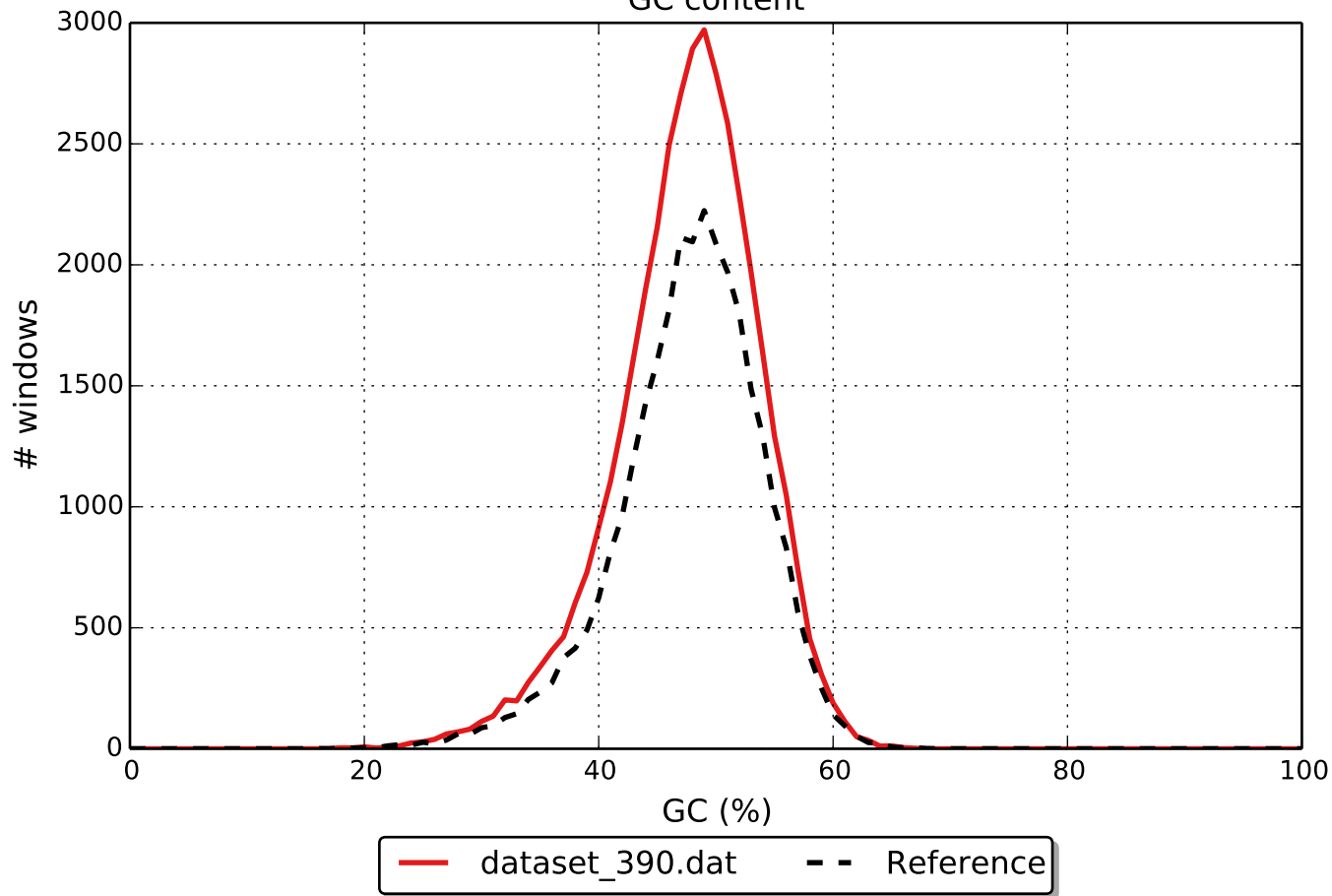
