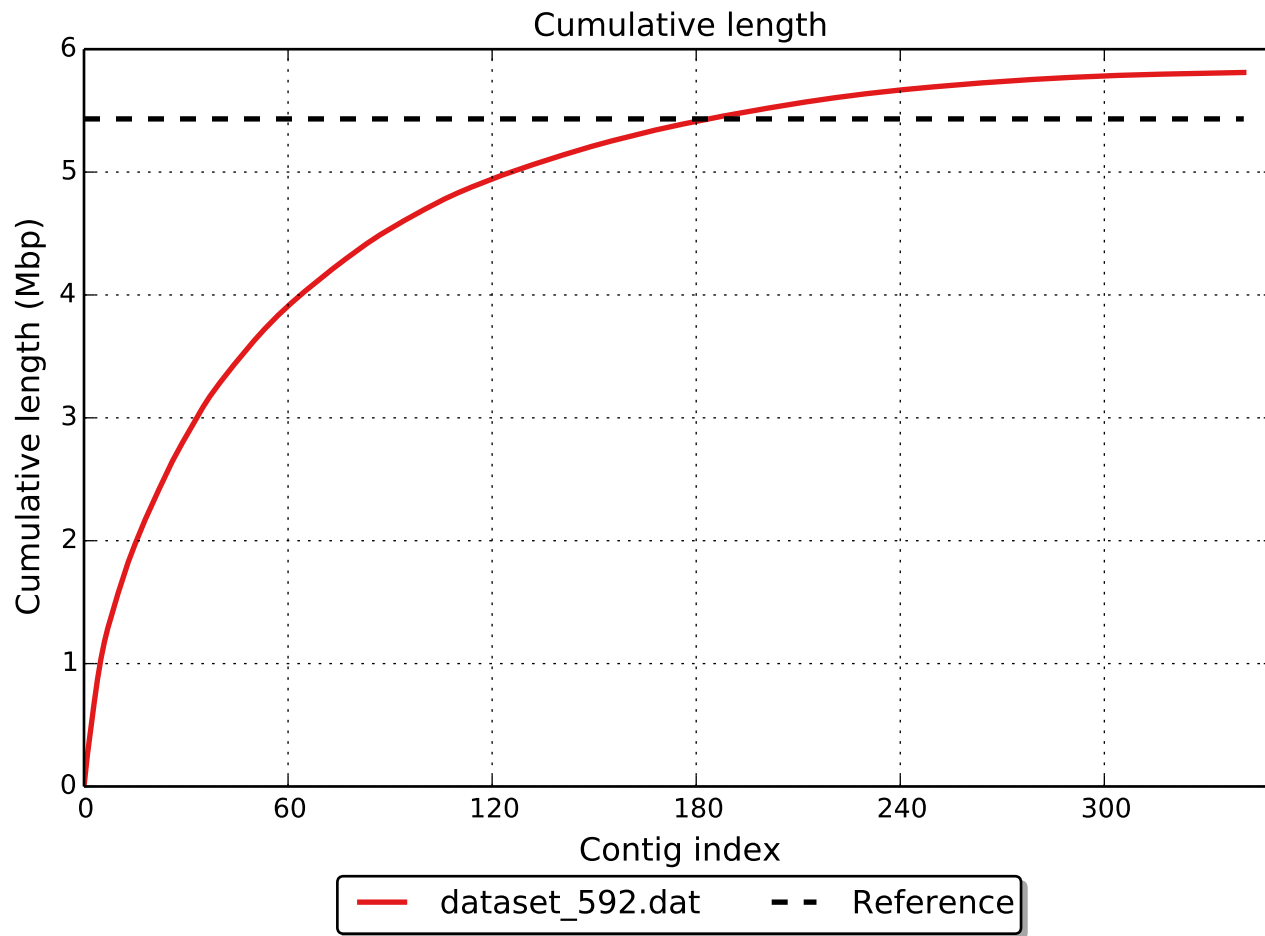
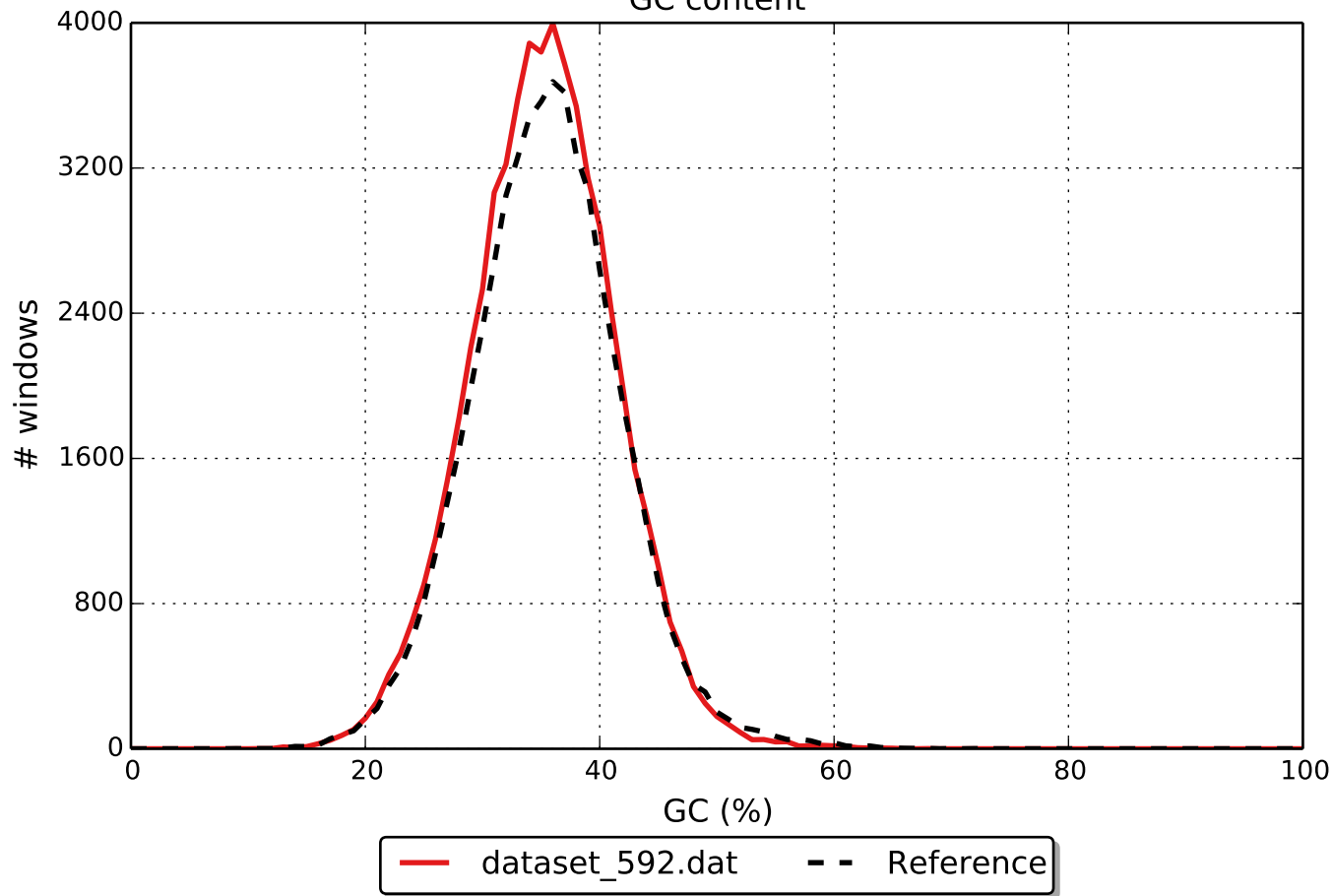


