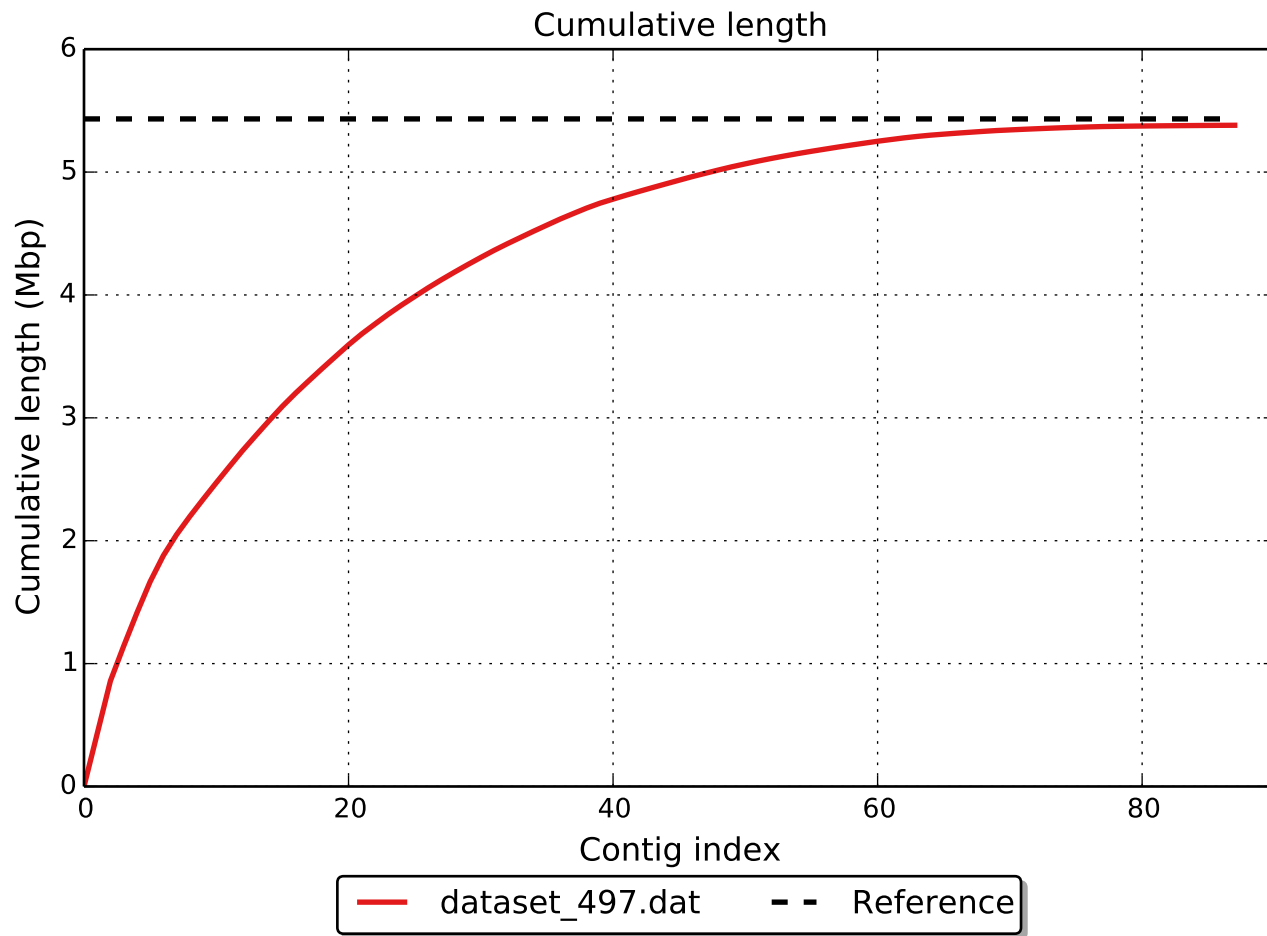
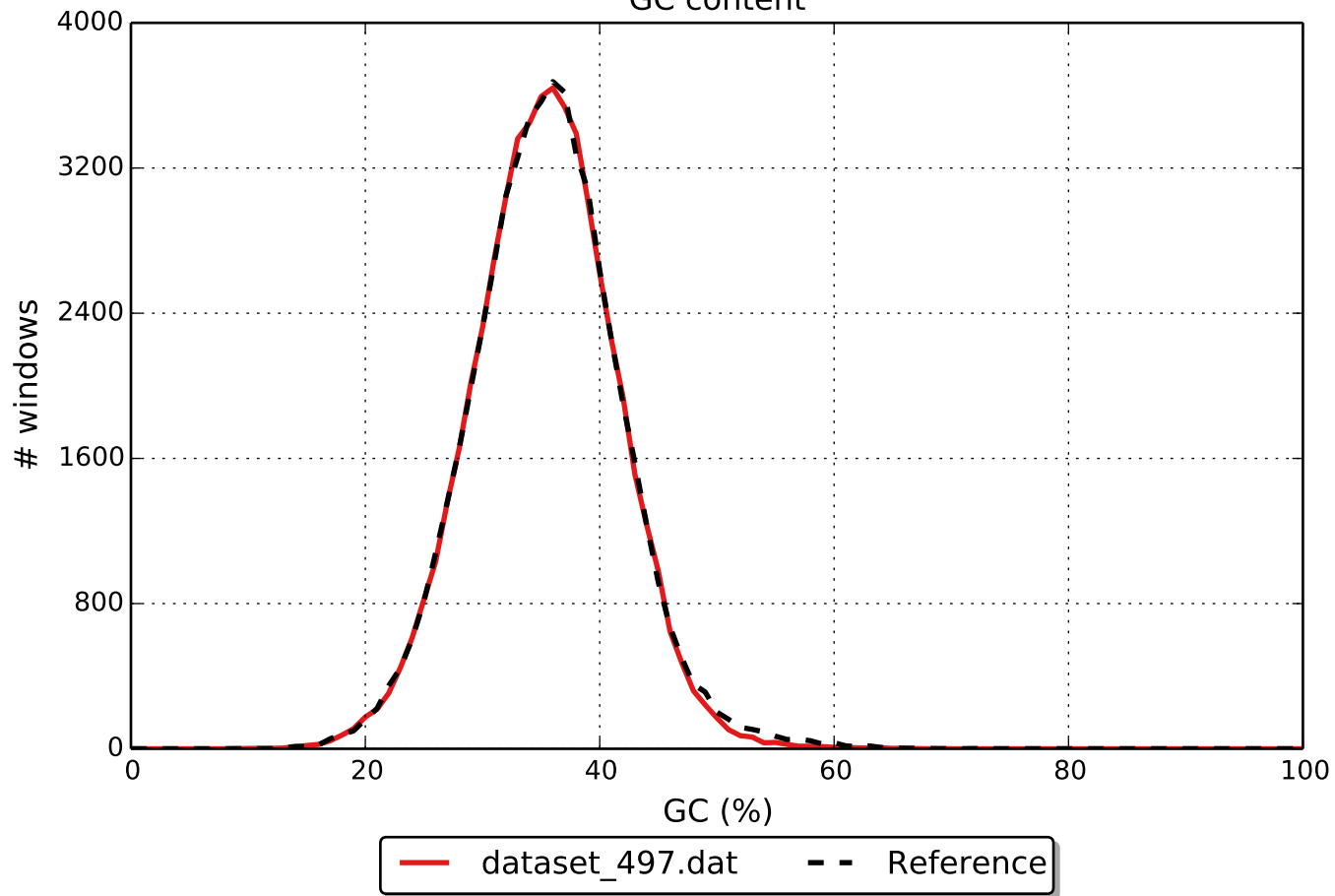


