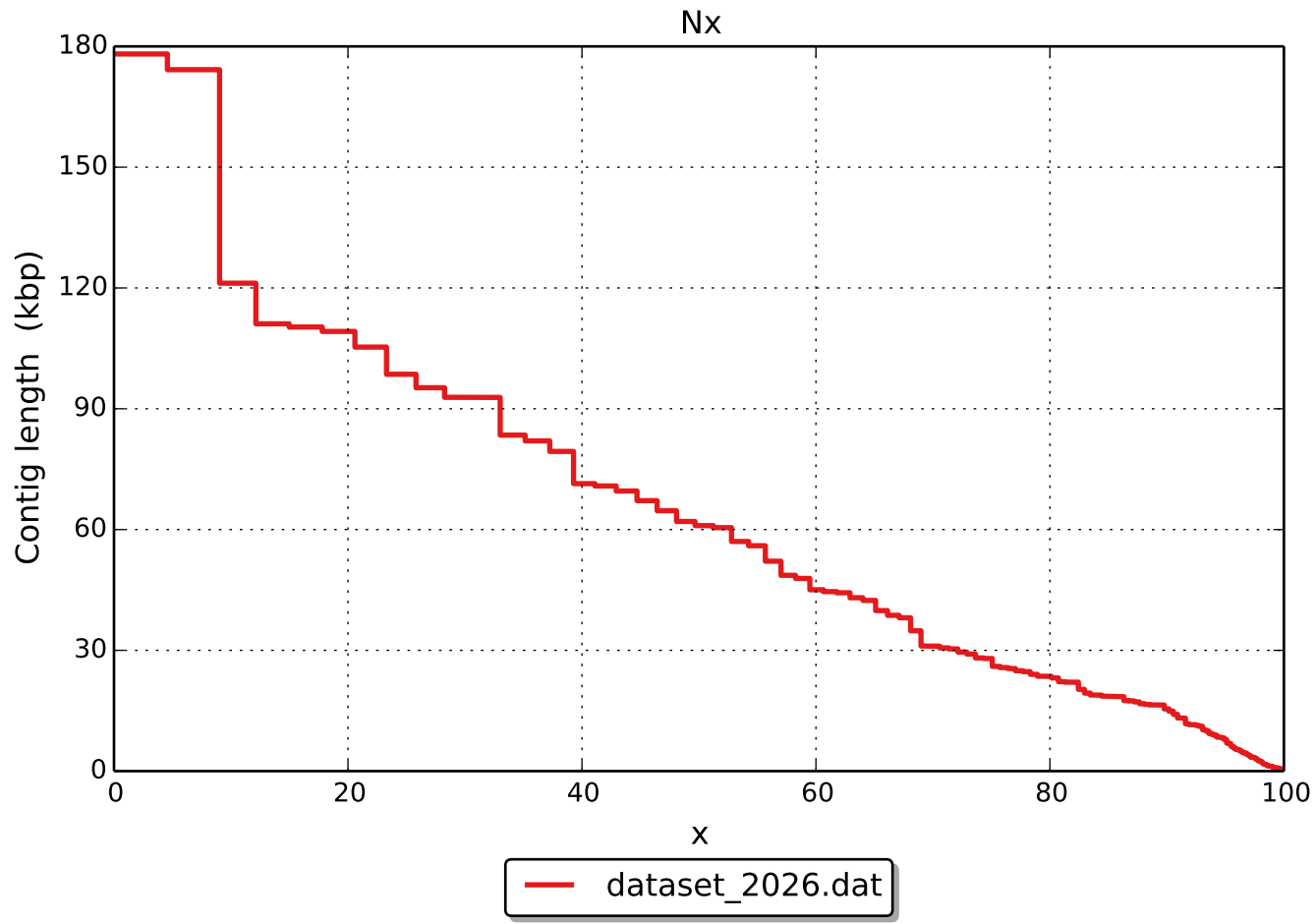