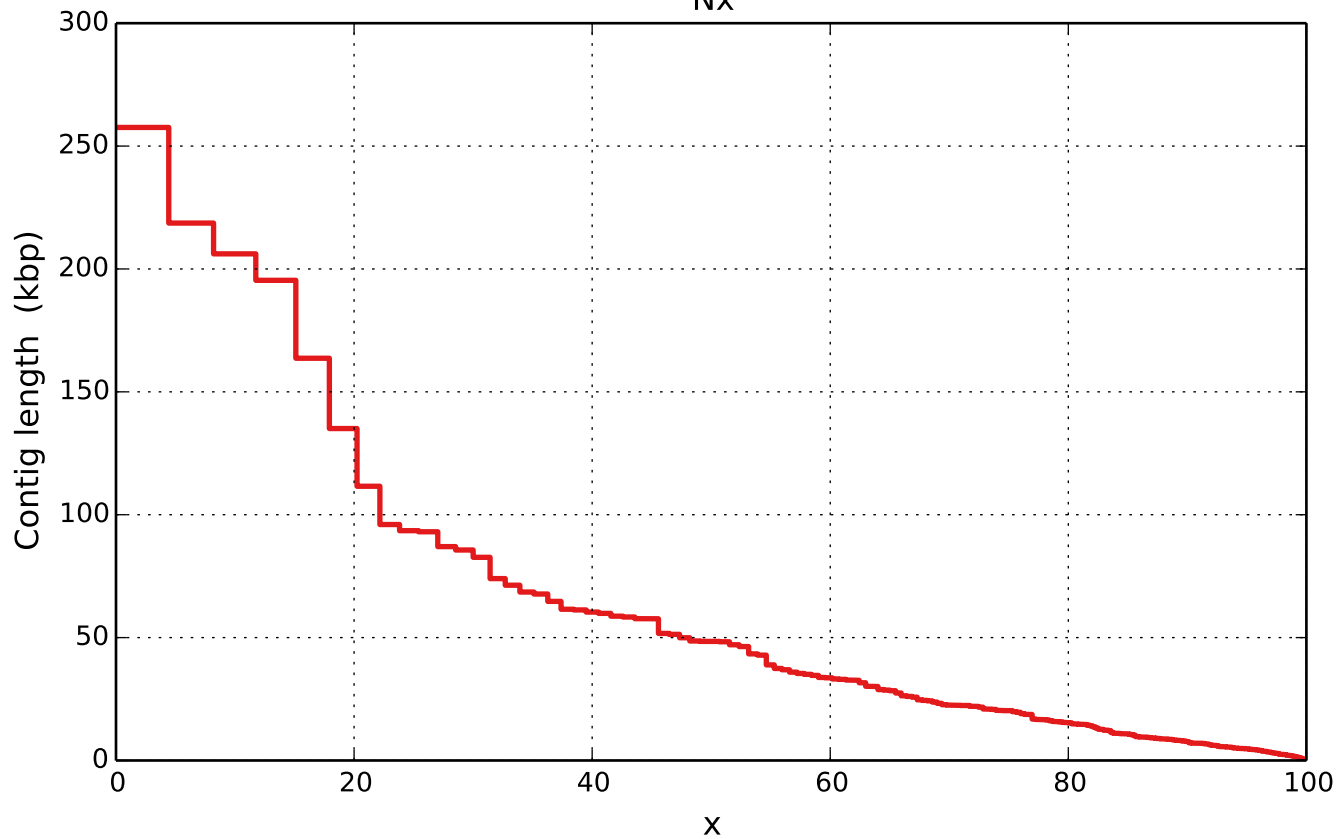
Contig



GC content

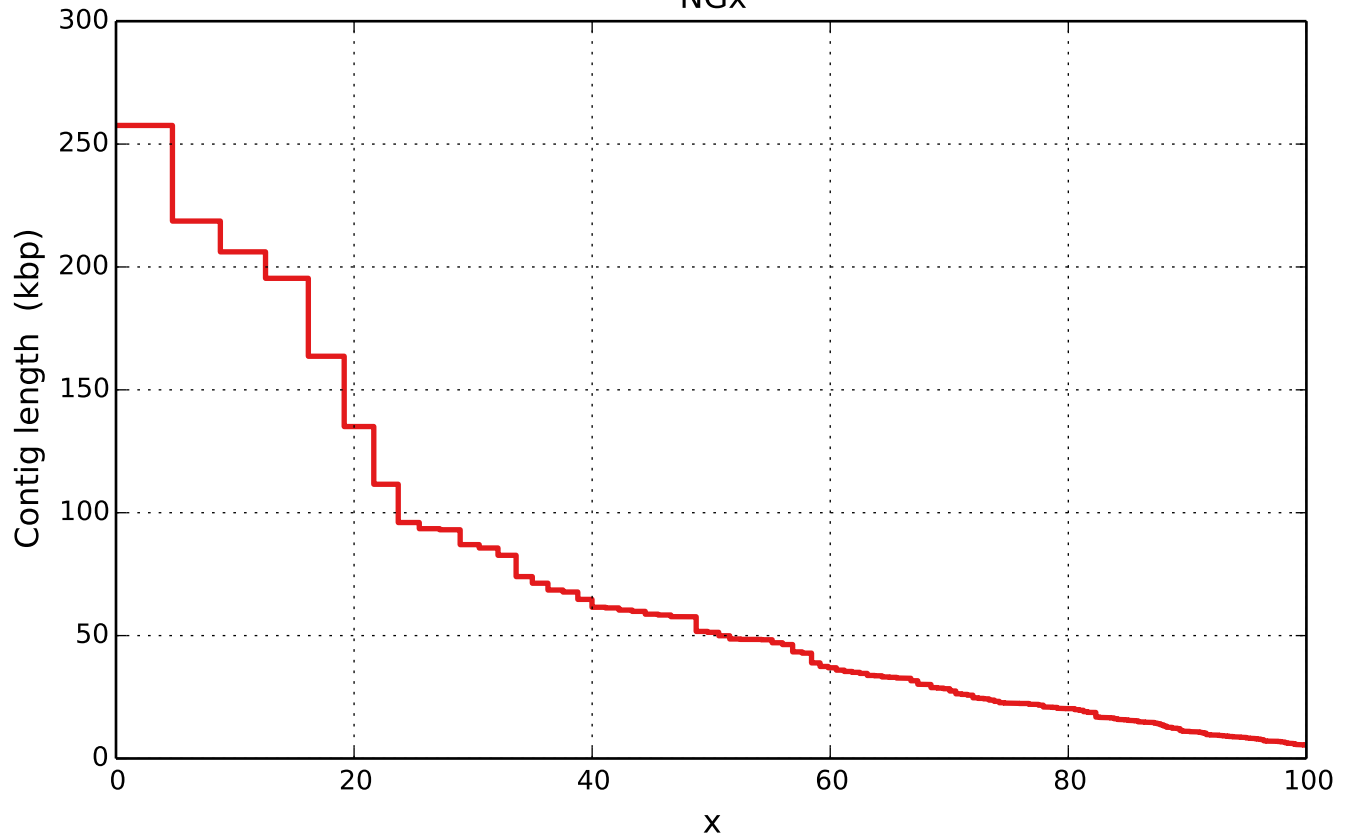


Nx

