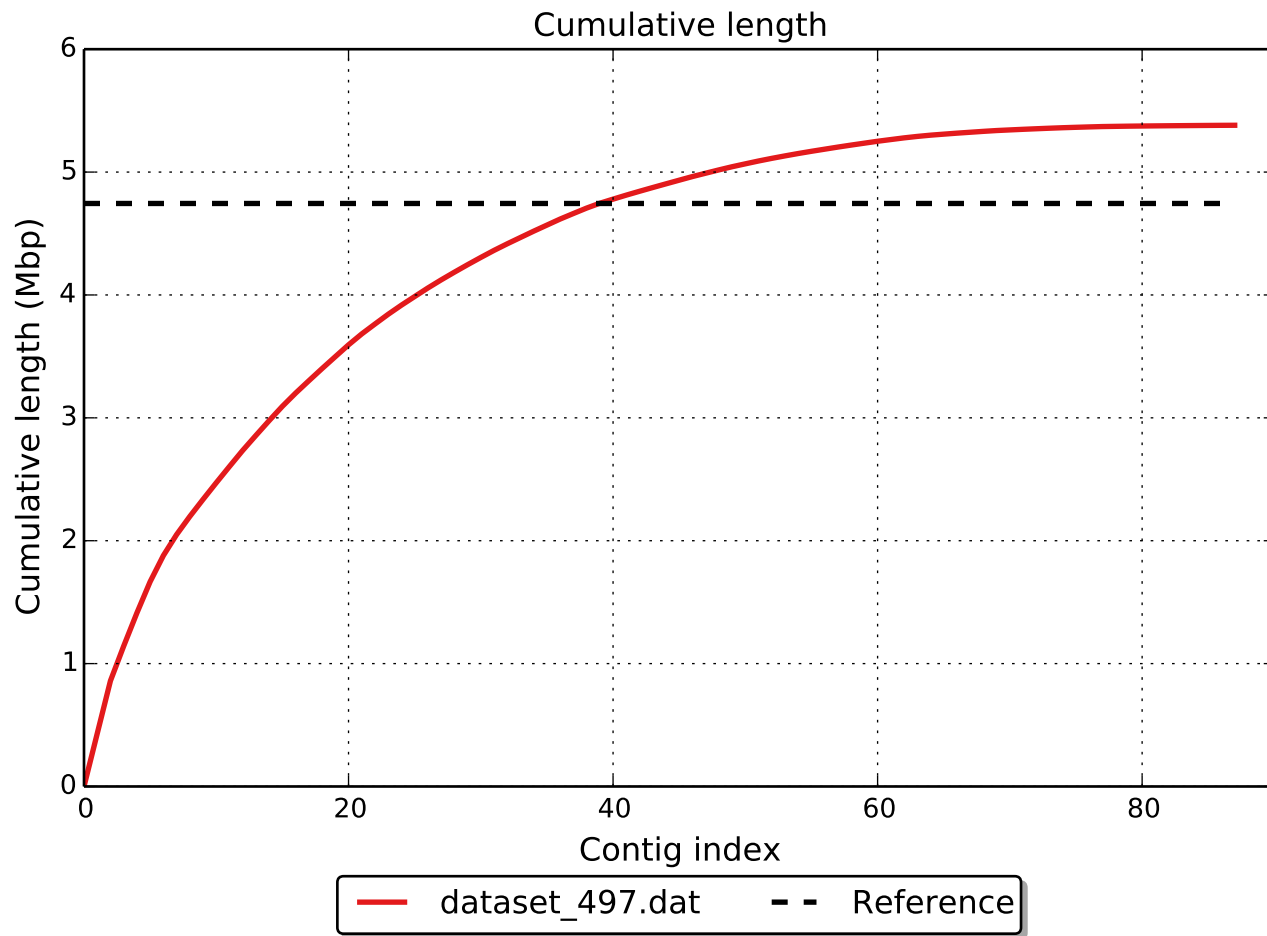
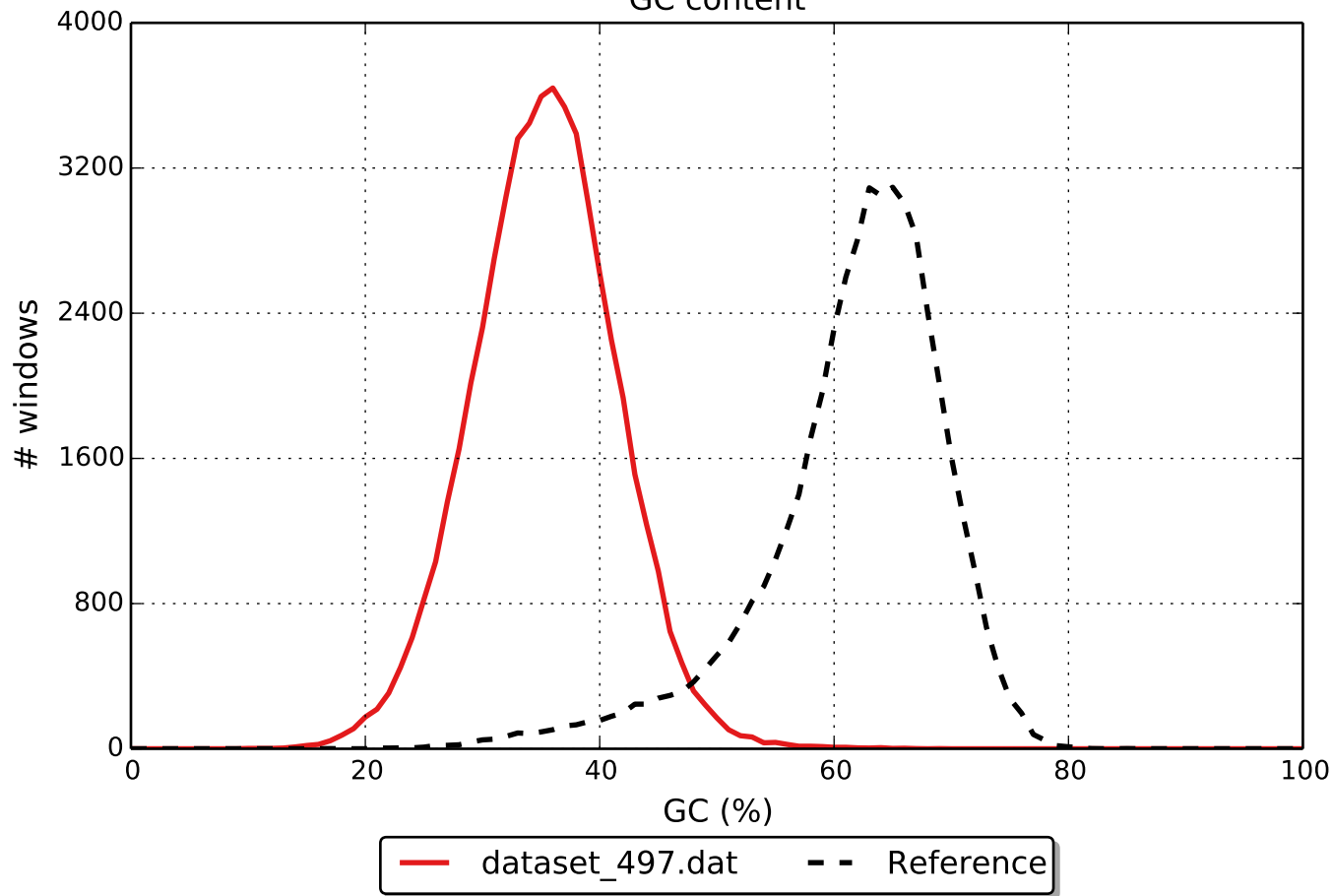