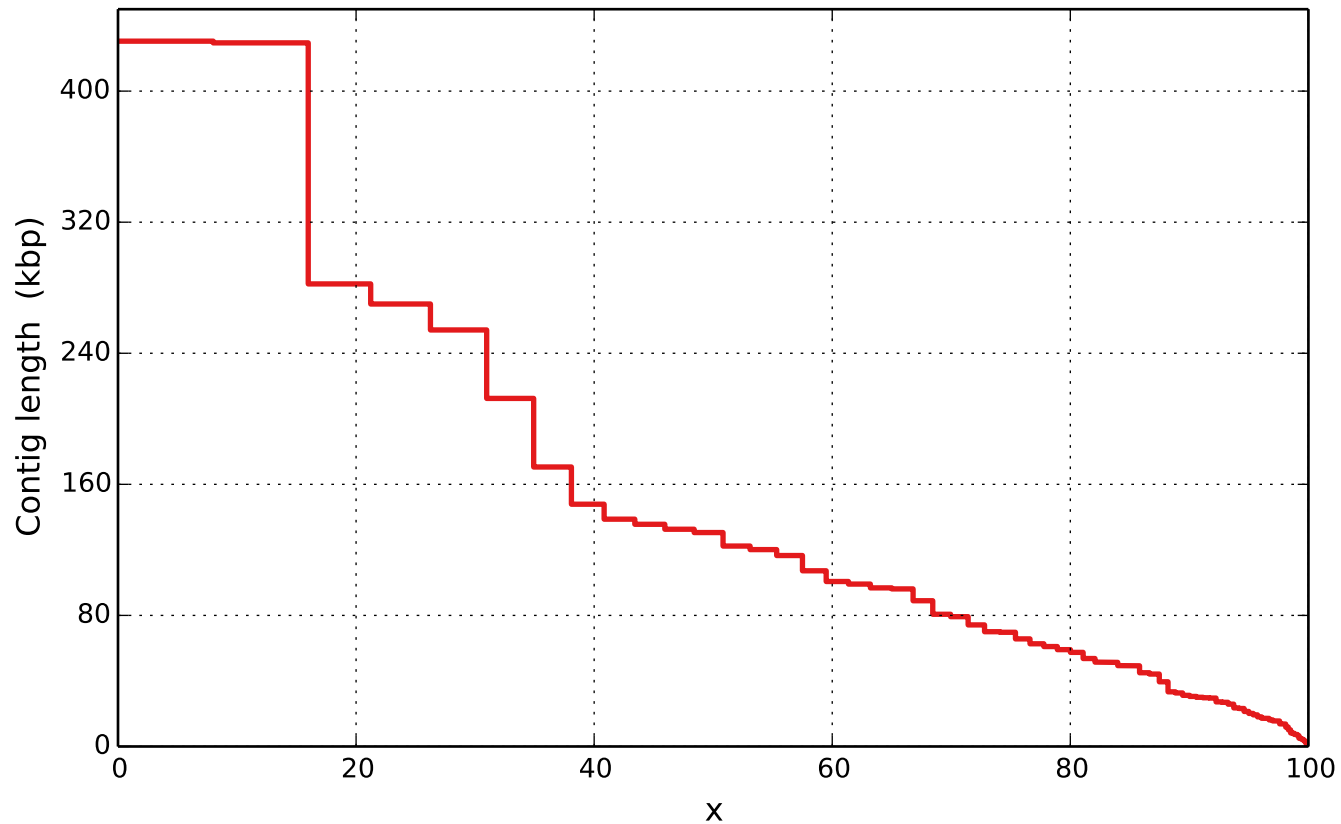