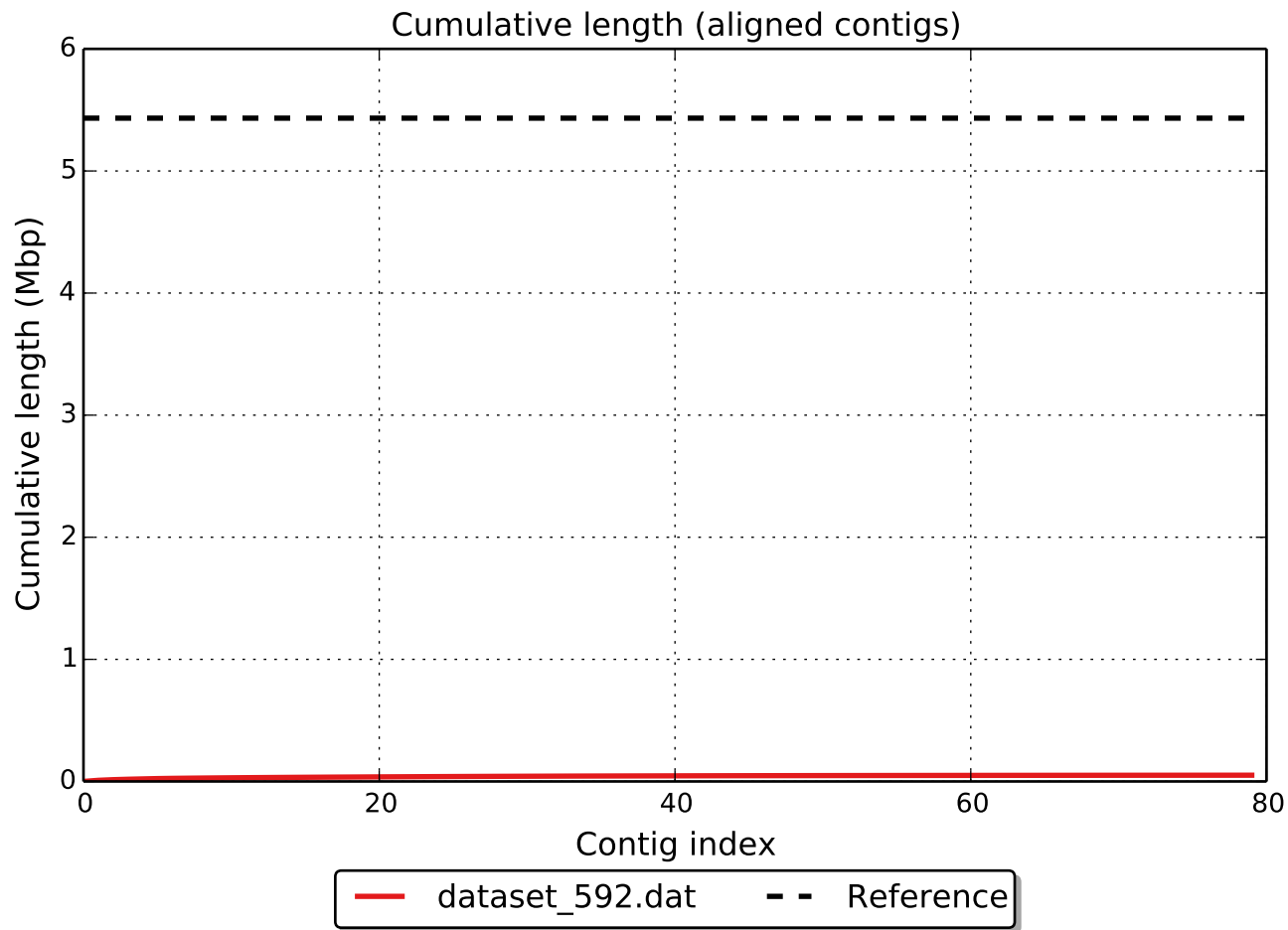


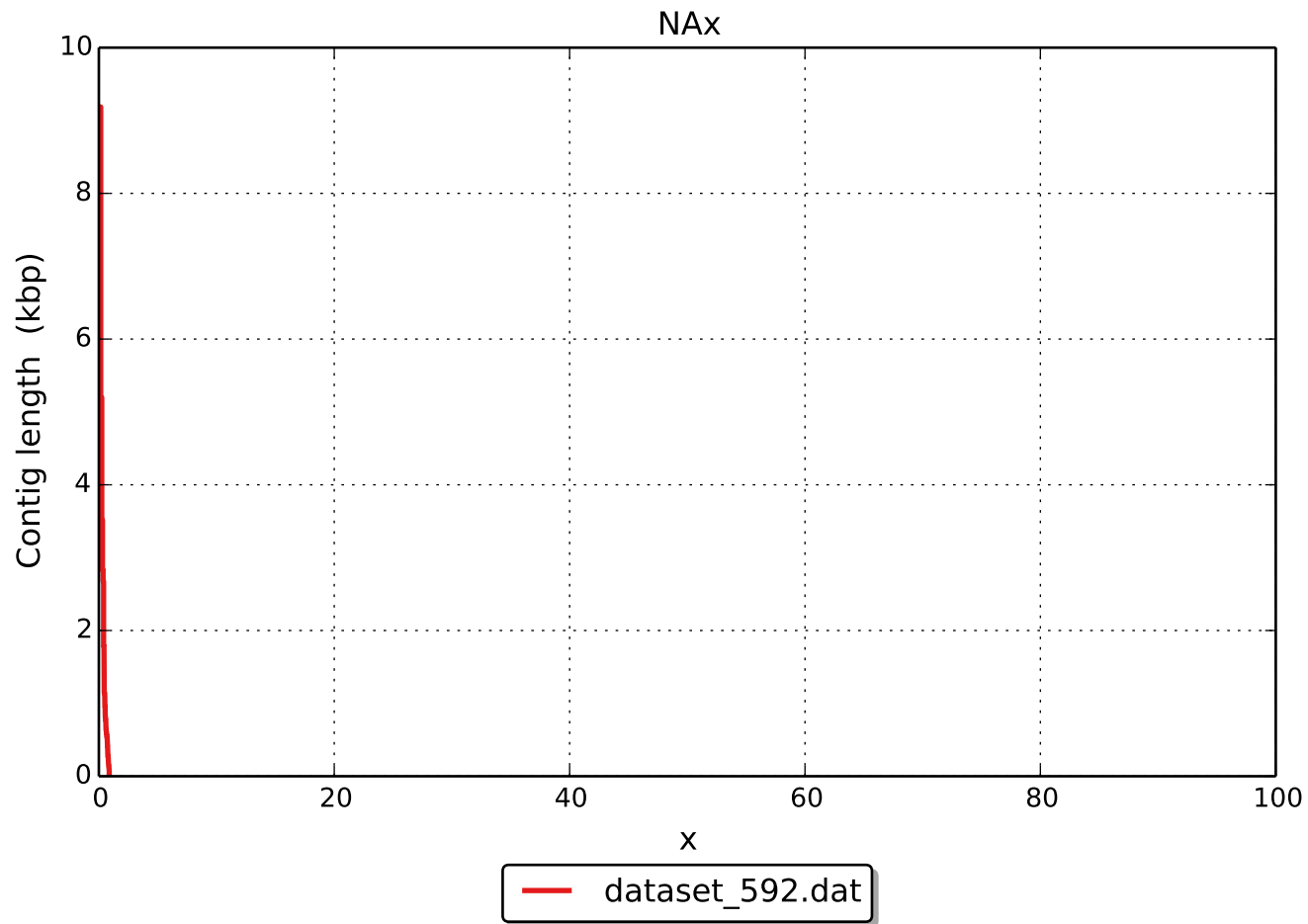
dataset_592.dat

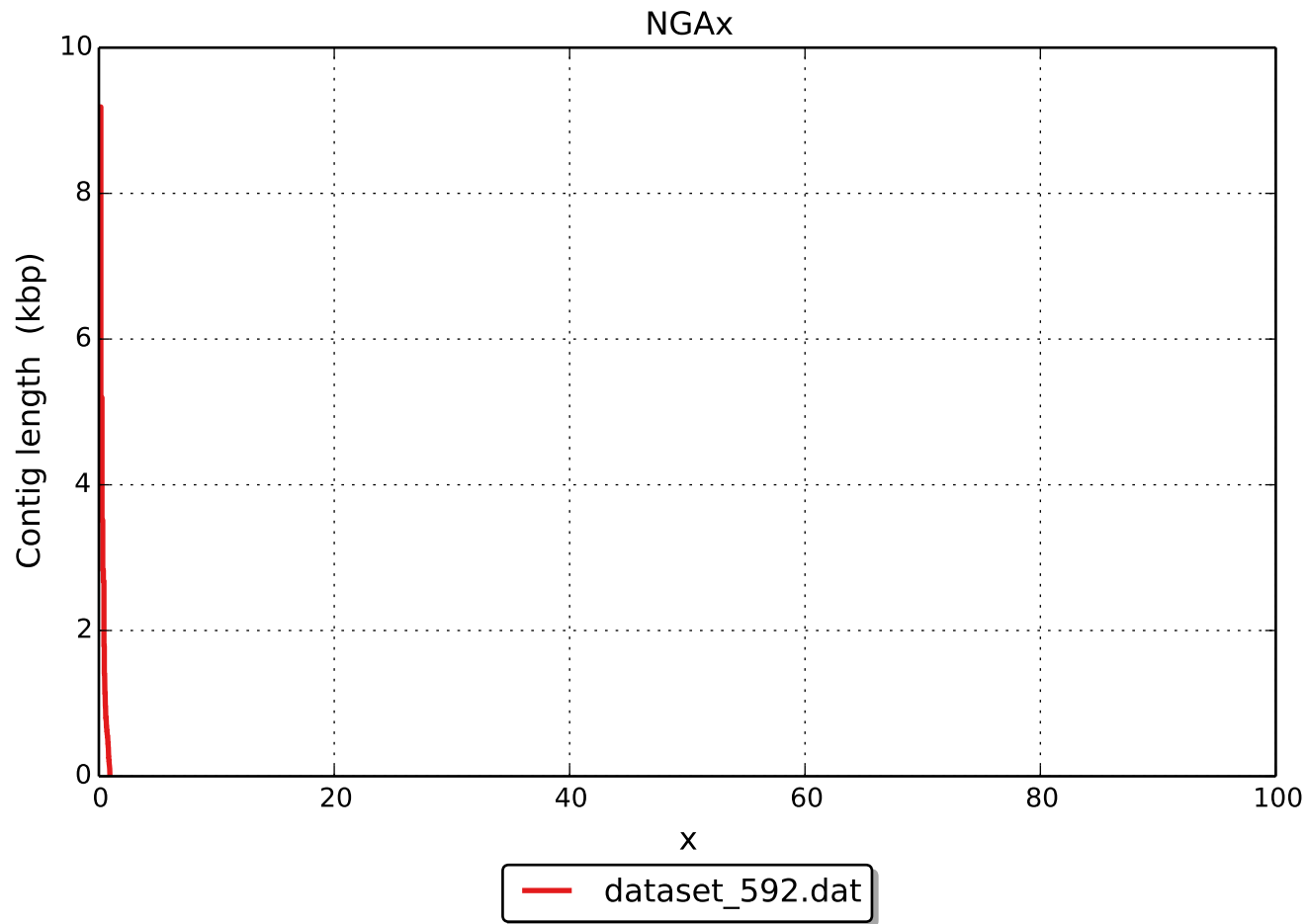