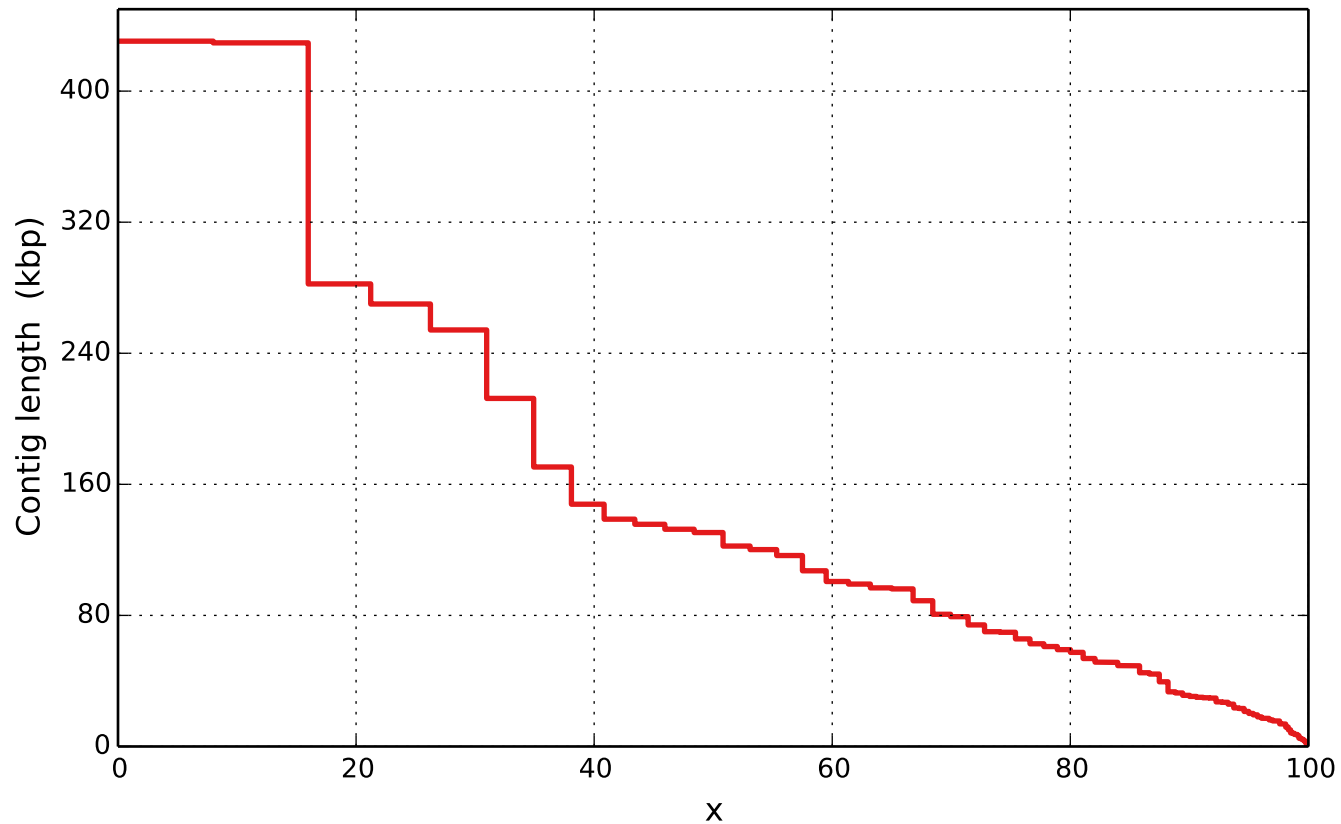
A_hydrophila_ATCC_7966



GC content

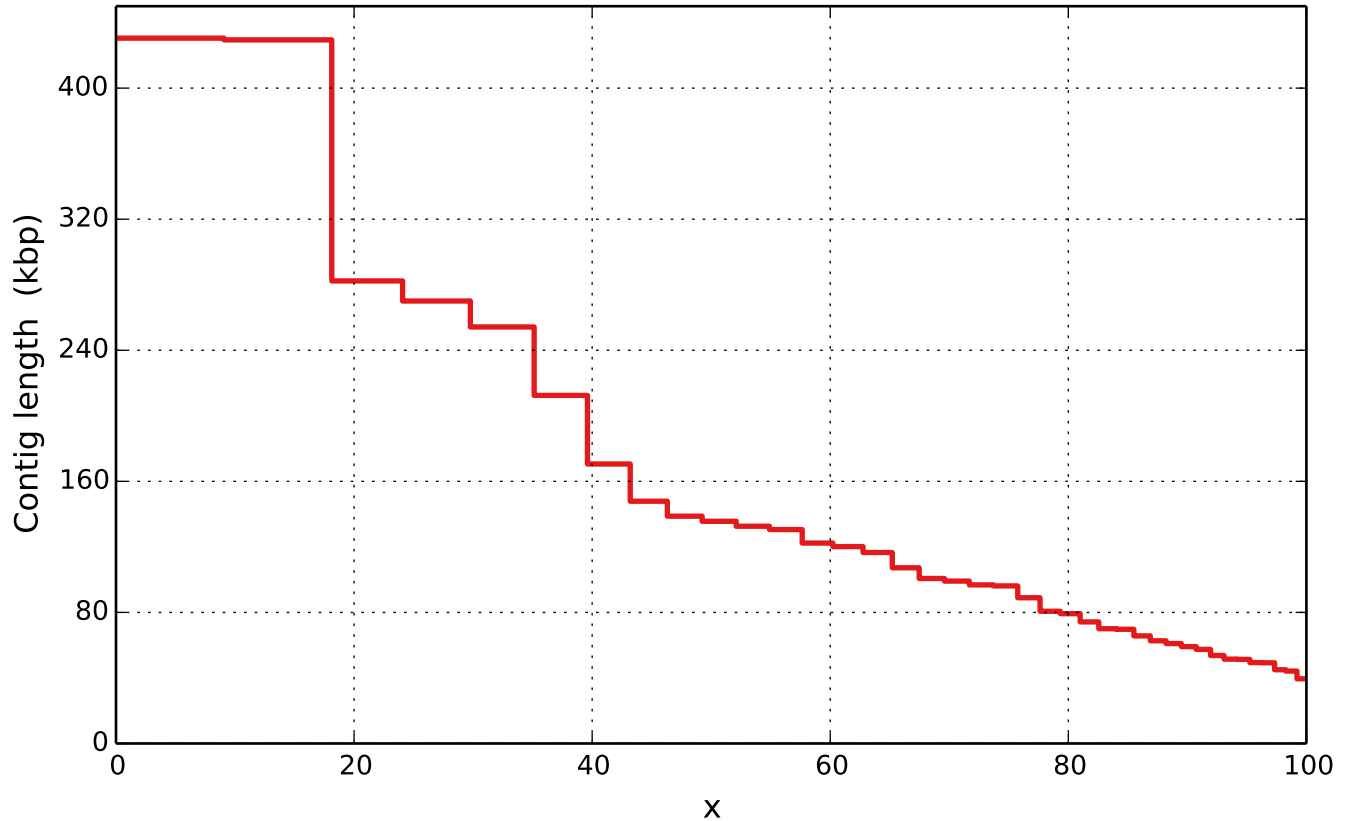


Nx



