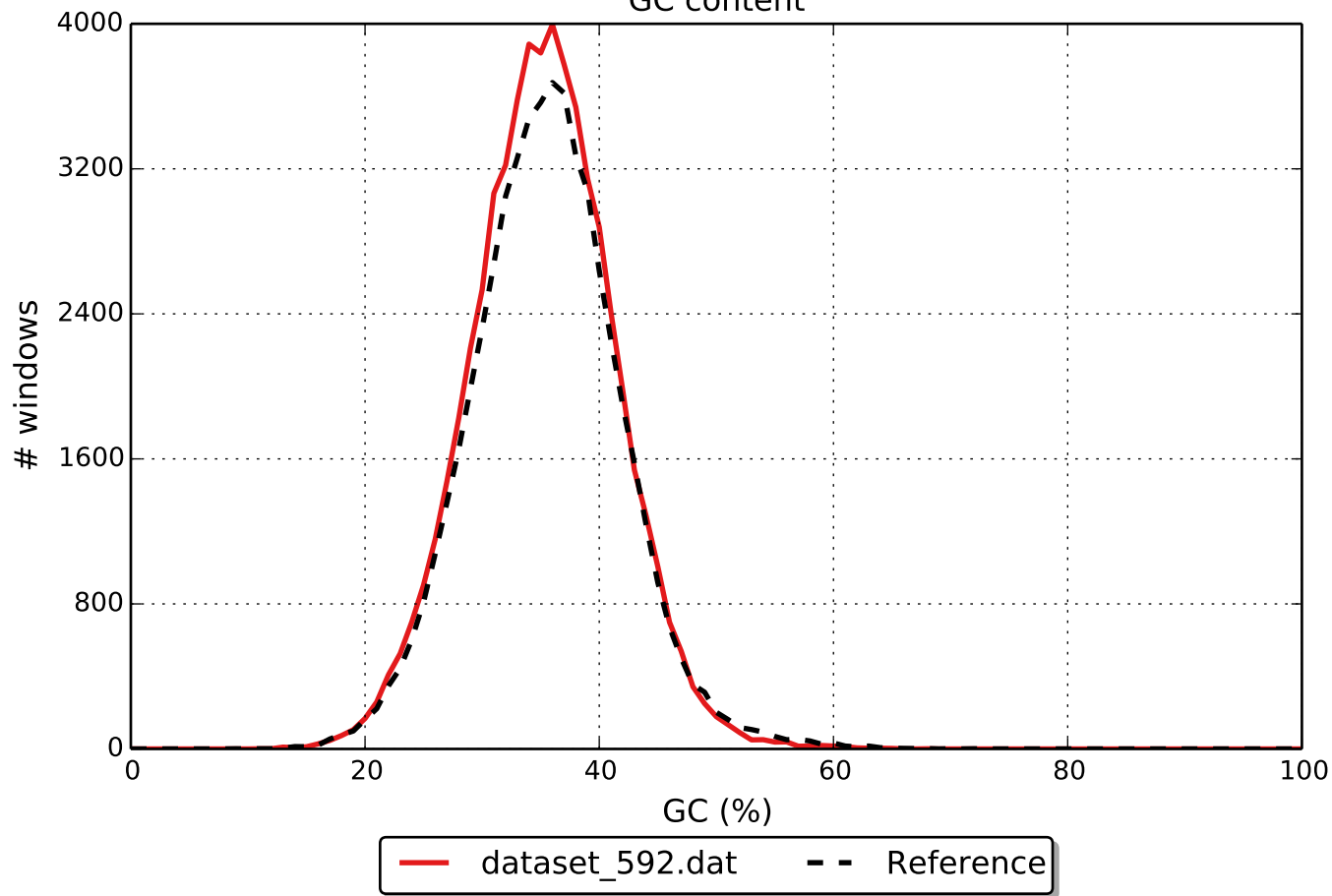




GC content

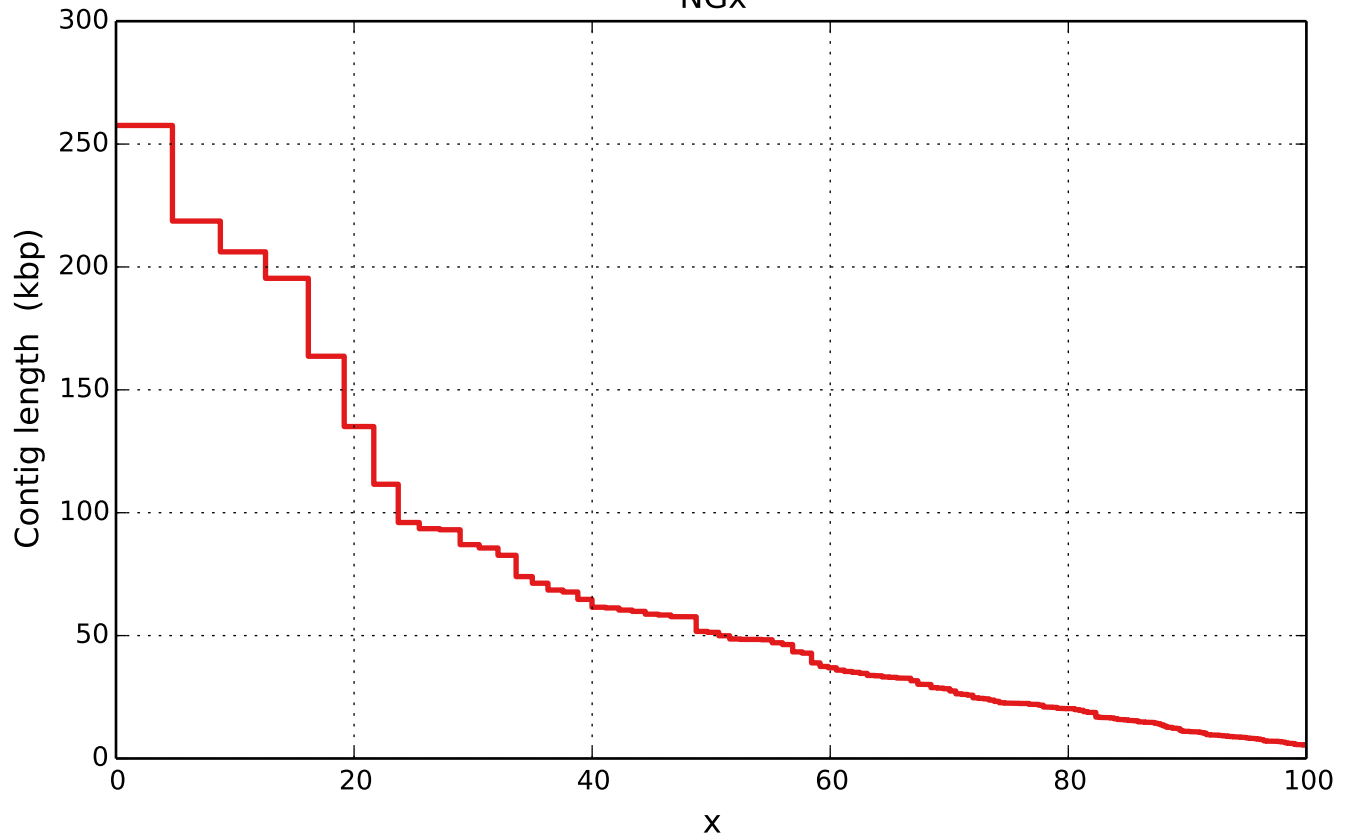