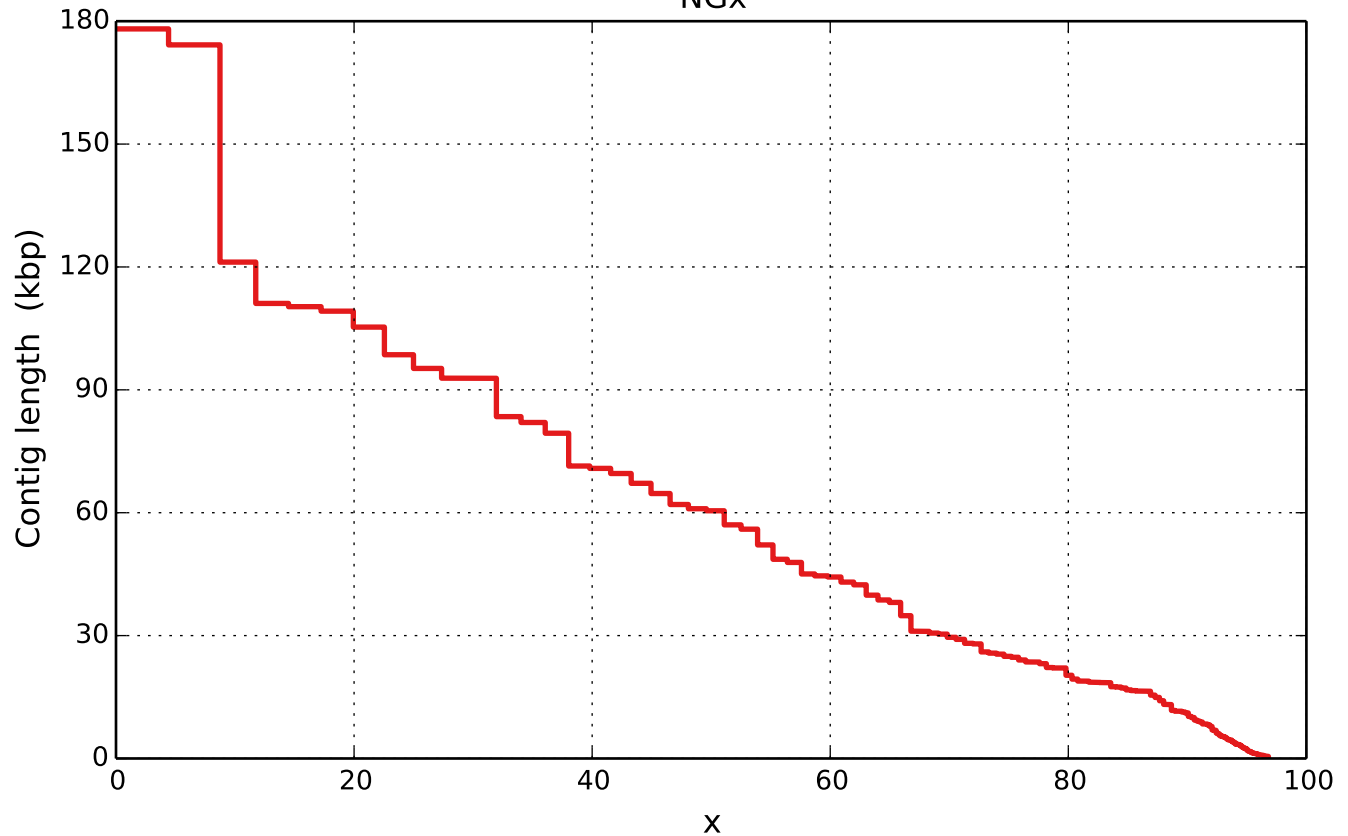