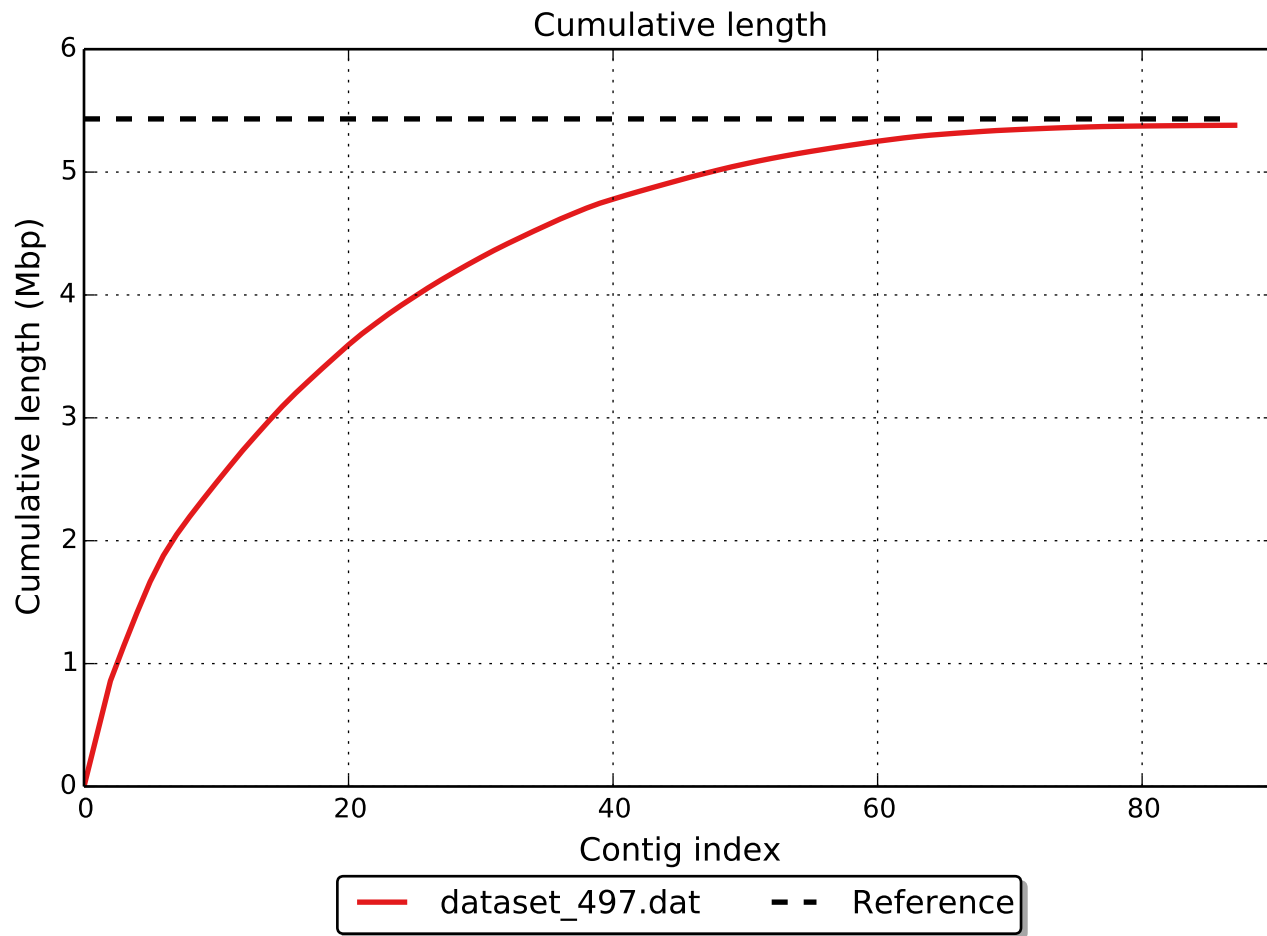
dataset_497.dat

NGx

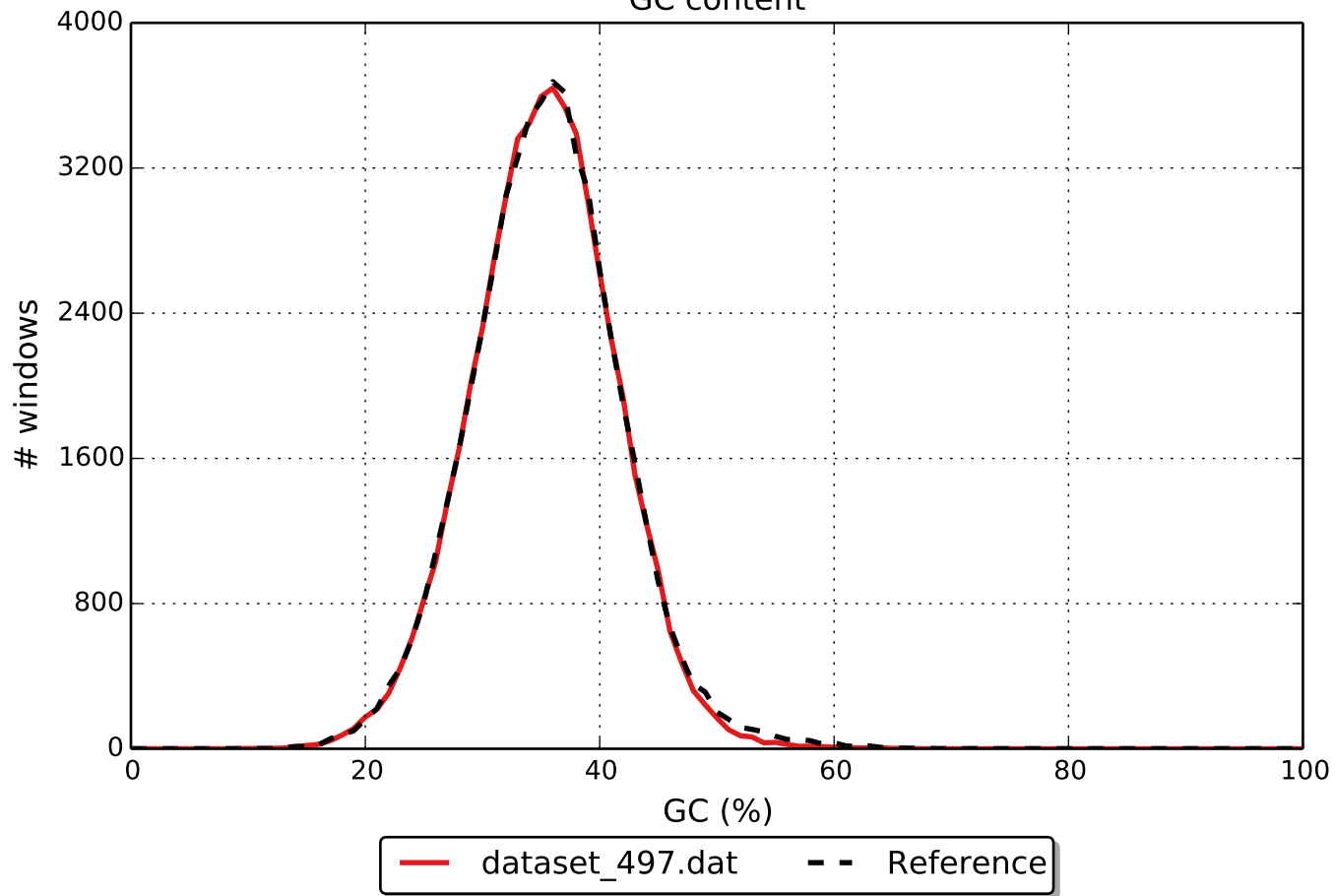


dataset_497.dat

B_cereus_ATCC_1098