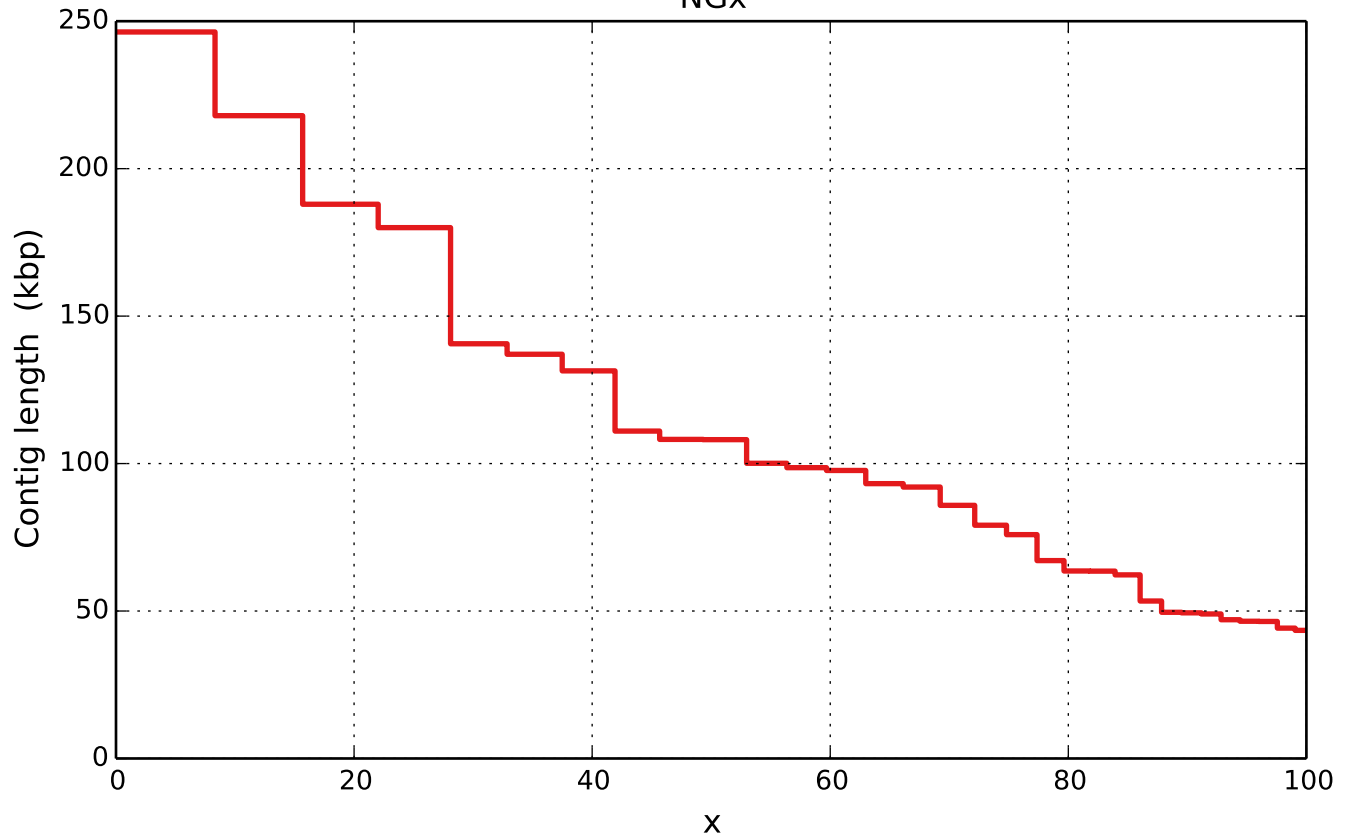