GC content



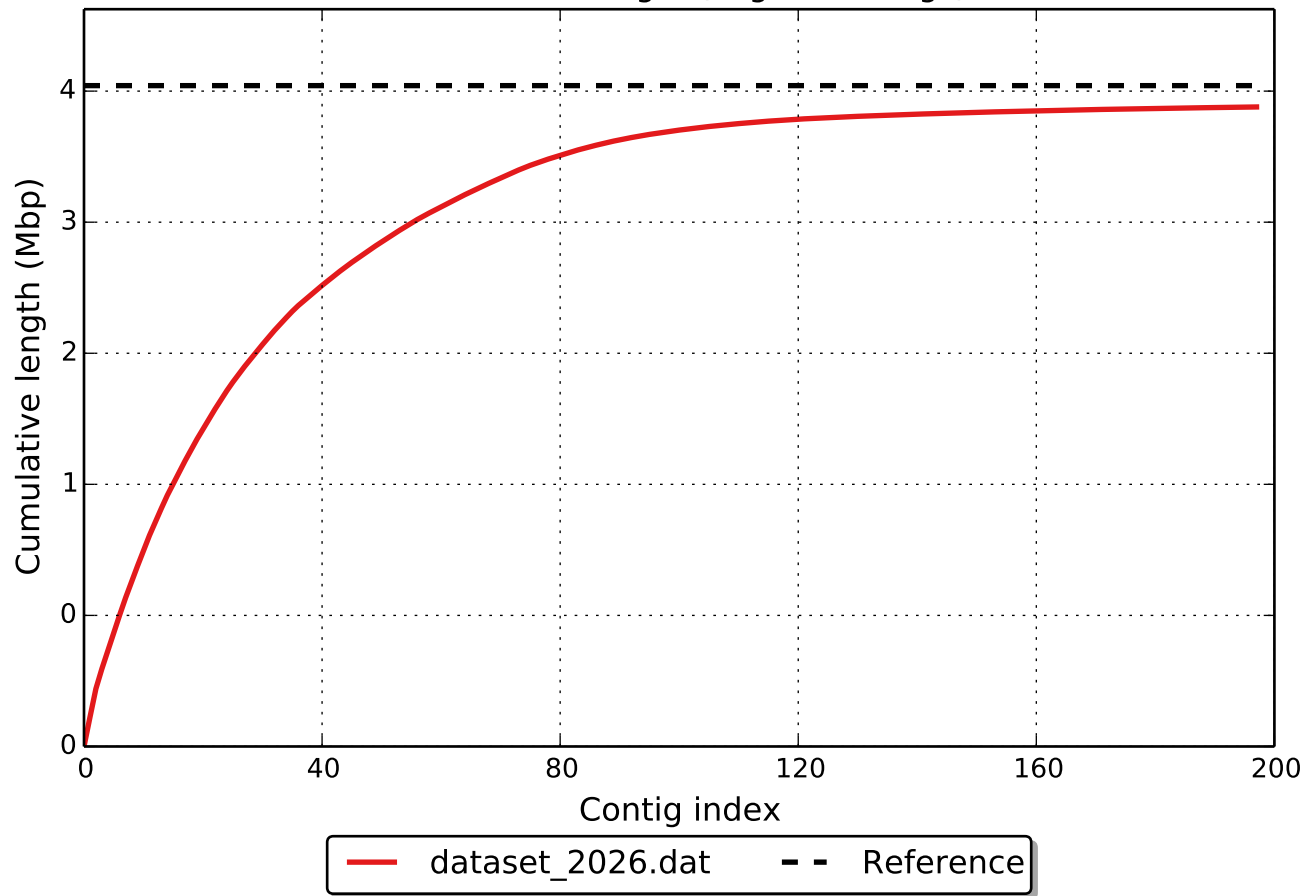


NGx