GC content

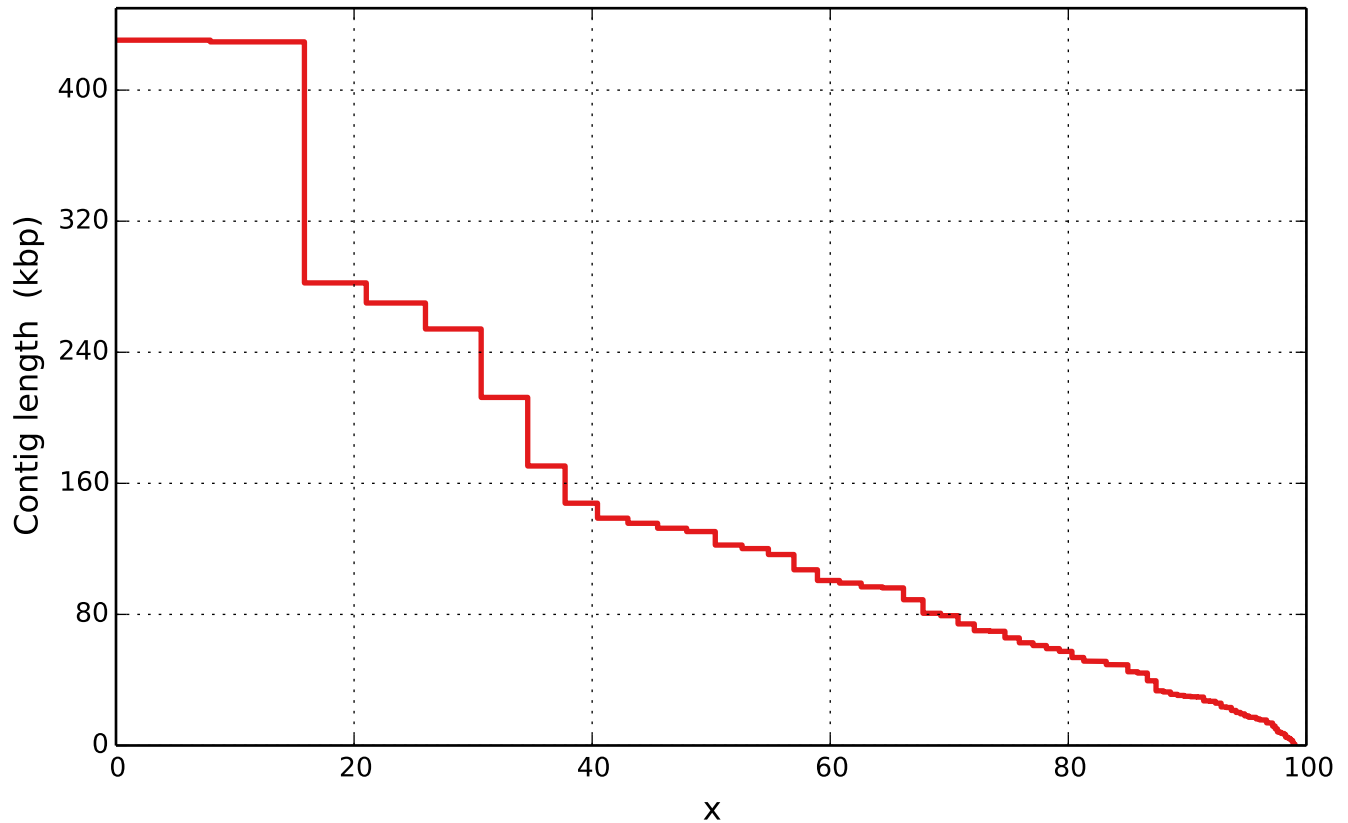


Nx

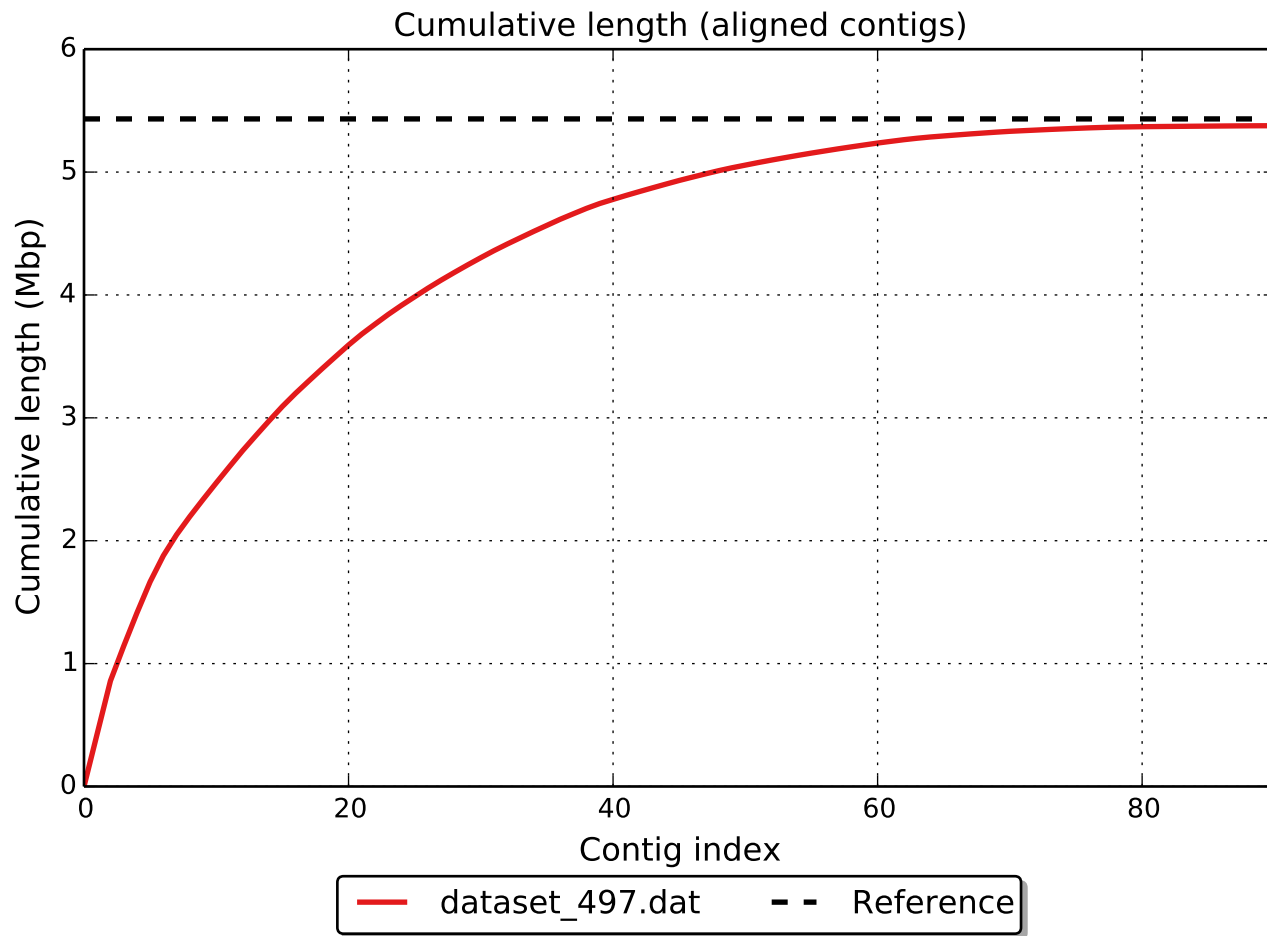


dataset_497.dat

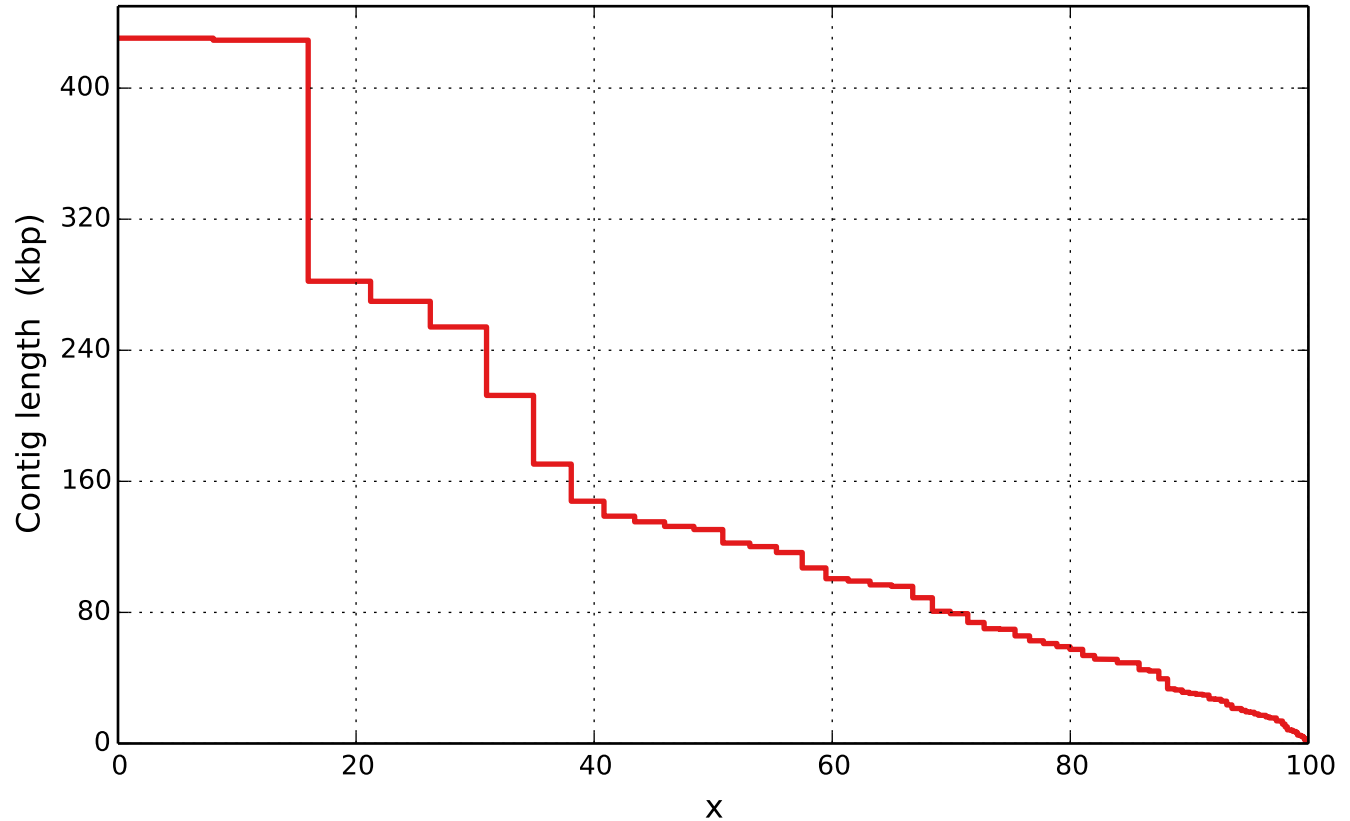