Nx



dataset_592.dat

NGx



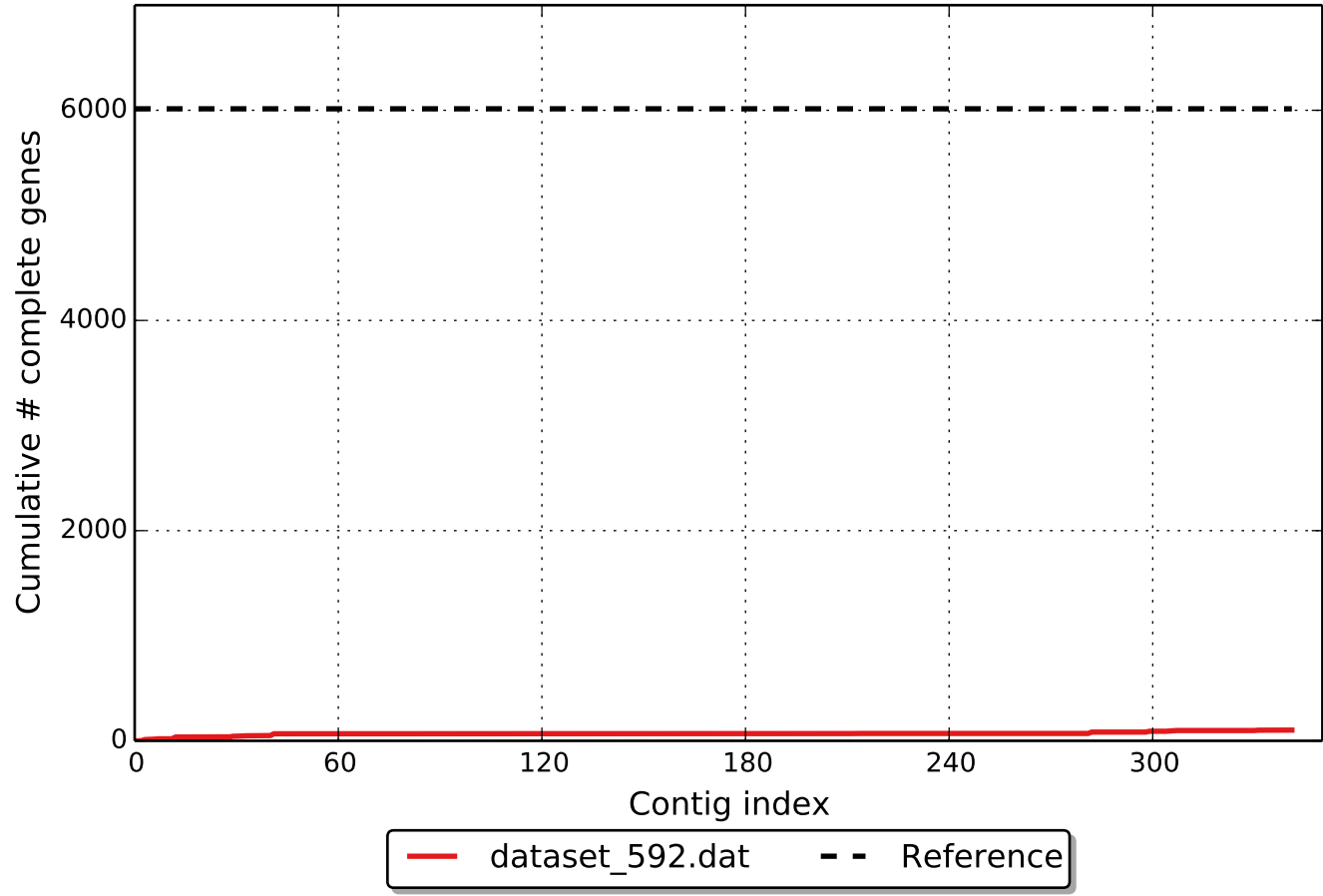