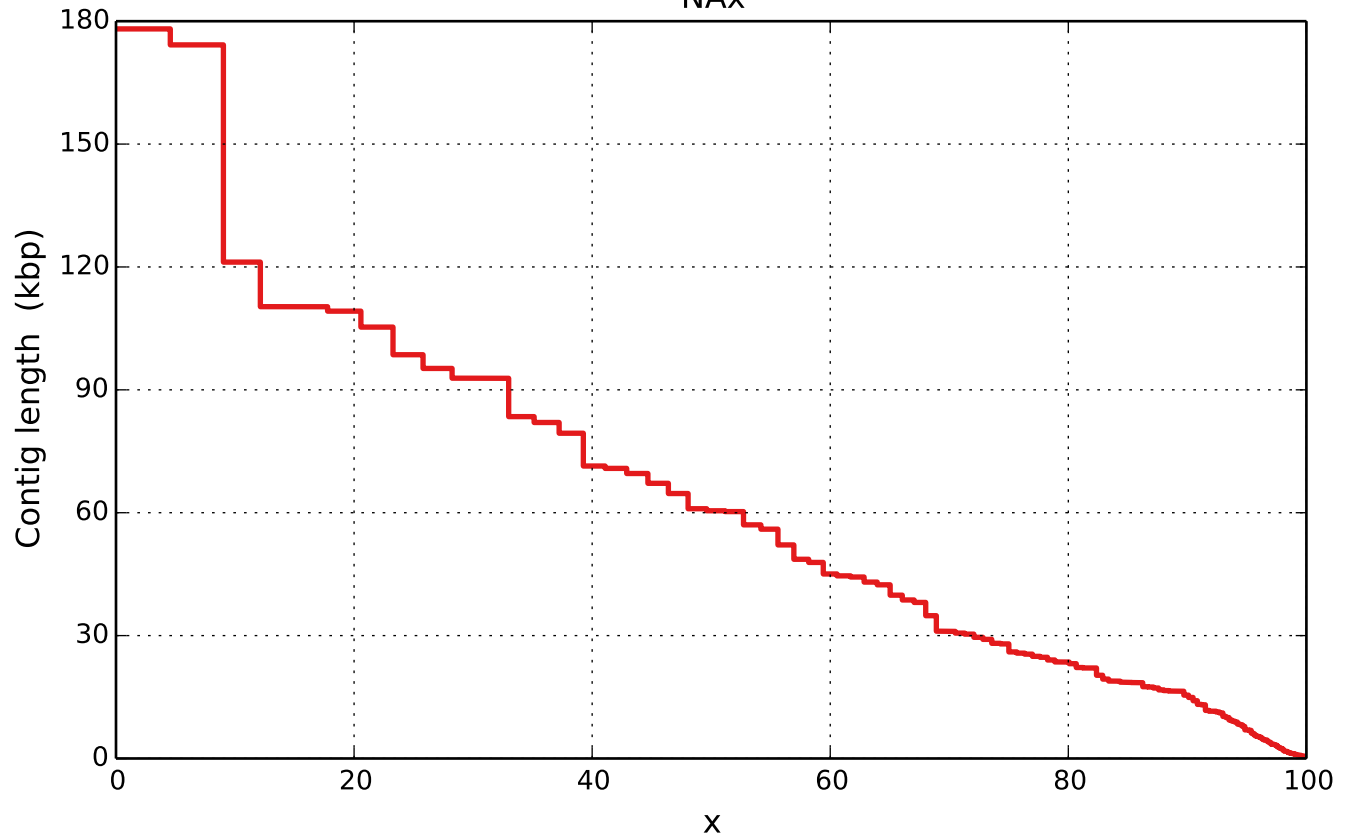


— dataset_2026.dat

Cumulative length (aligned contigs)