NGx



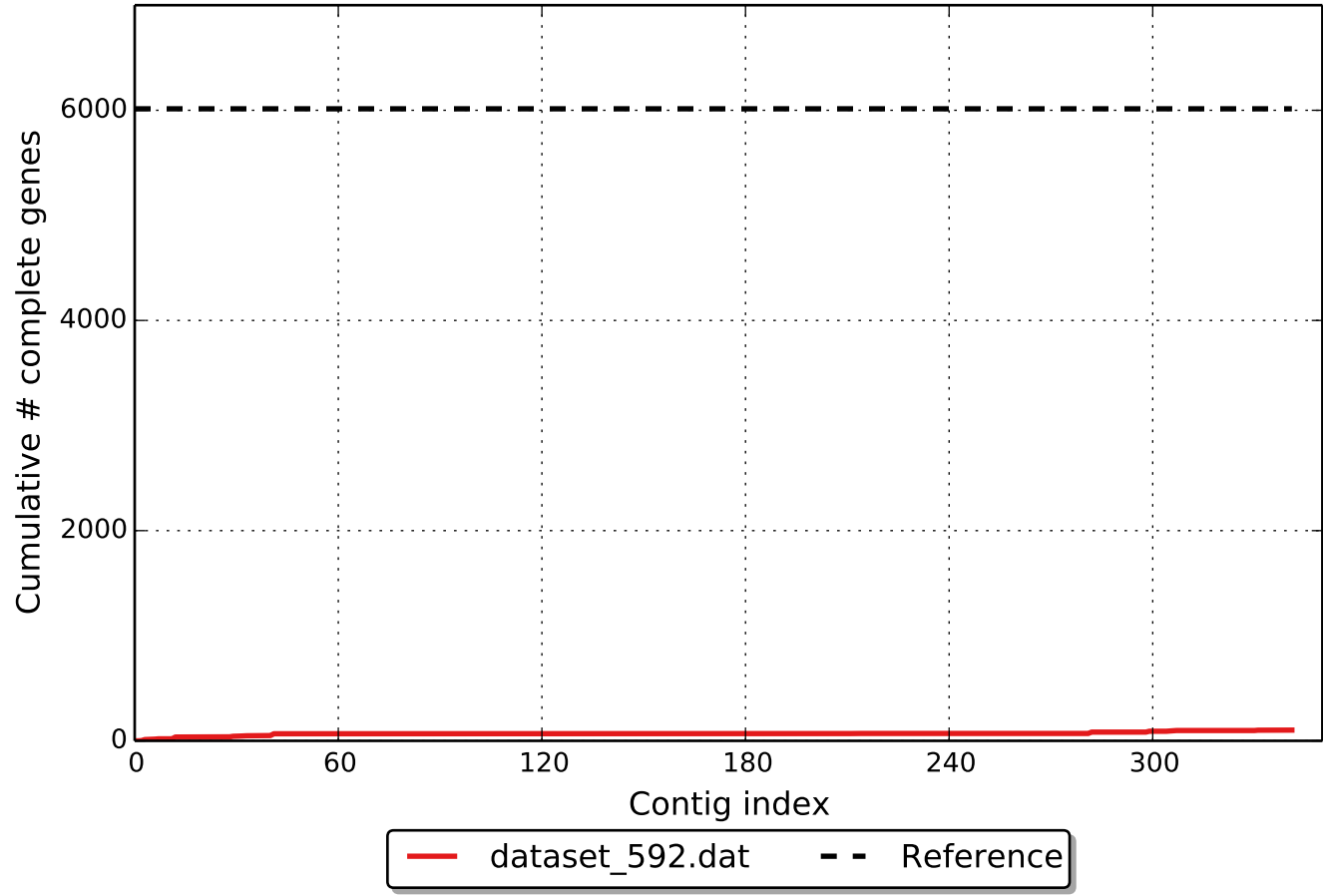
dataset_592.dat

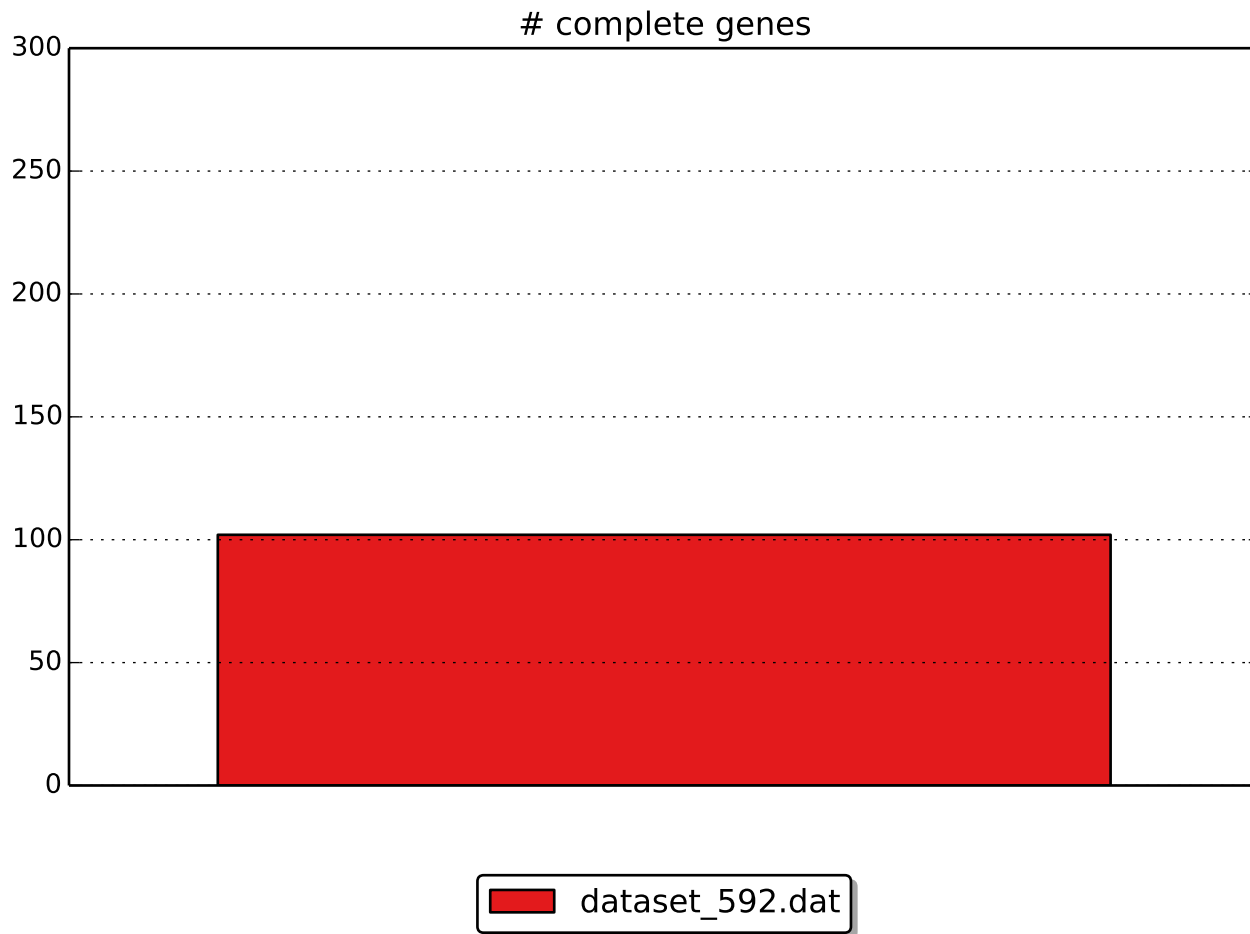