Contig



GC content

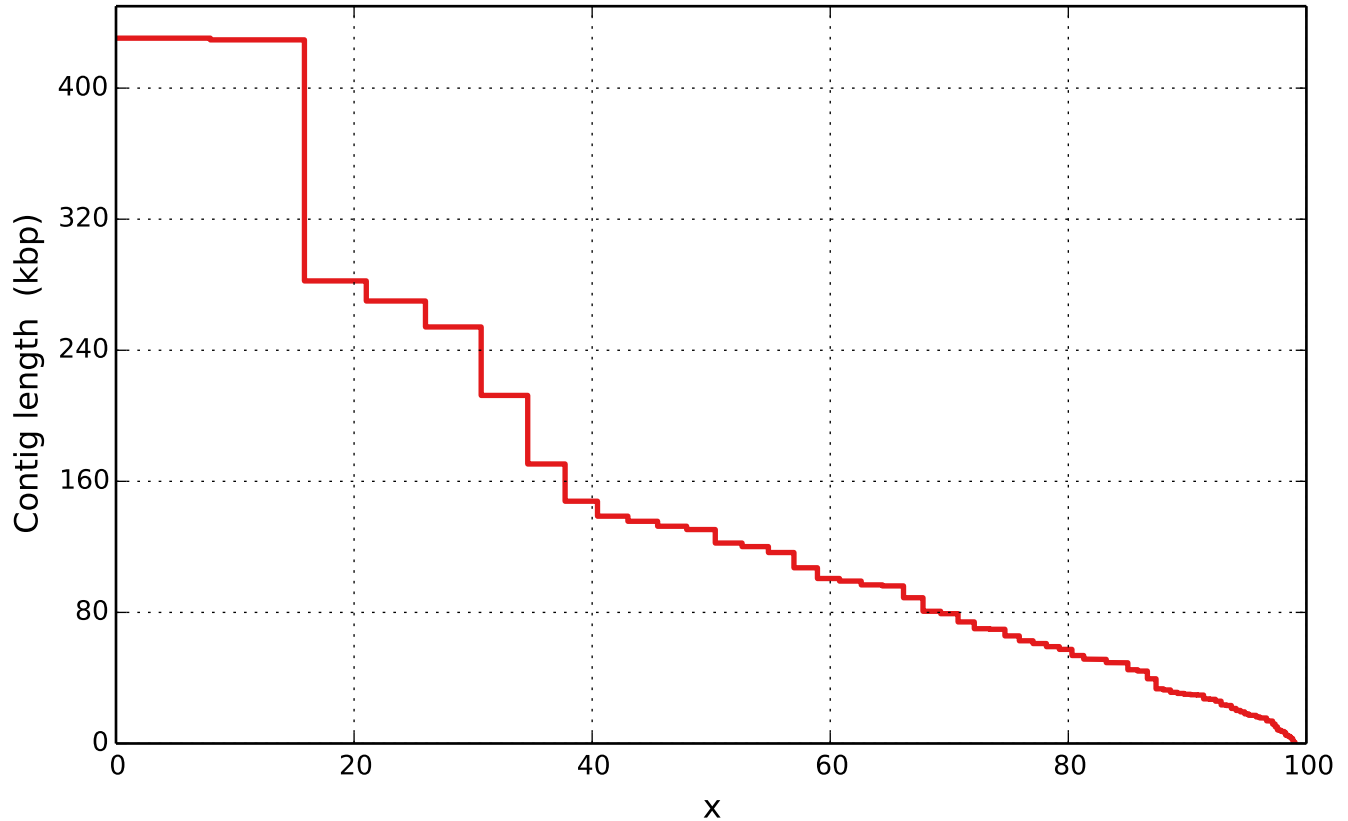