NGx



— dataset_497.dat

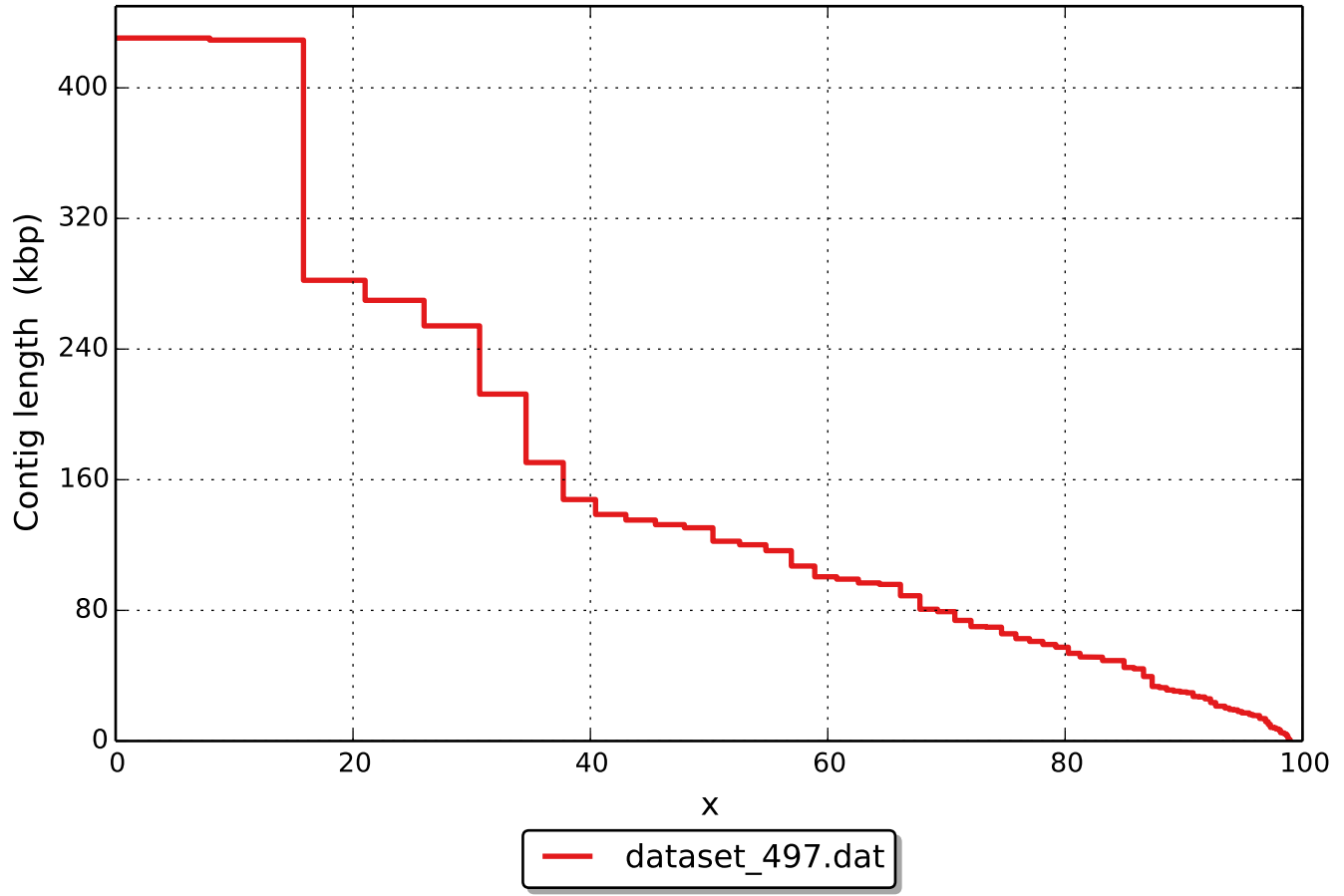


NAx

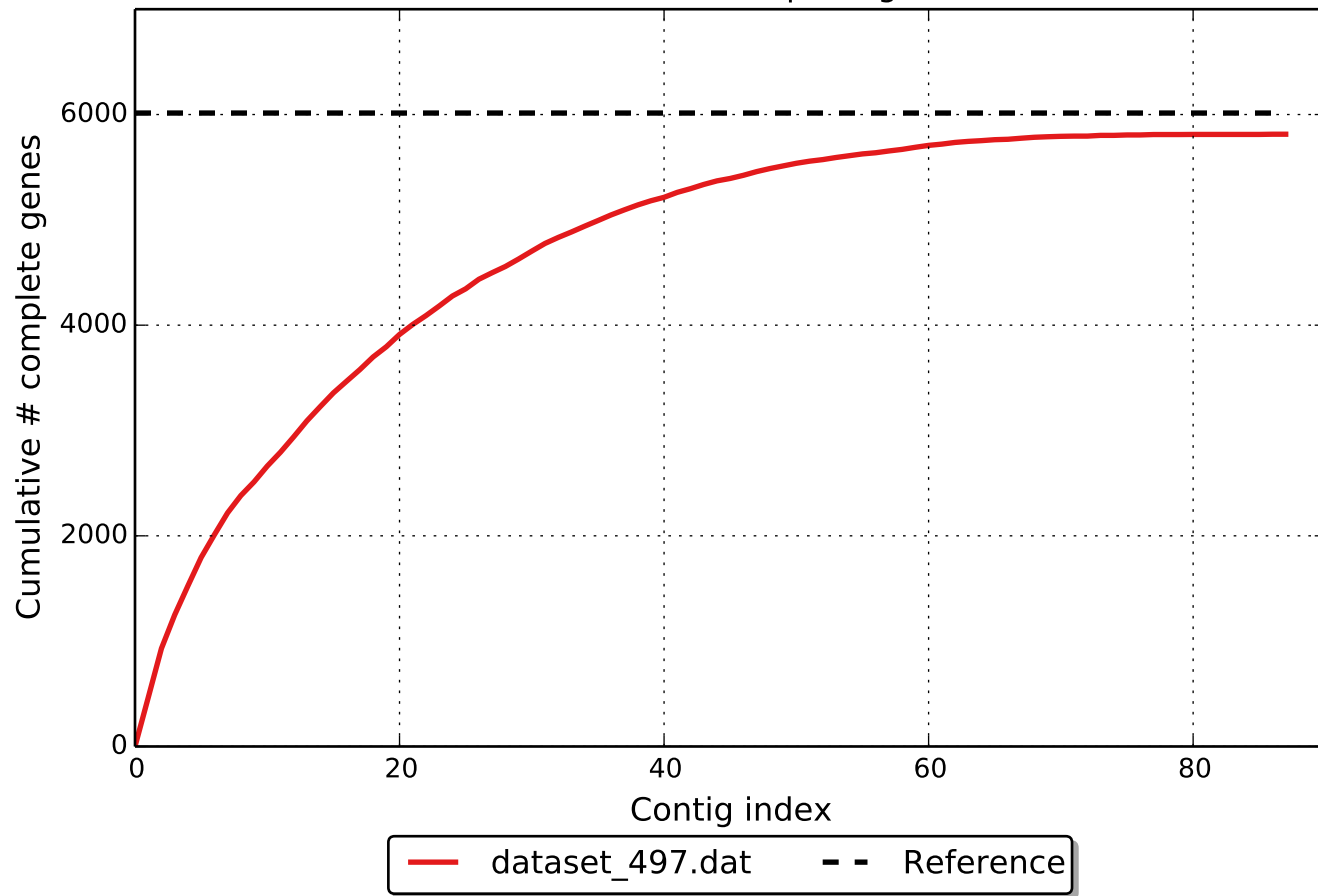