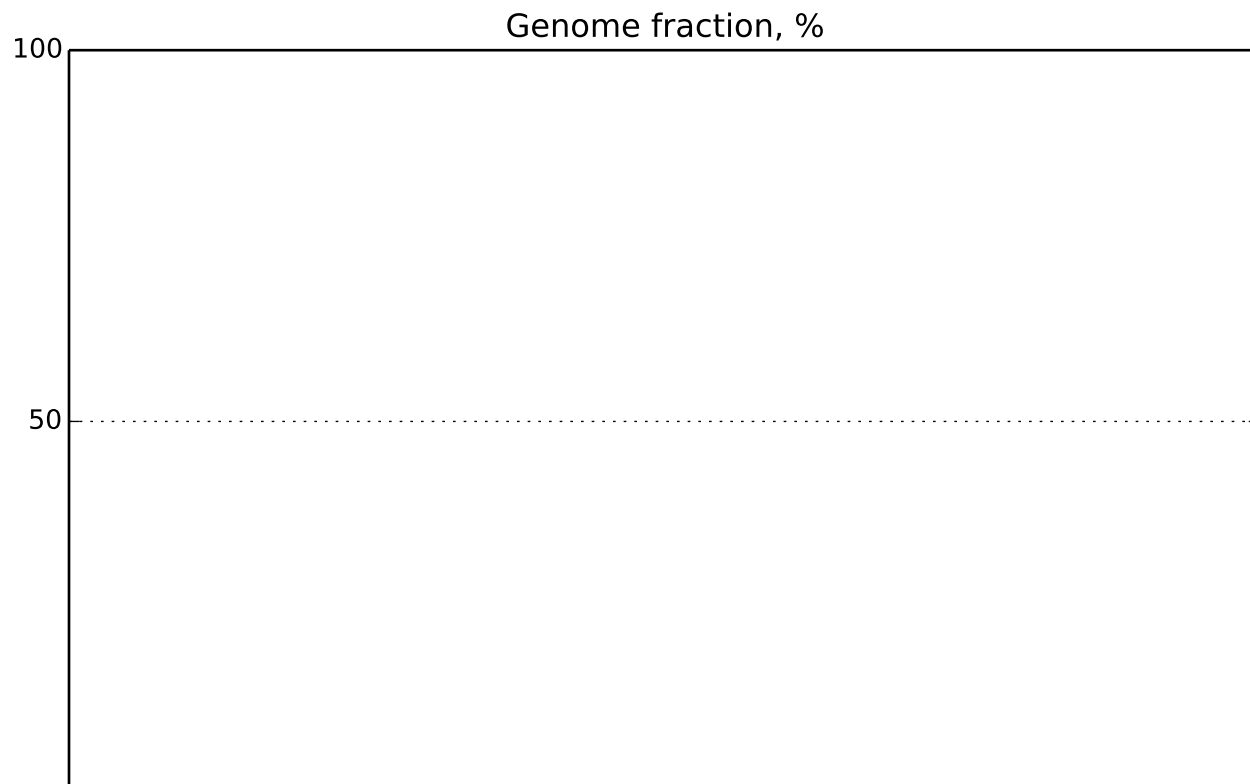





Cumulative # complete genes

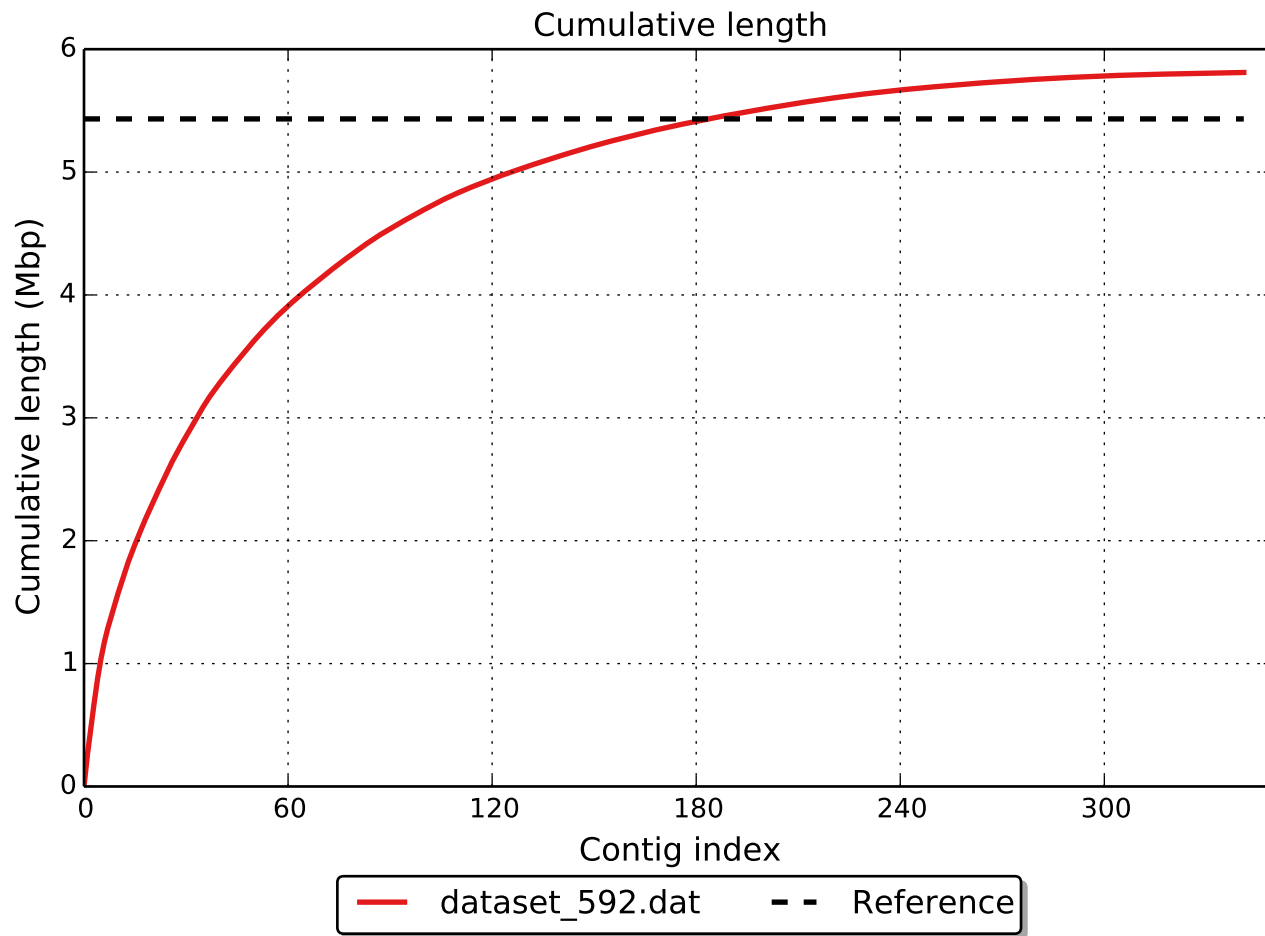