Nx

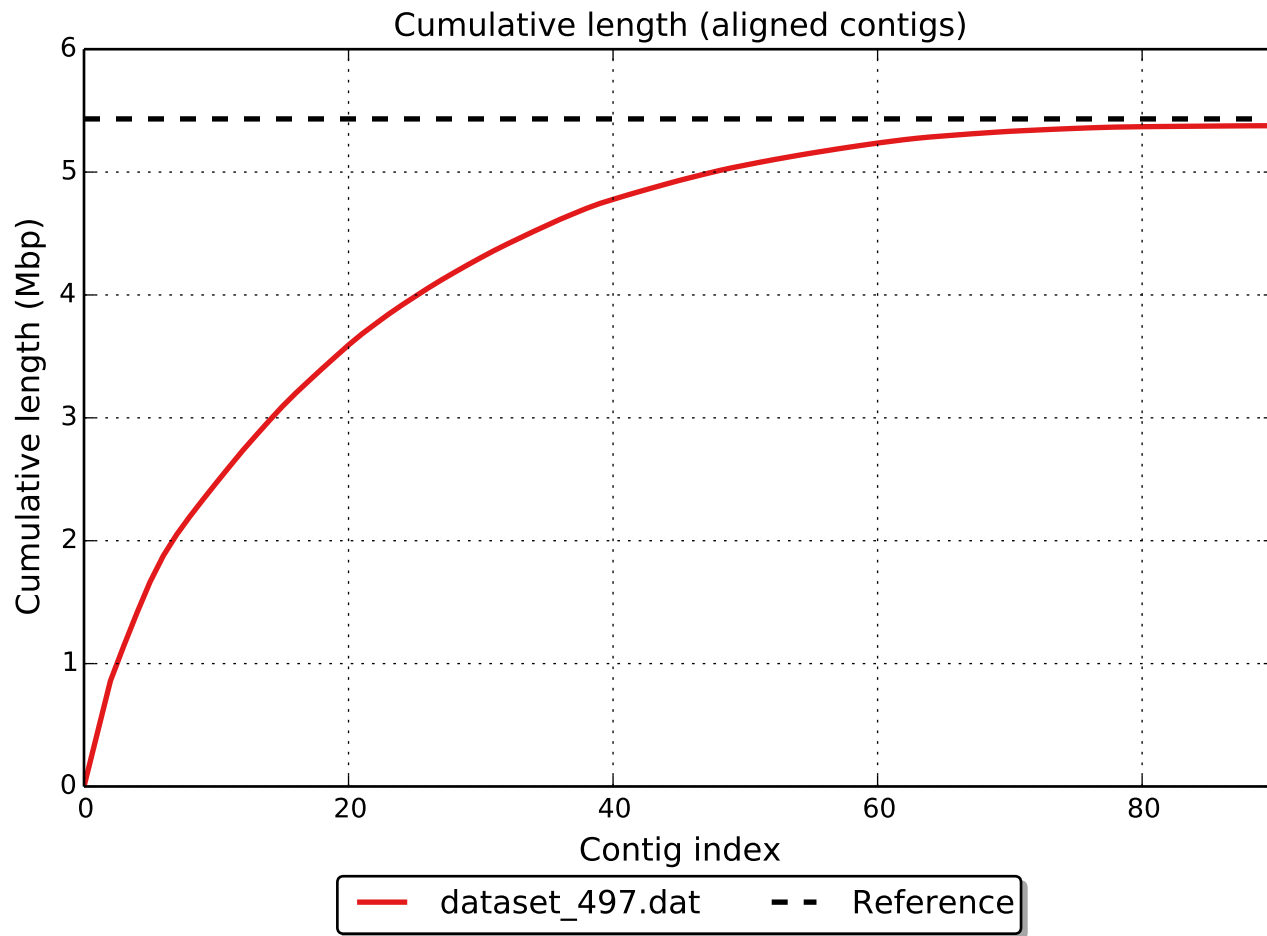


dataset\_497.dat