dataset_497.dat

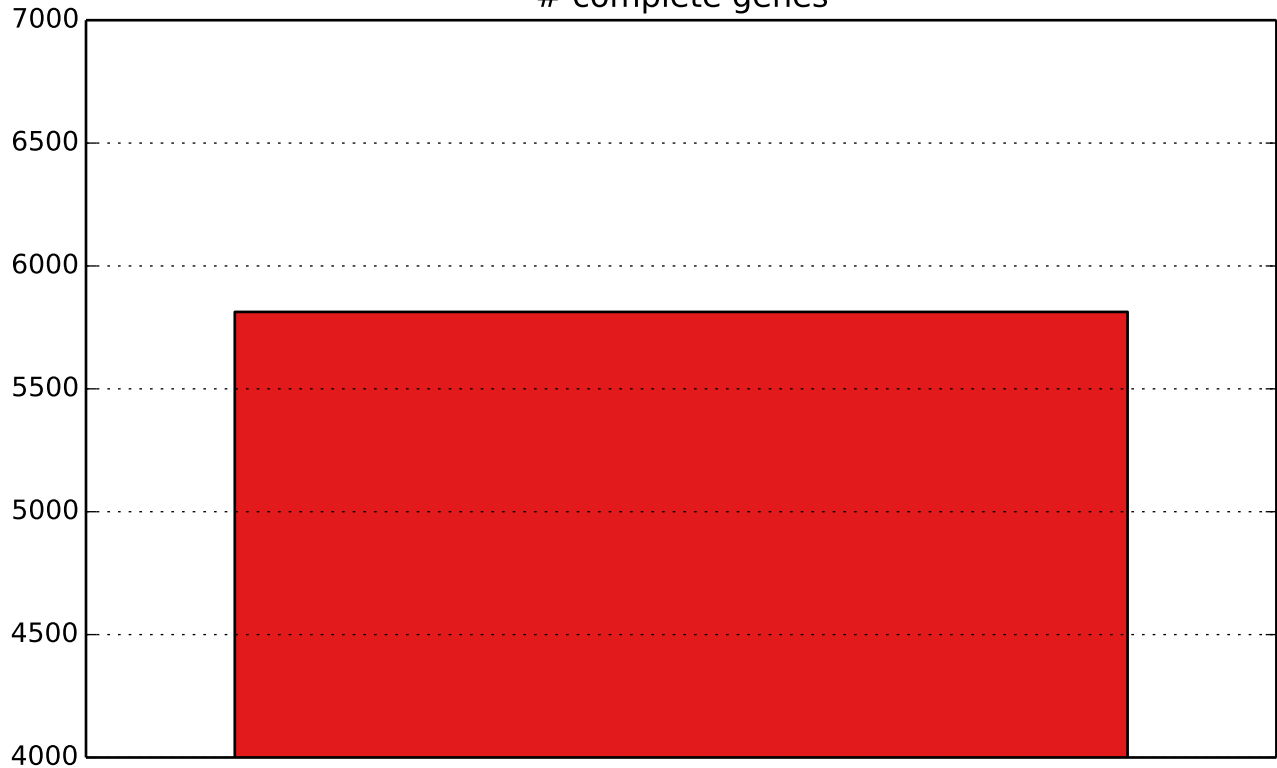
NGAx



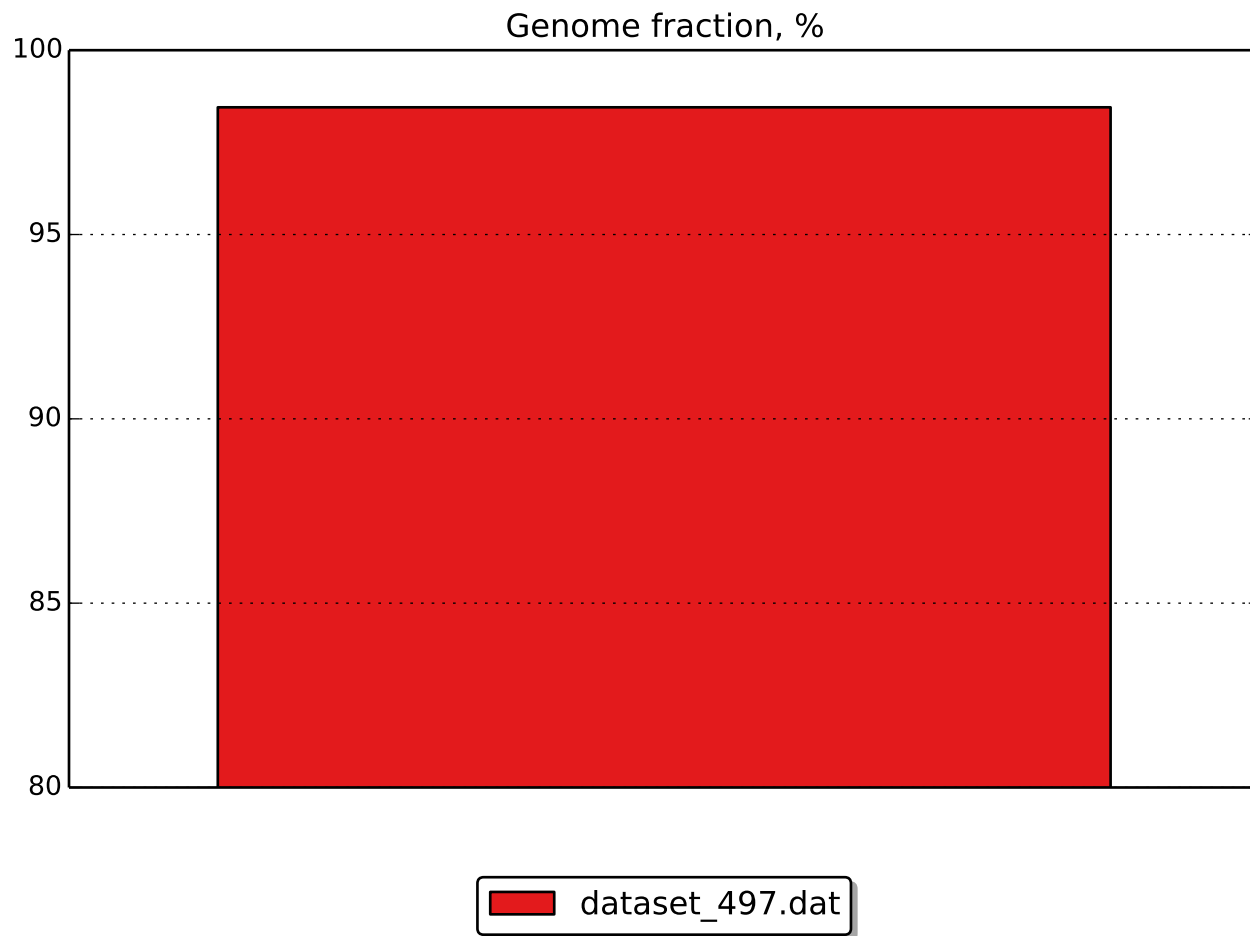
Cumulative # complete genes