GC content



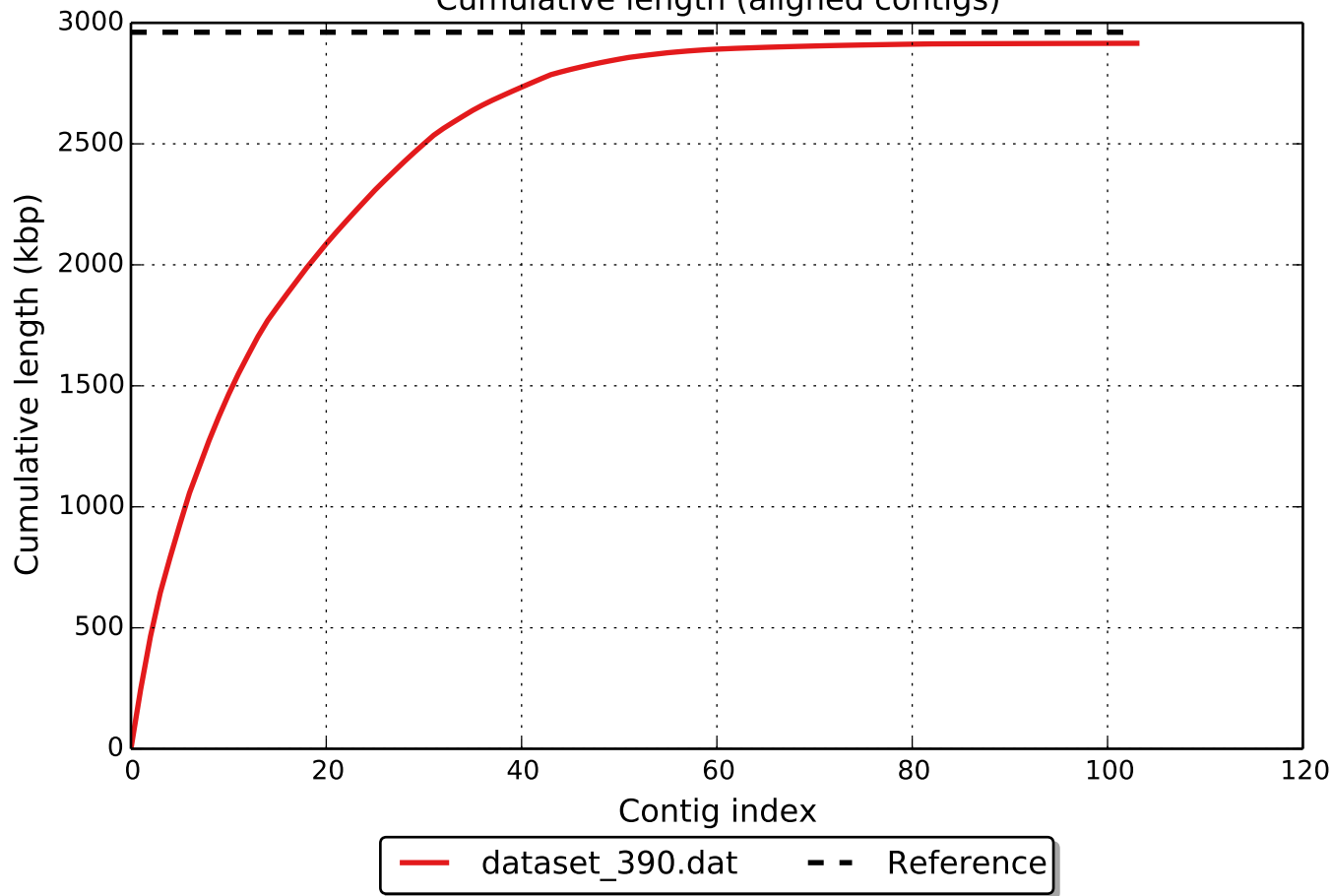


NGx