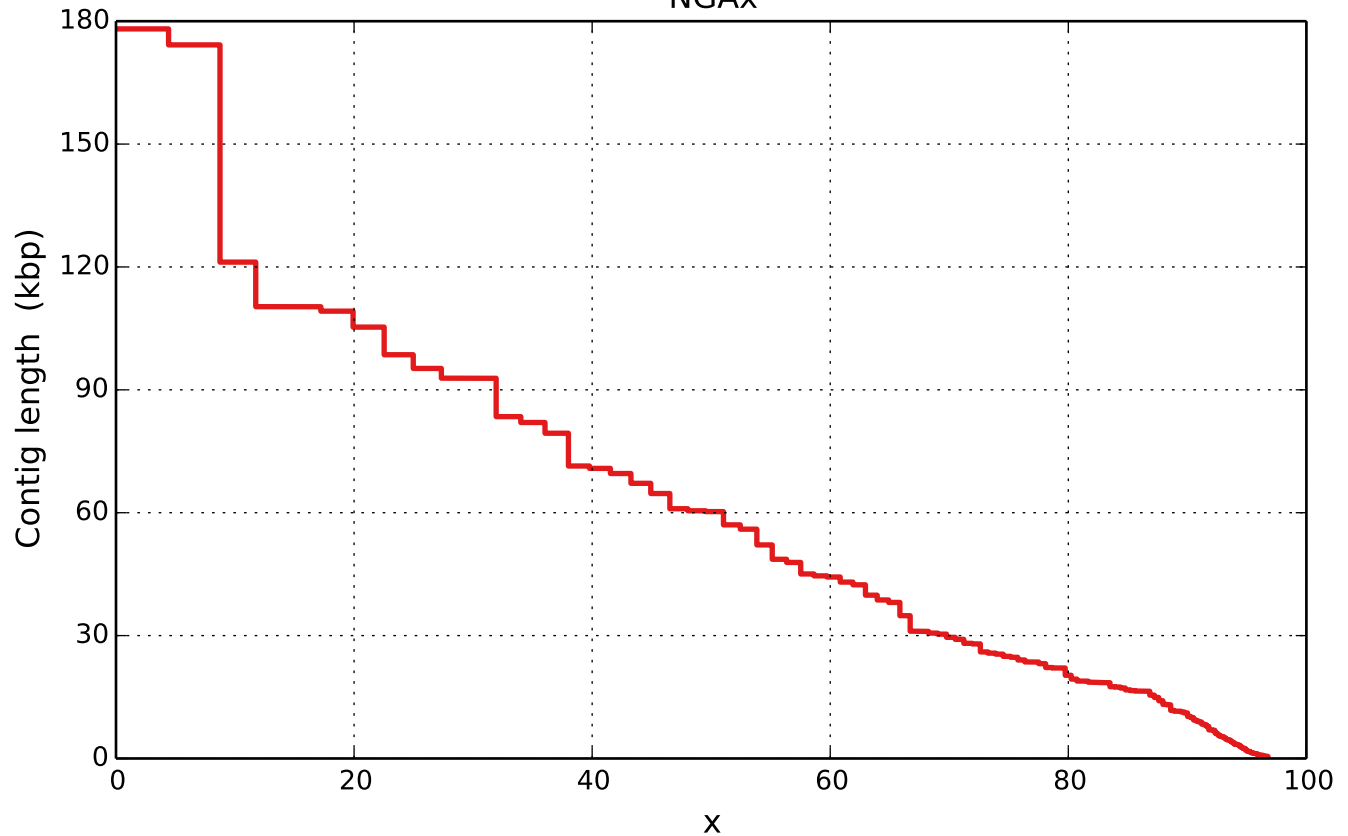


NAx



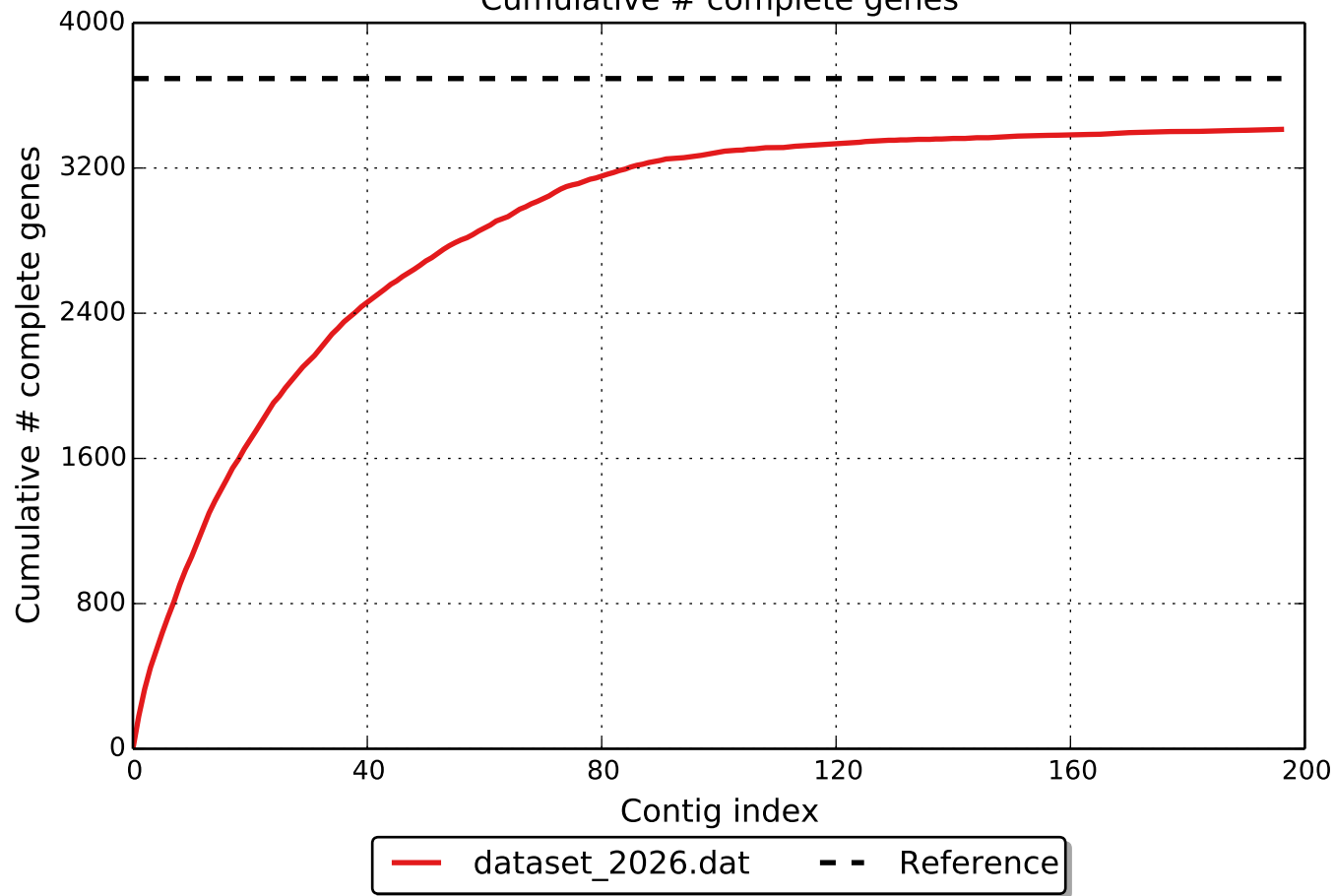
— dataset_2026.dat

NGAx