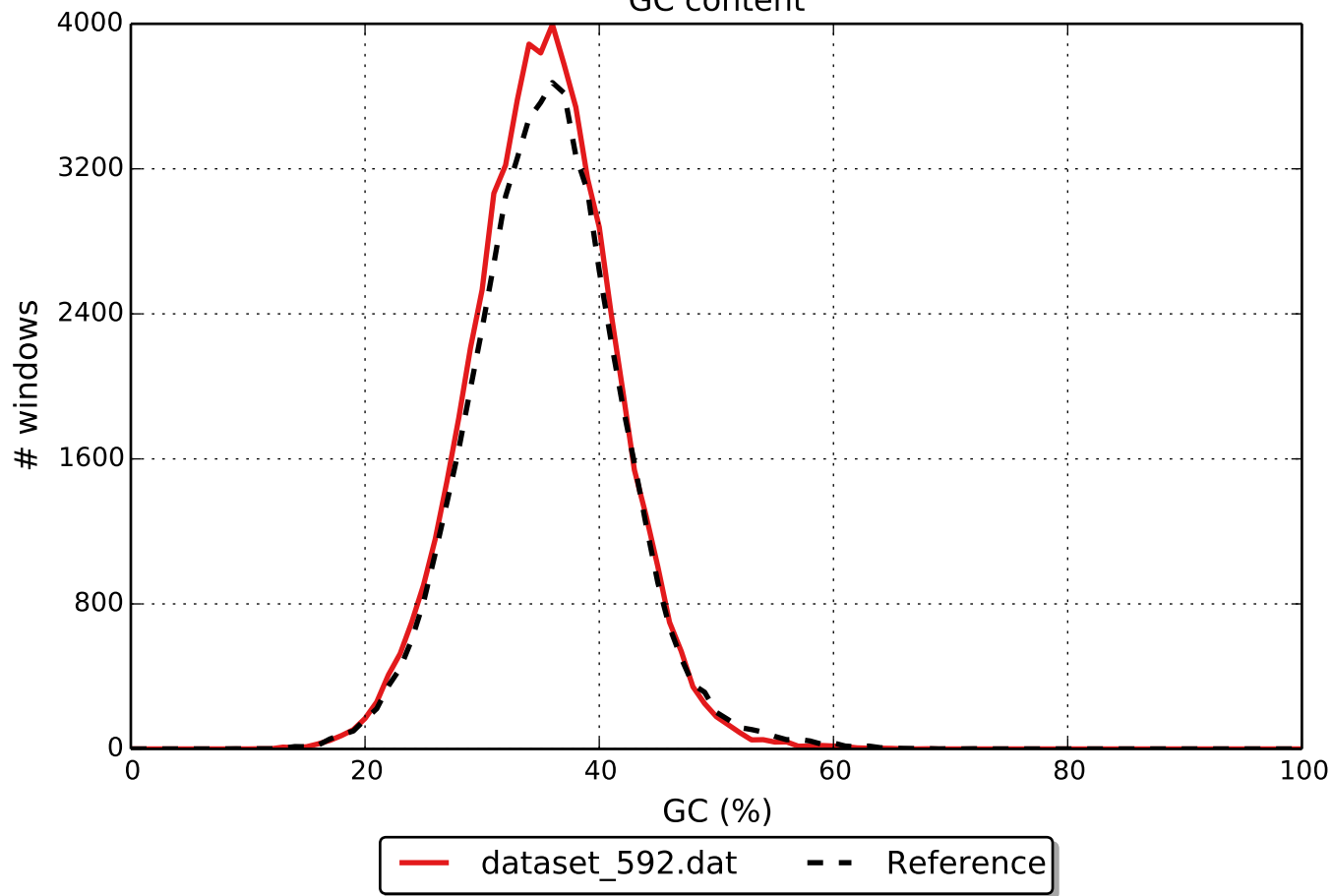


 dataset_592.dat

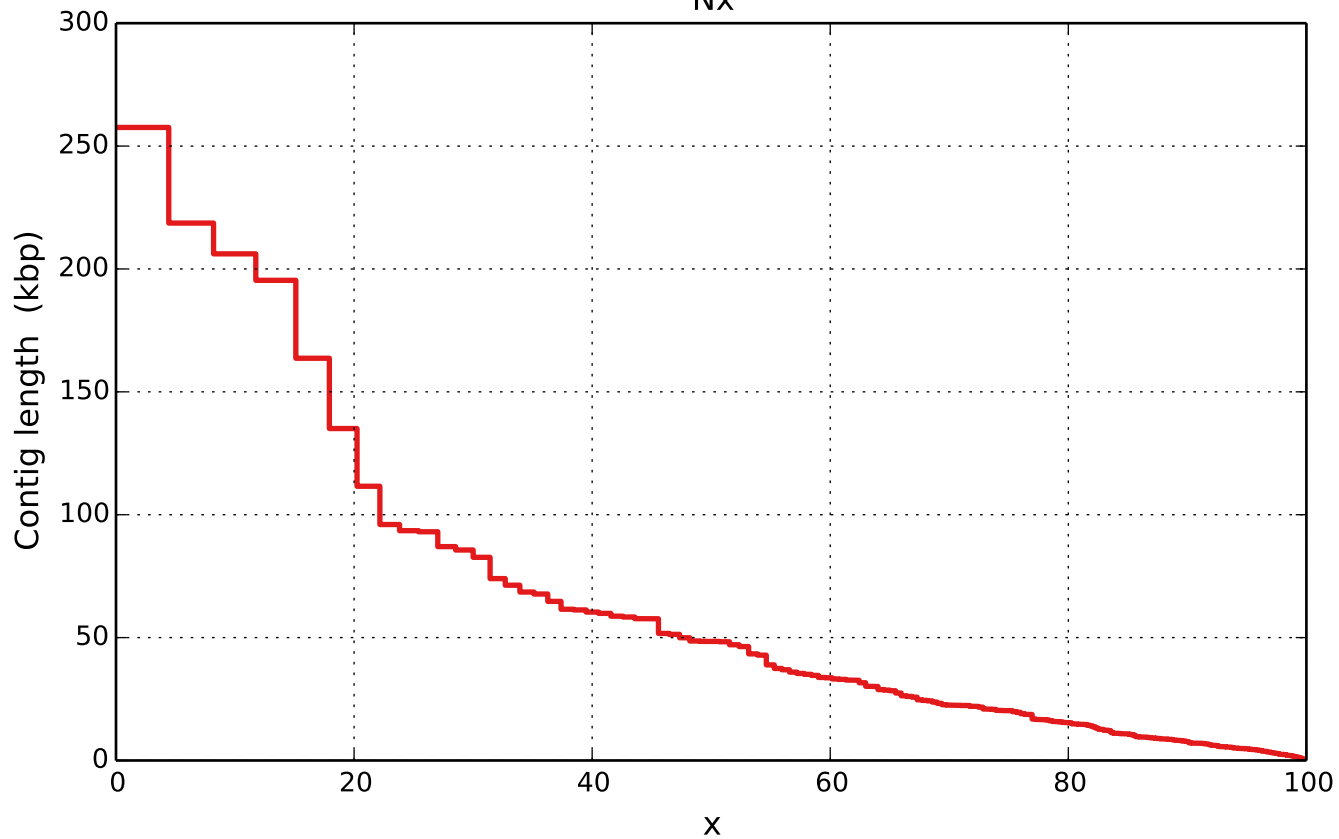
Contig



GC content