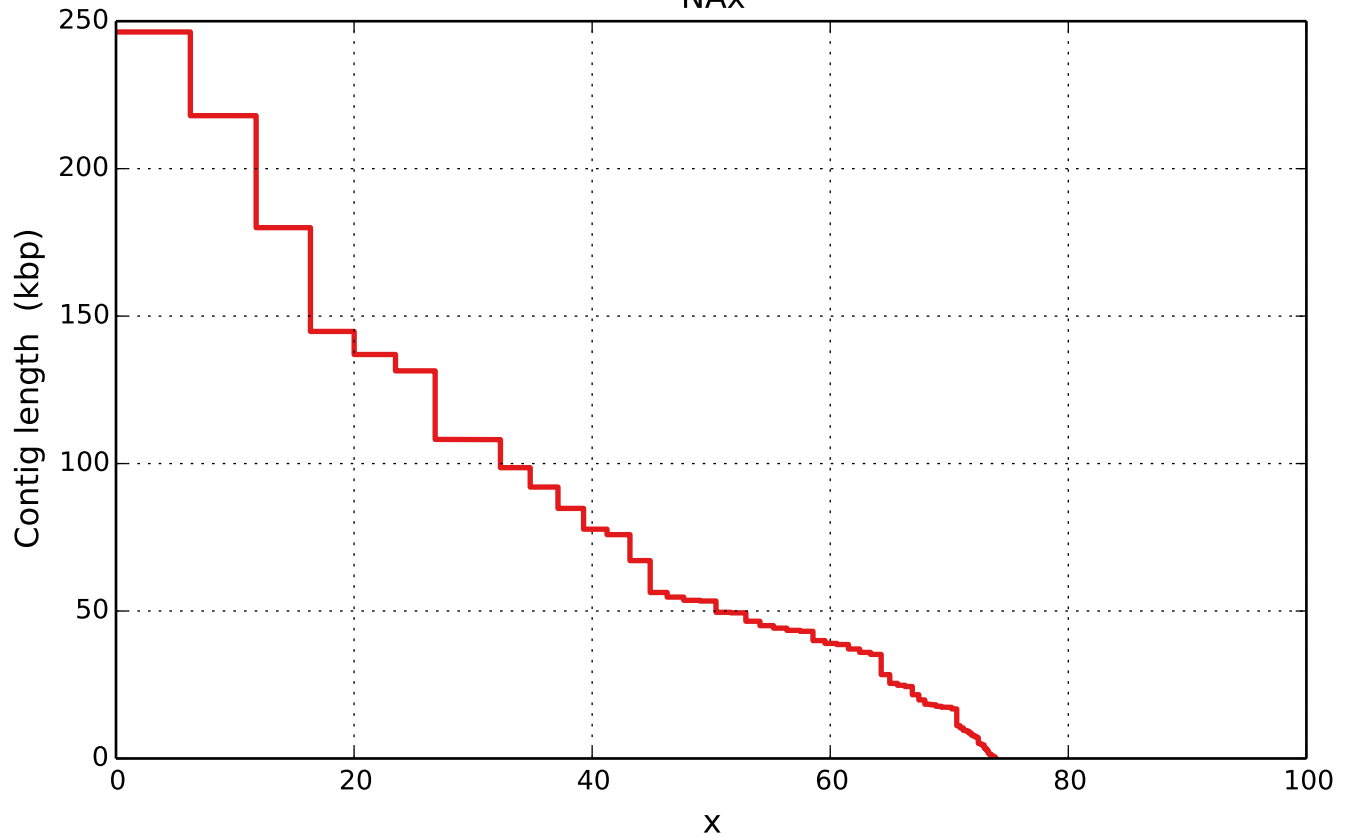


dataset_390.dat

Cumulative length (aligned contigs)