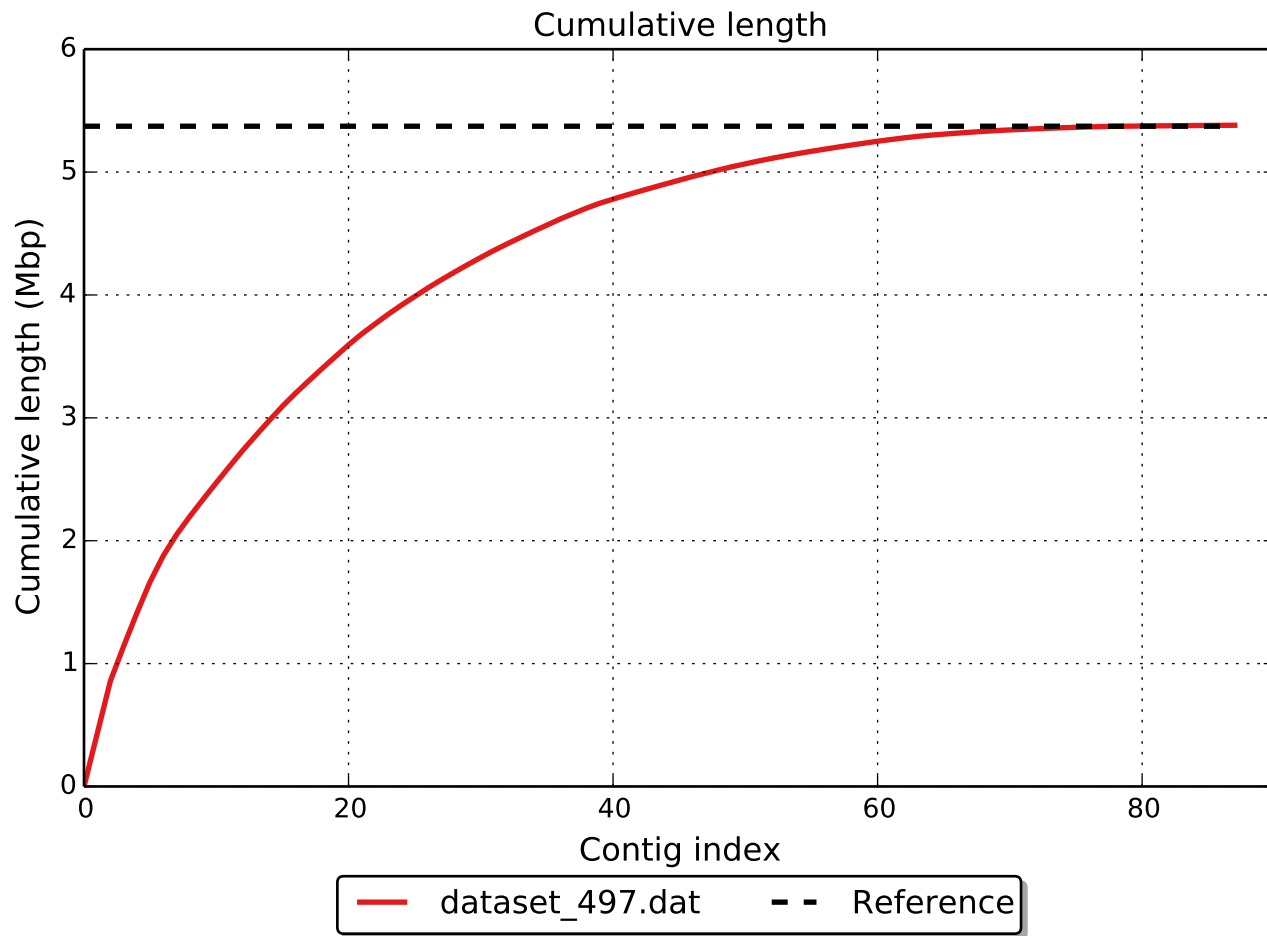
complete genes



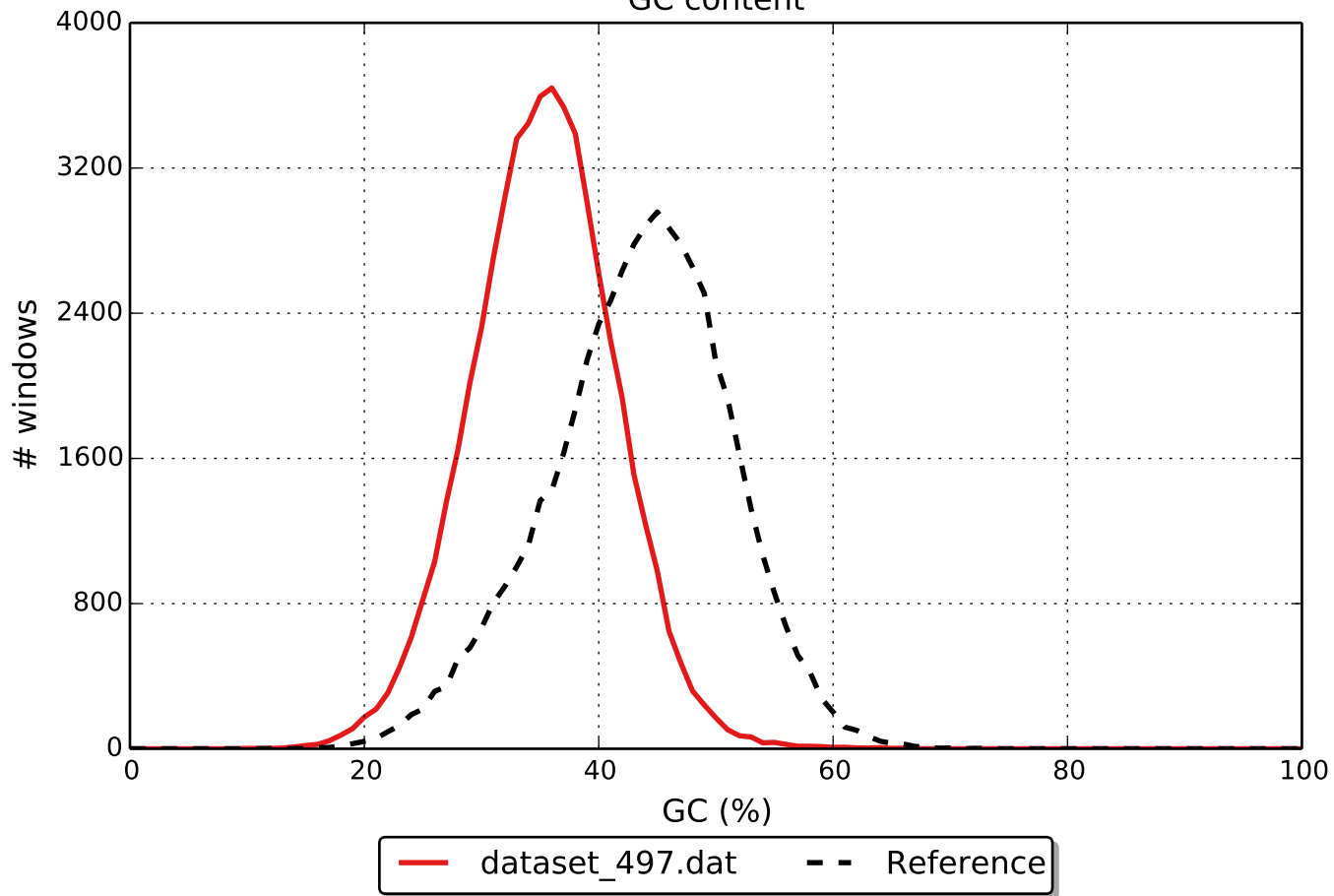
dataset_497.dat



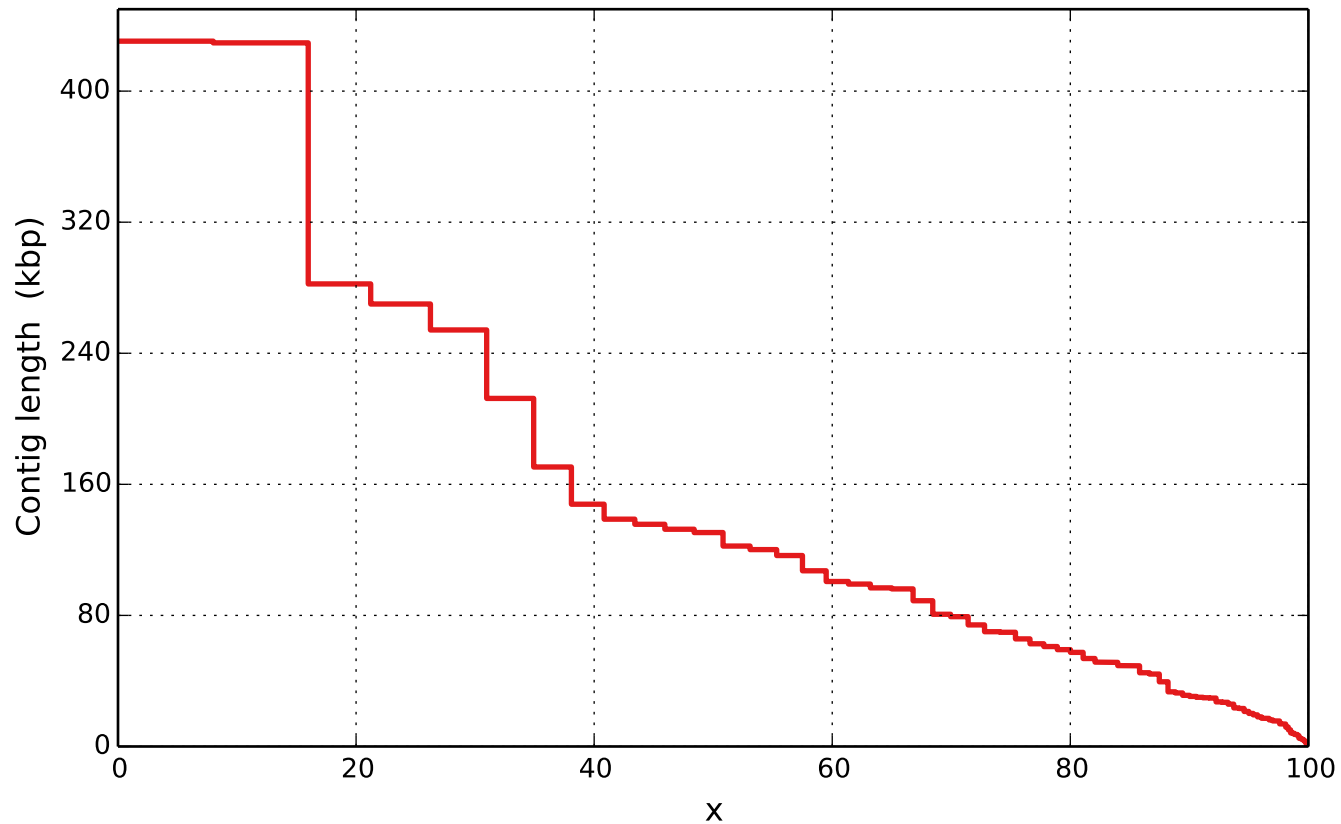