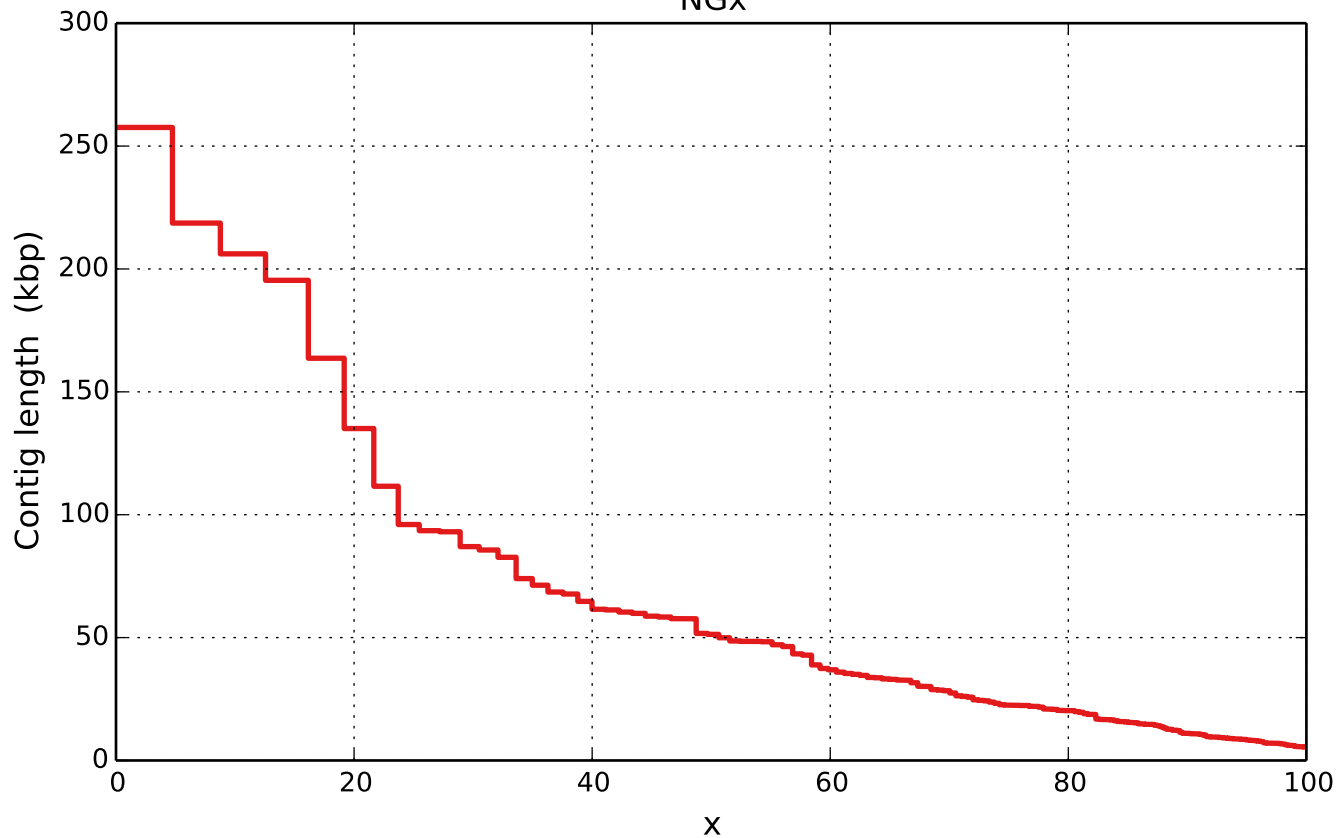


Nx



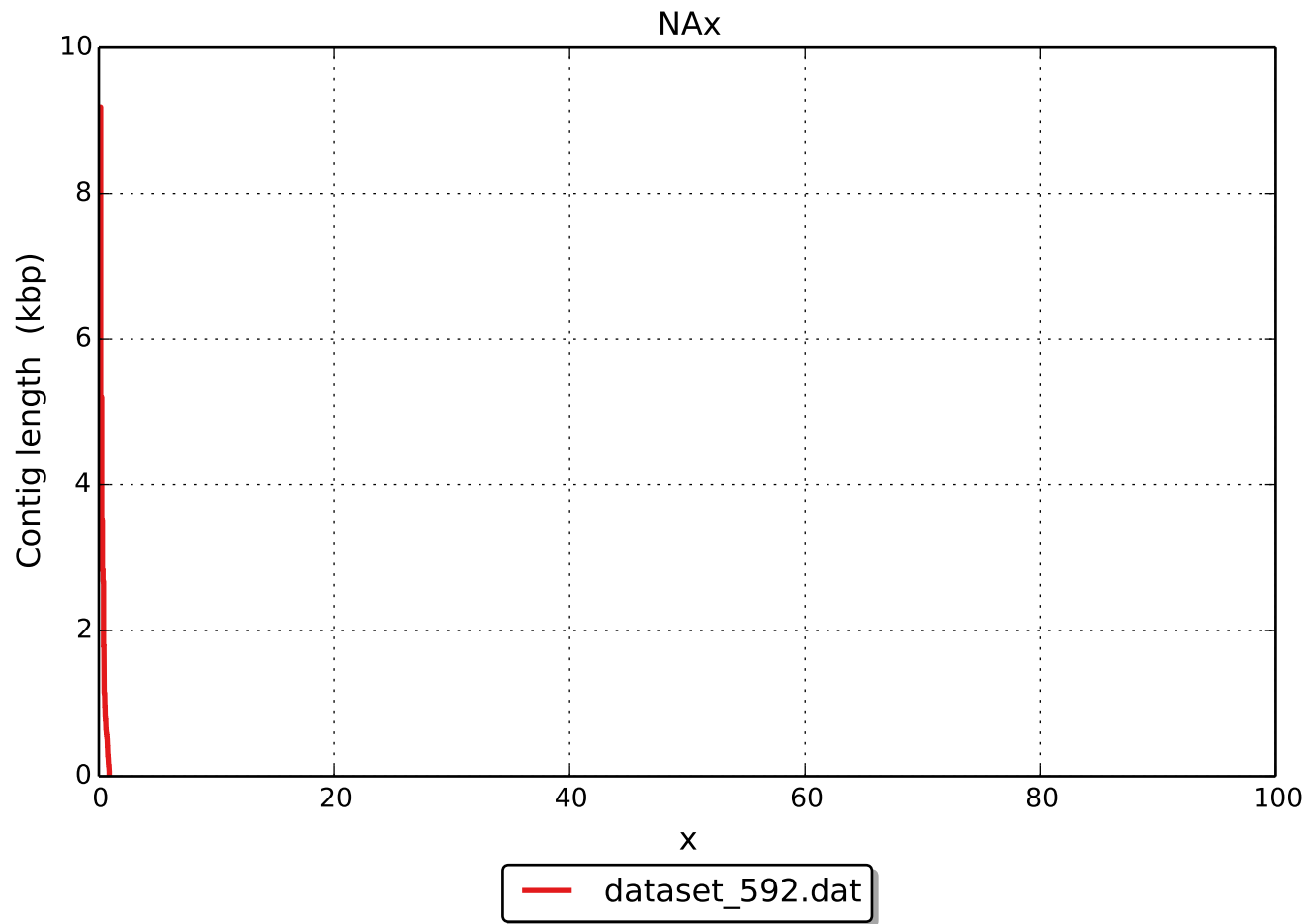
— dataset_592.dat