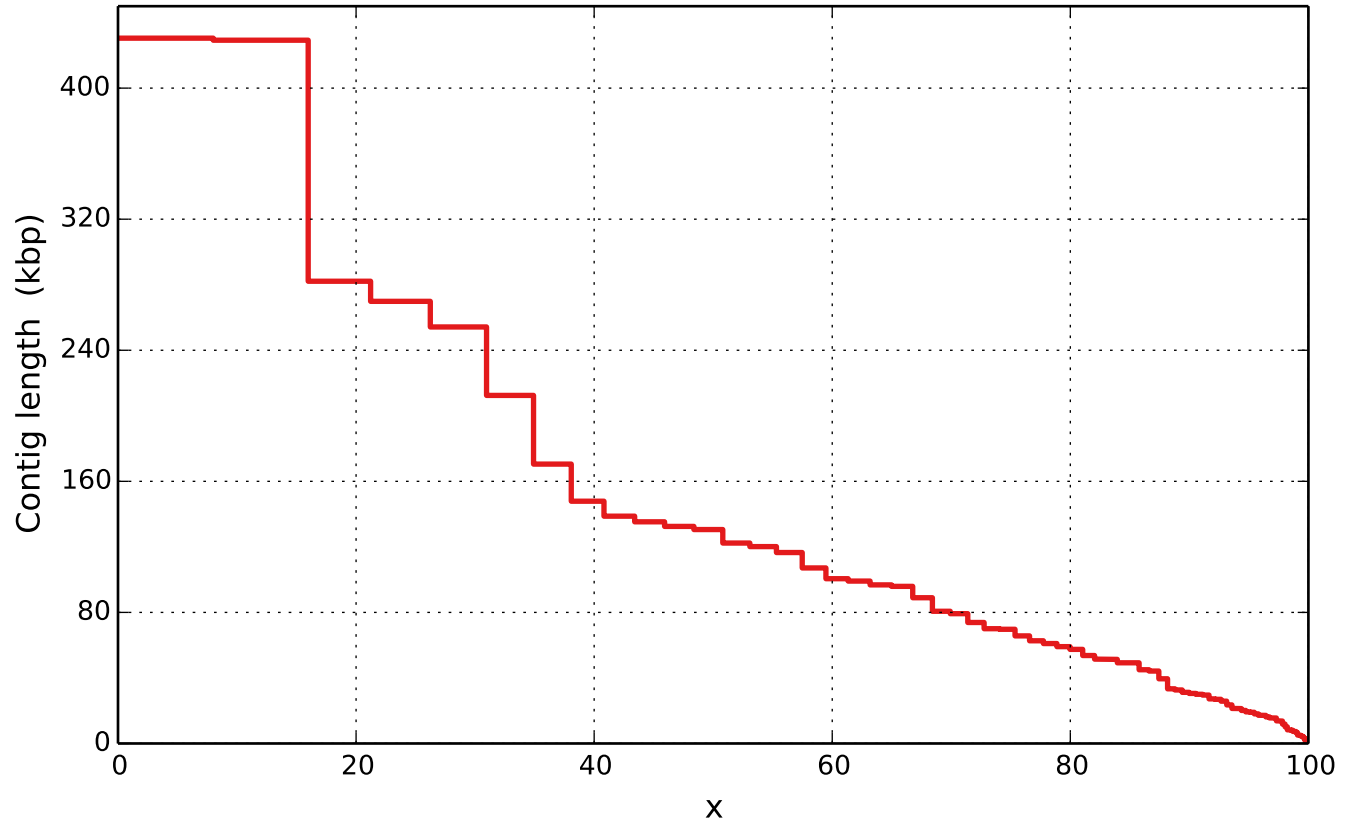
NGx