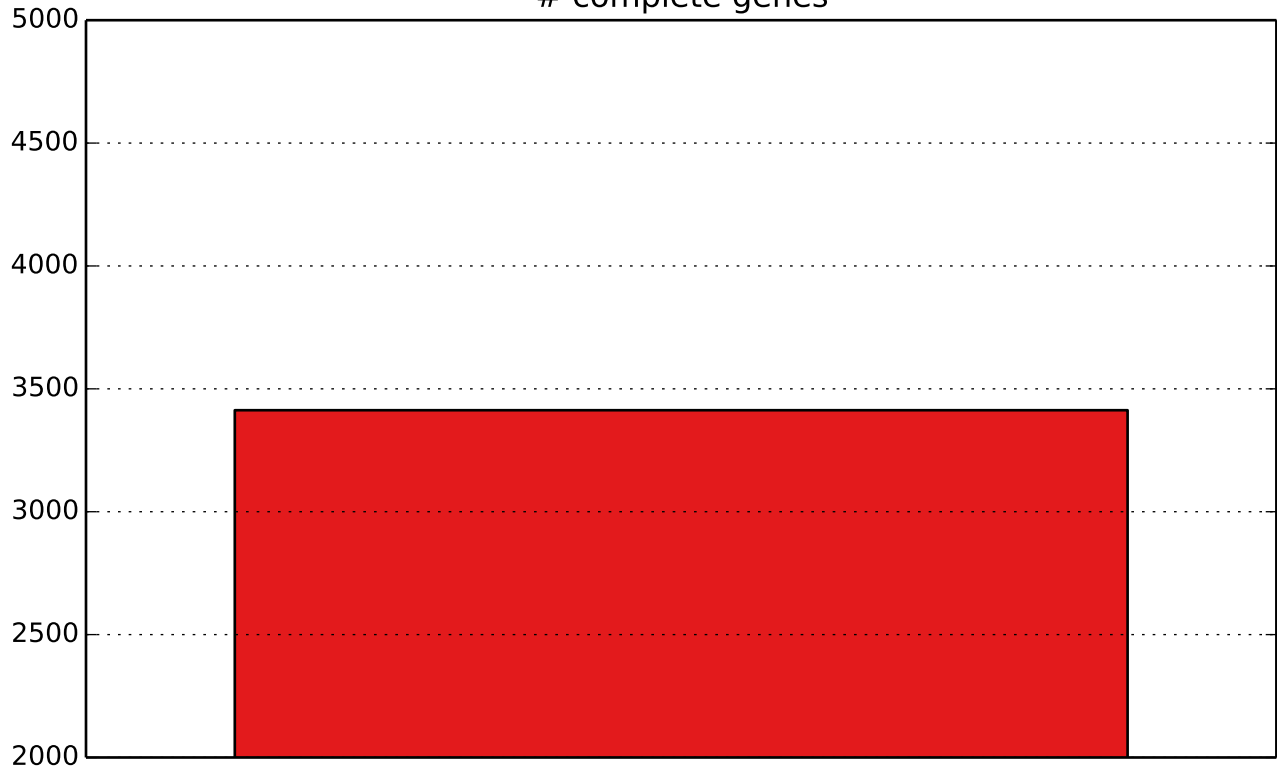


— dataset_2026.dat

Cumulative # complete genes



complete genes



dataset_2026.dat