— dataset_592.dat

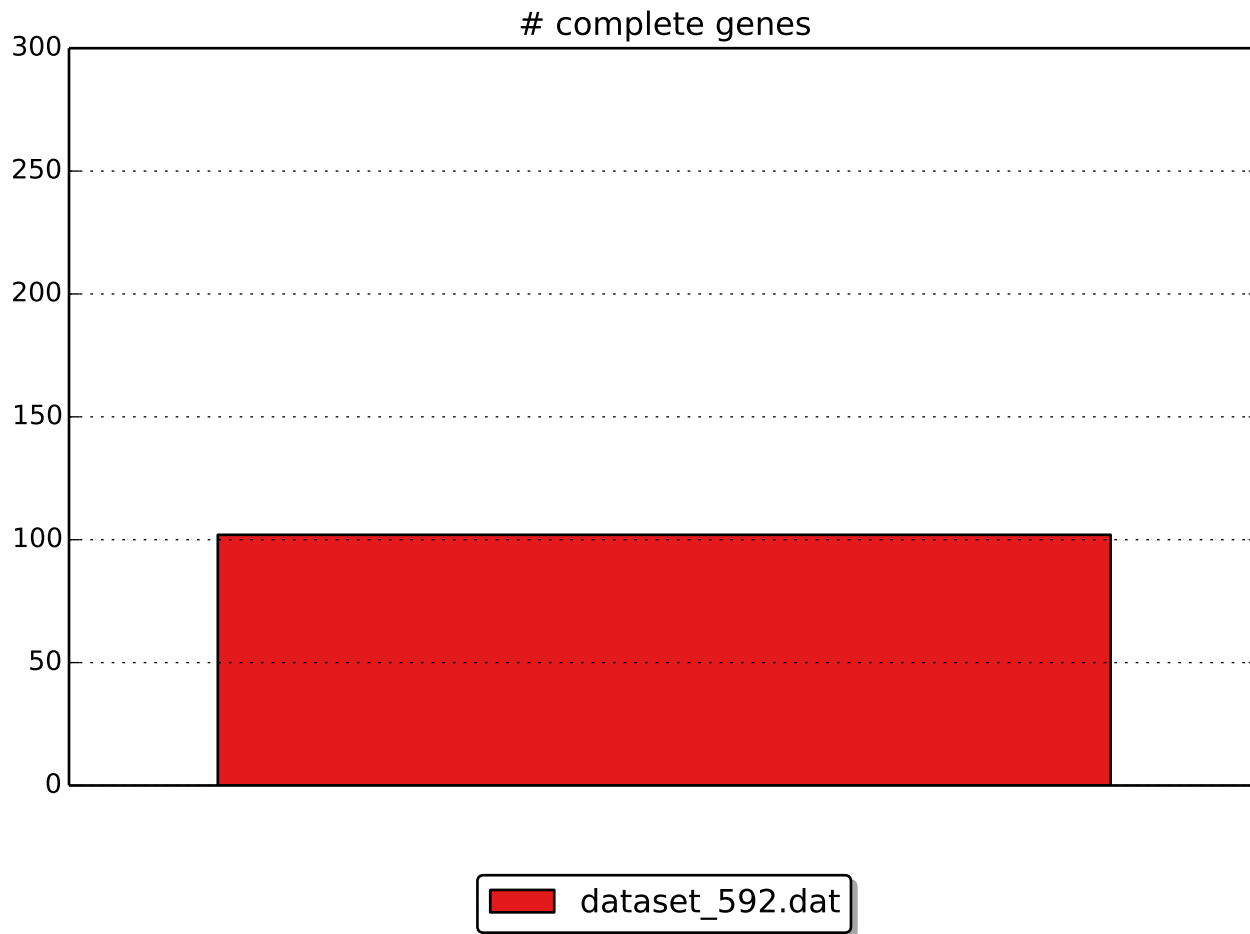