B_fragilis_638R



GC content

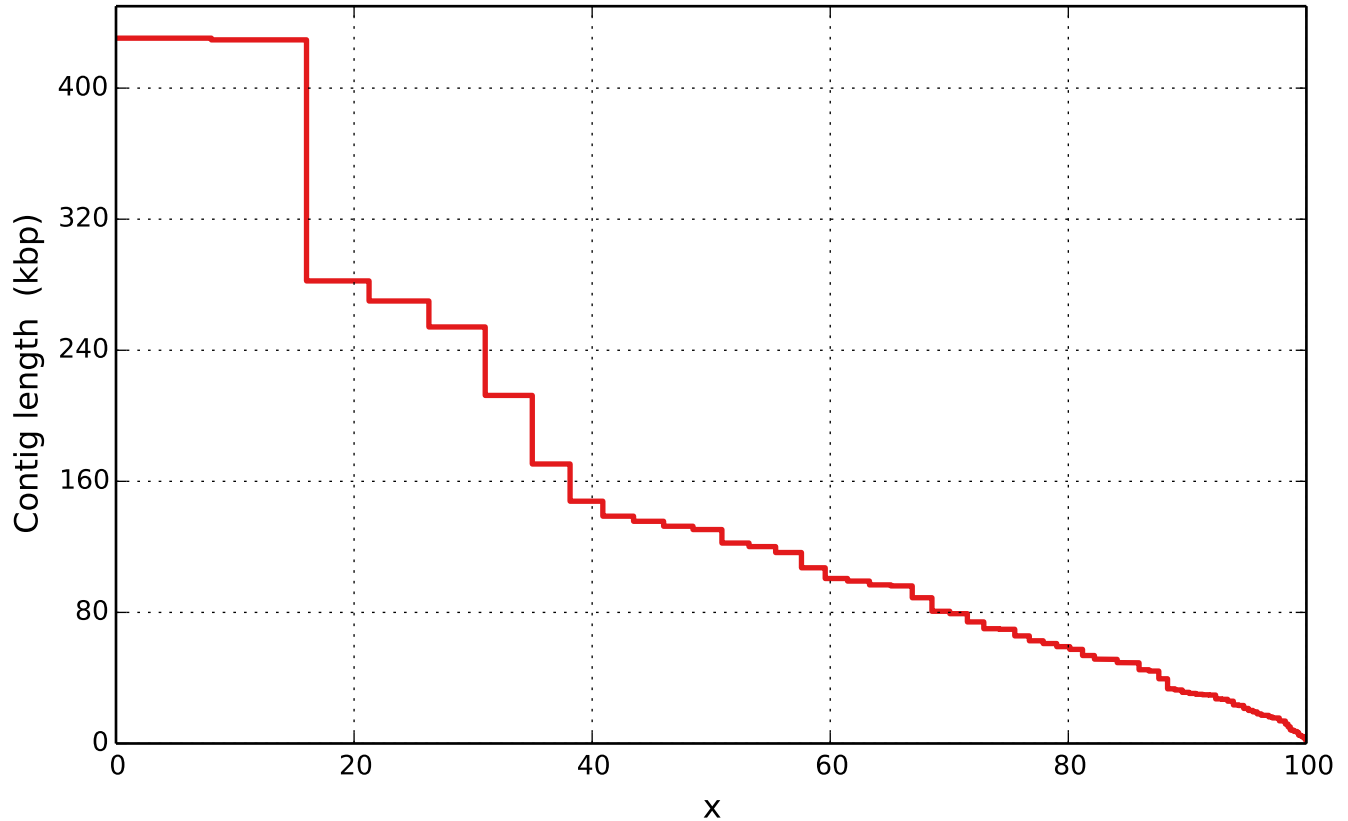


Nx



— dataset_497.dat

NGx



— dataset_497.dat