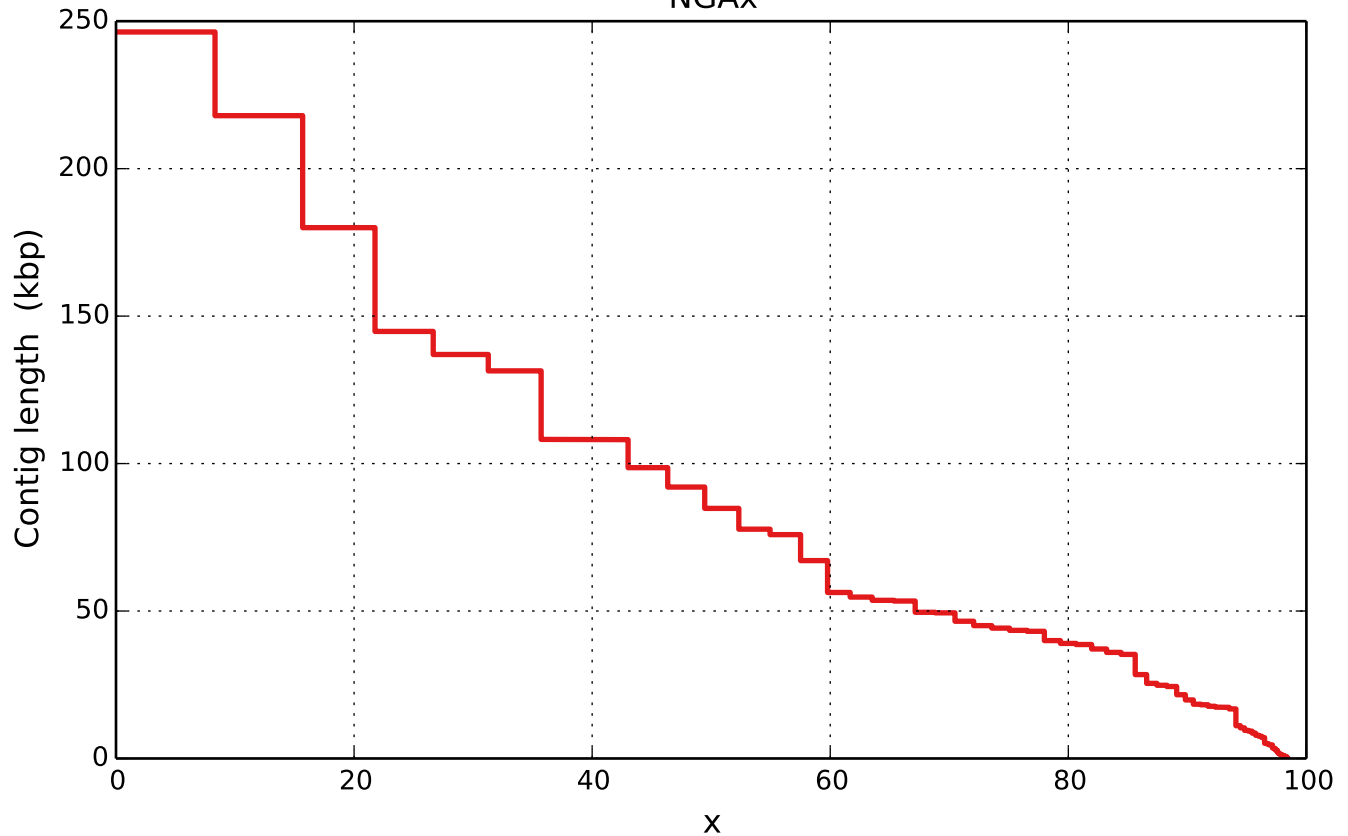


NAx



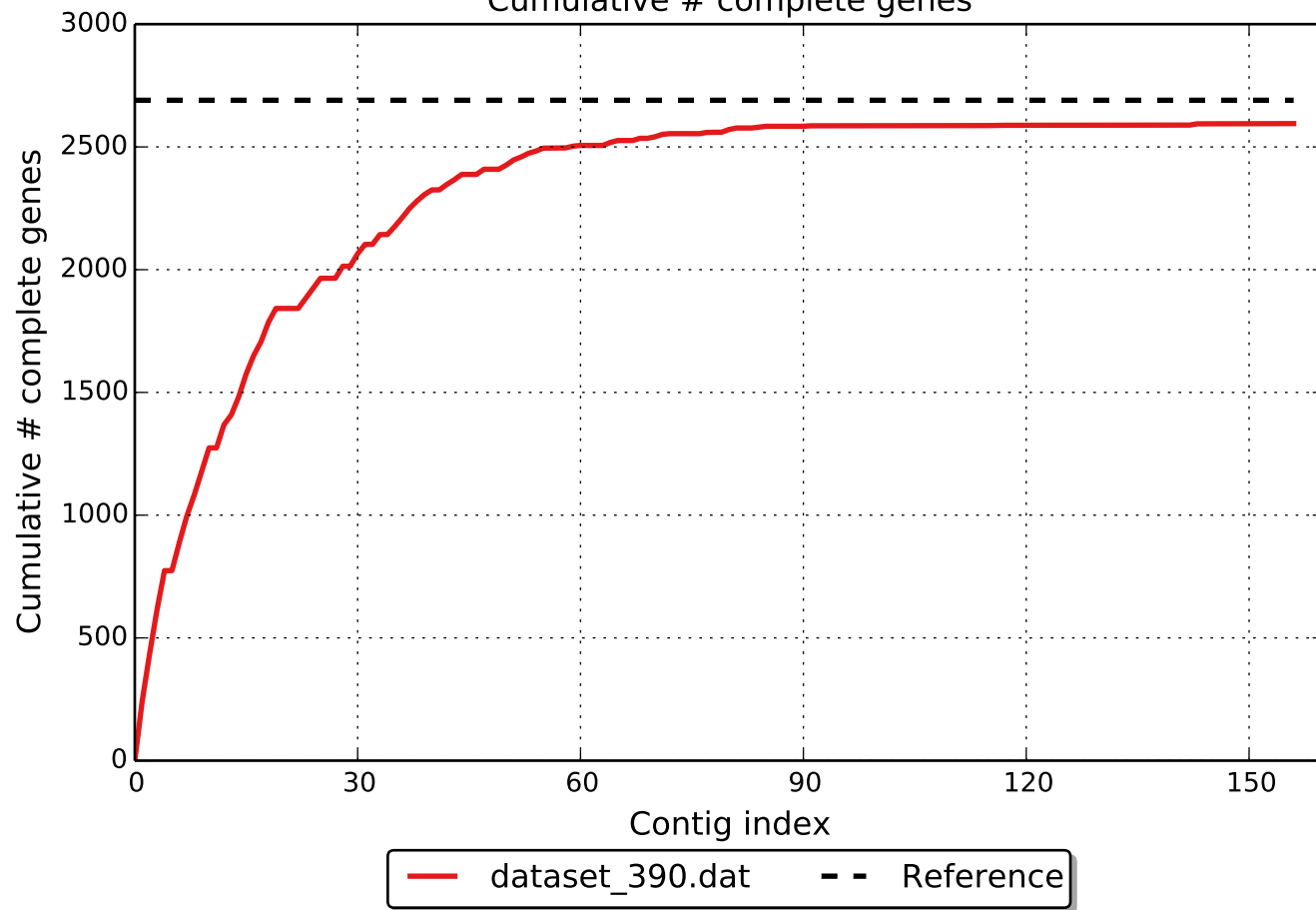