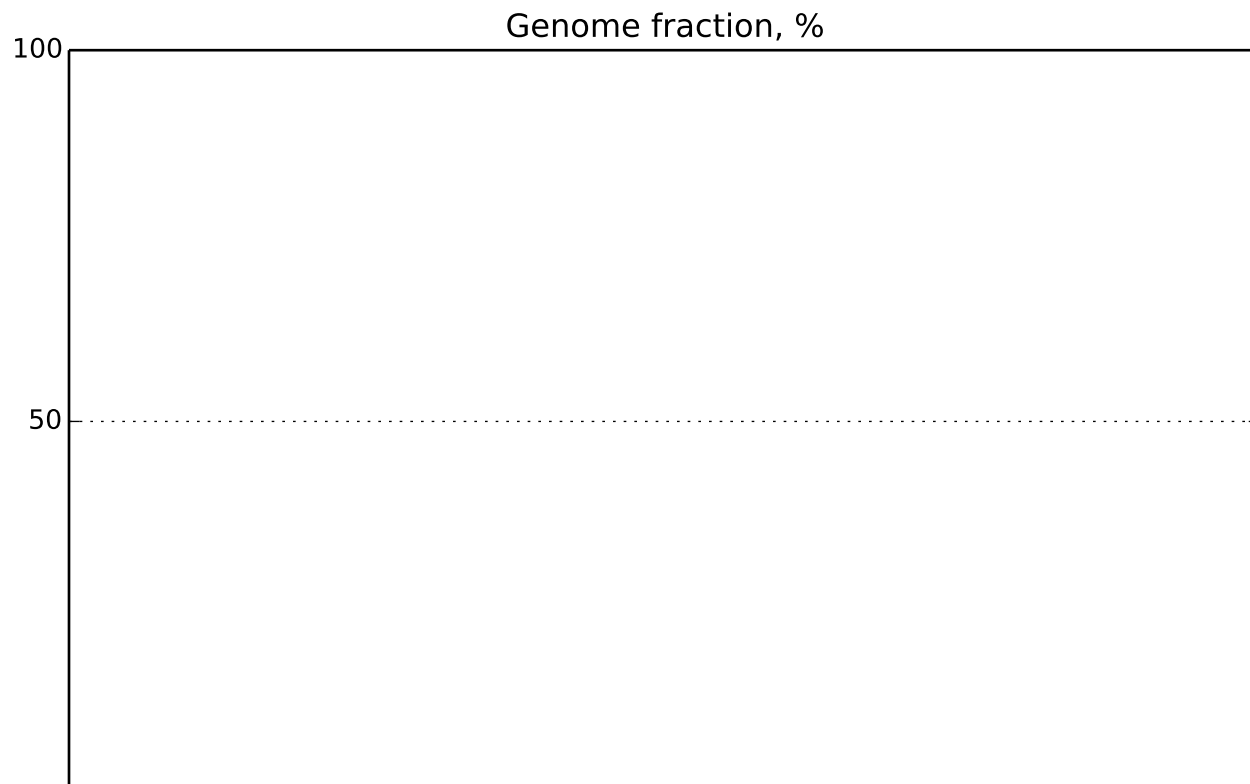




Cumulative # complete genes







 dataset_592.dat