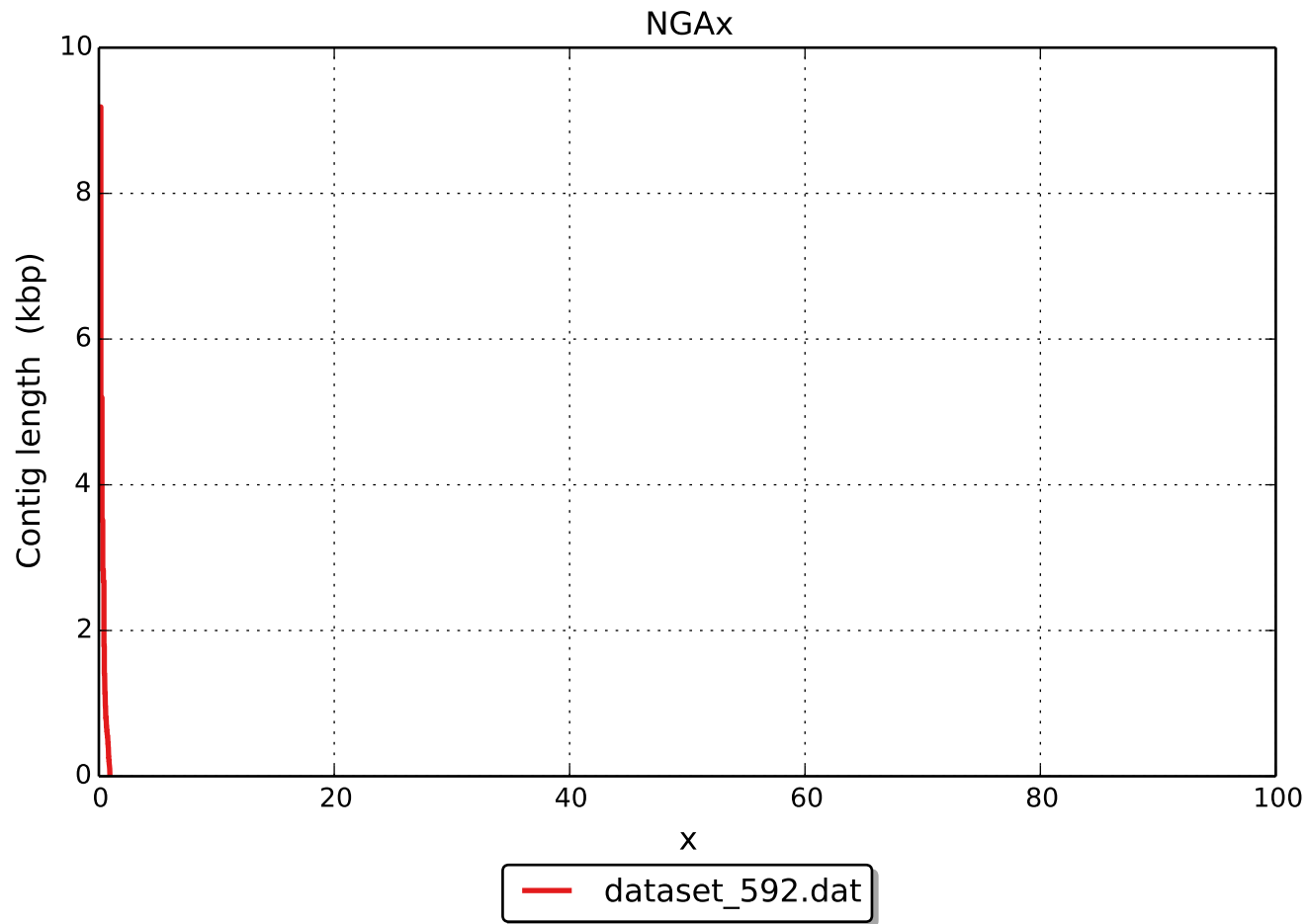
NGx

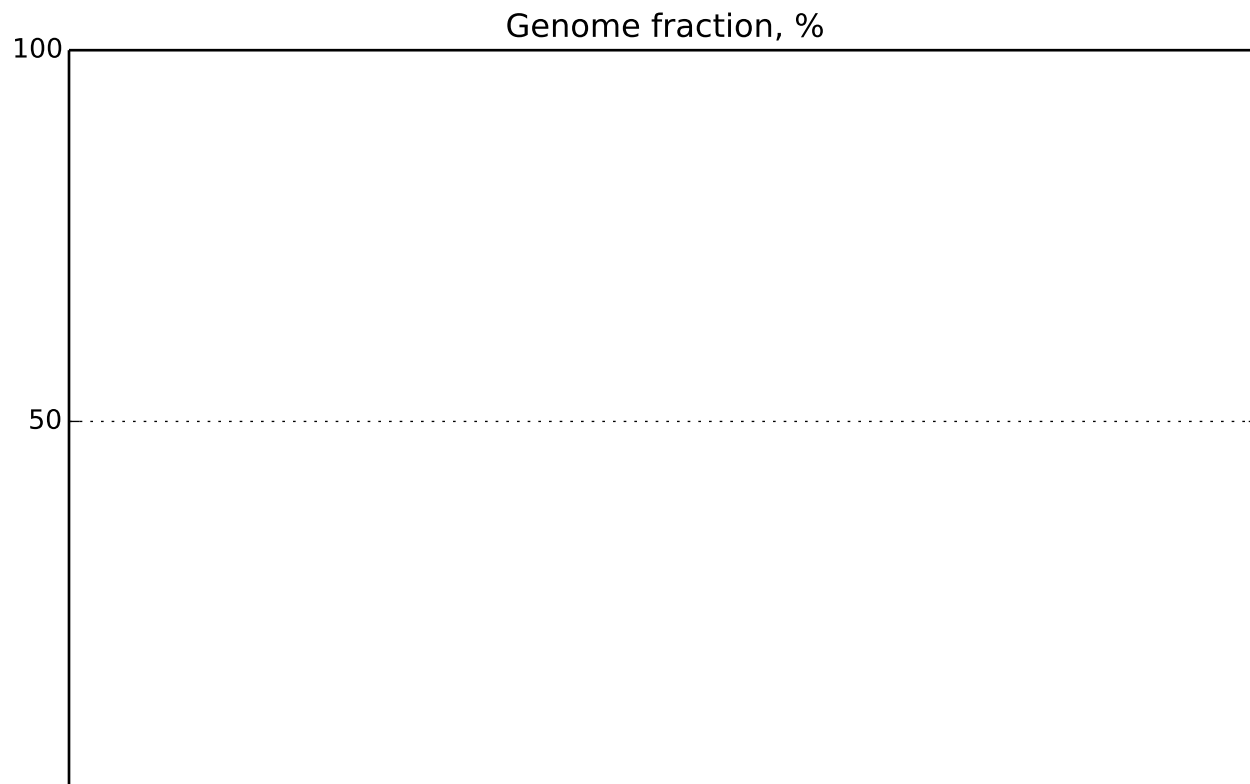


— dataset_592.dat









 dataset_592.dat