— dataset_390.dat

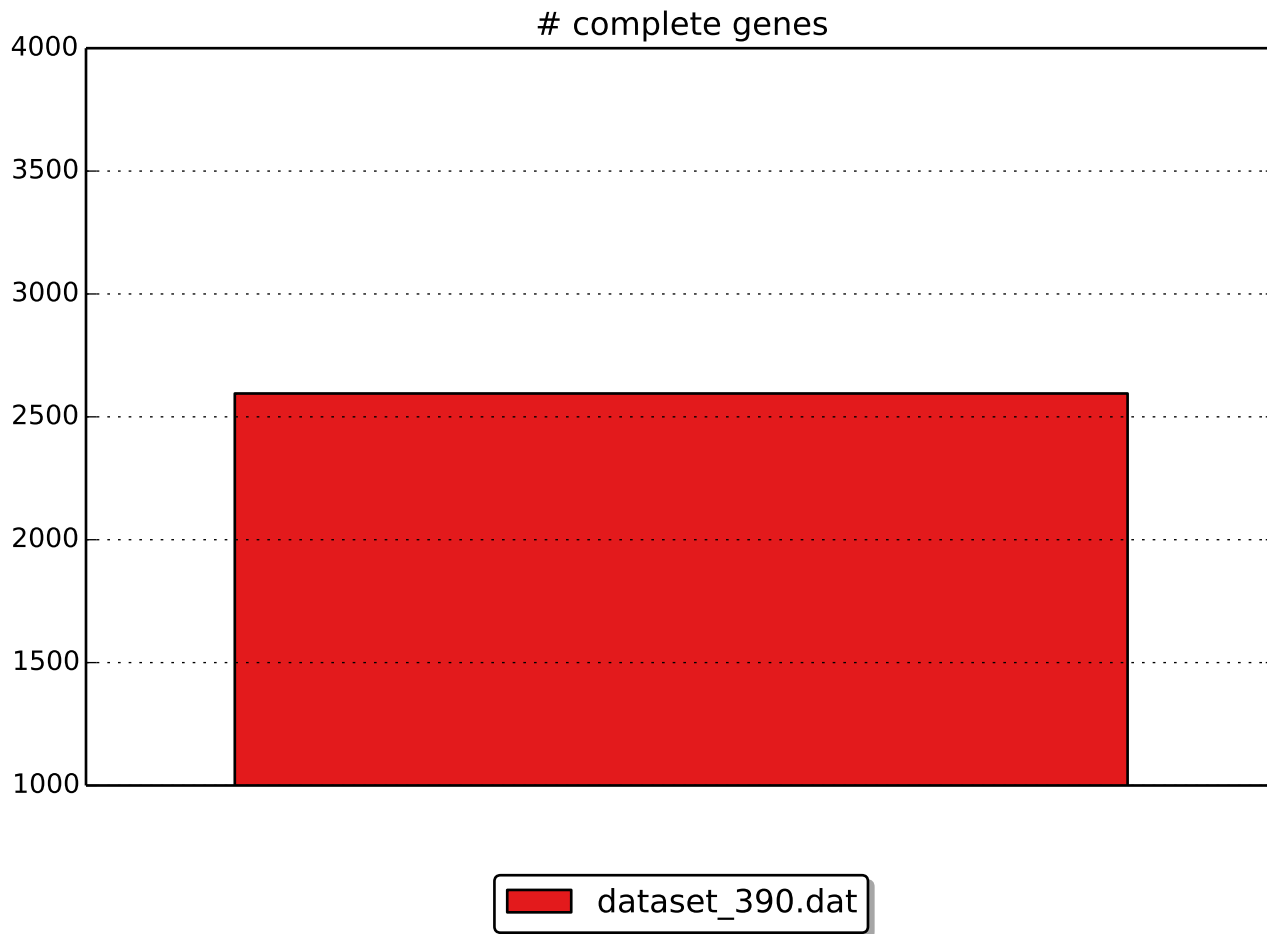
NGAx

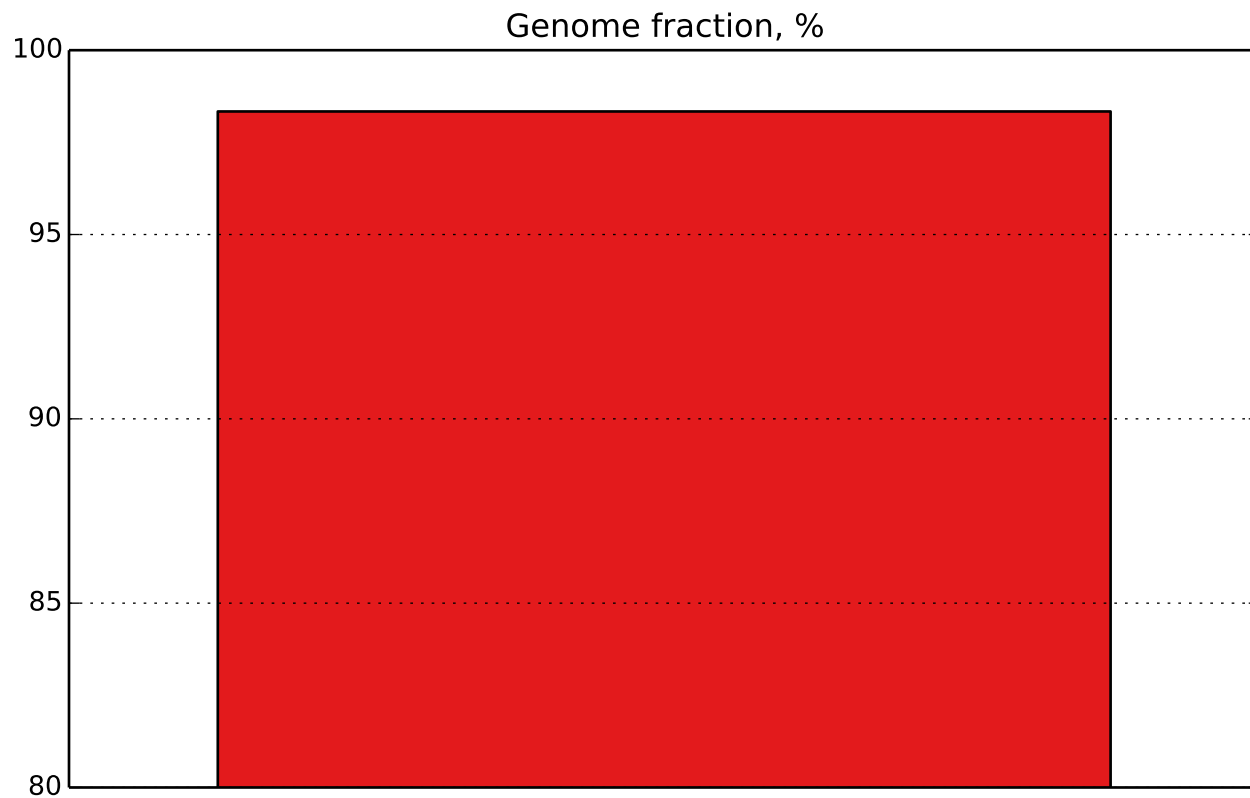


dataset_390.dat

Cumulative # complete genes







dataset_390.dat