— dataset\_497.dat

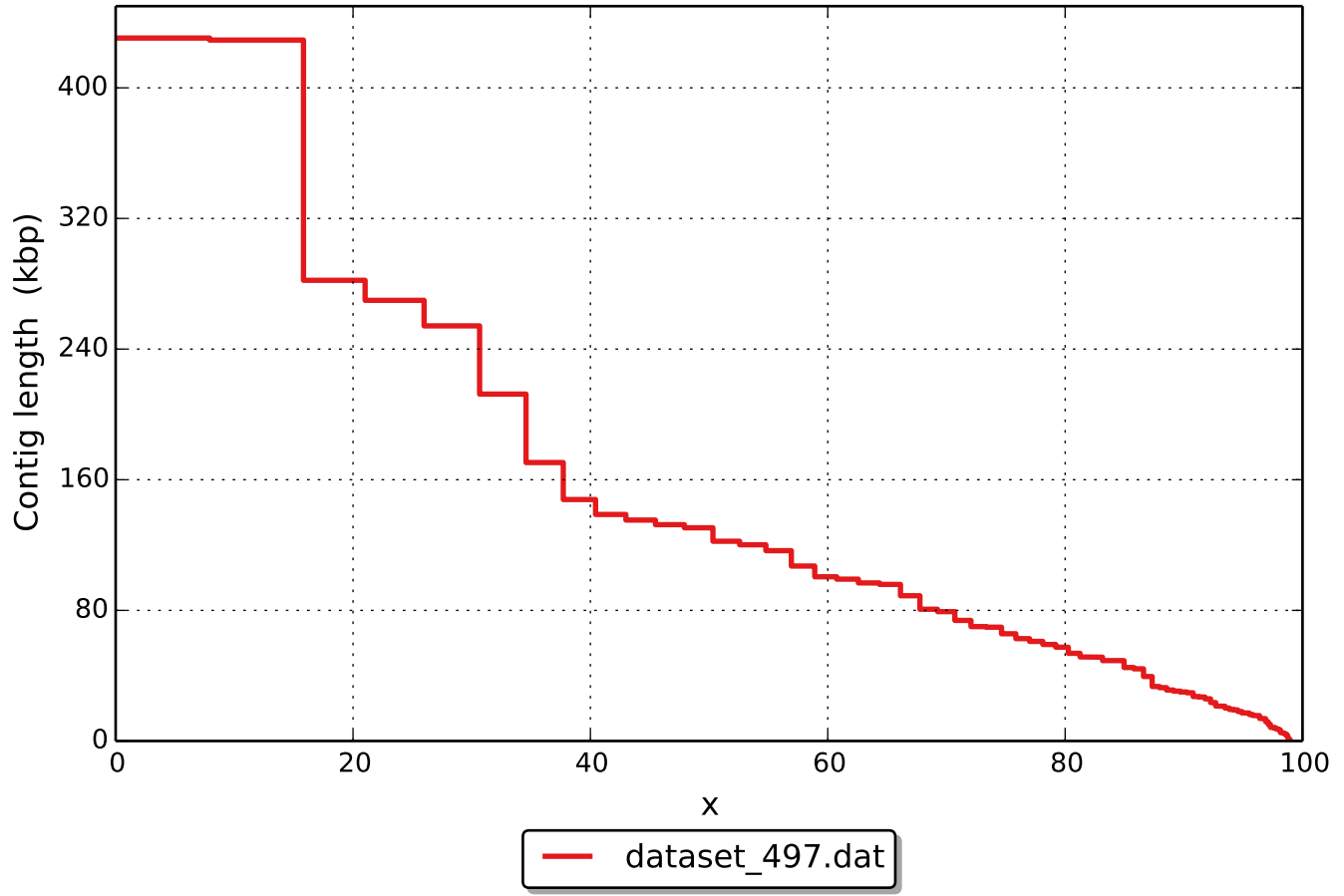


NAx



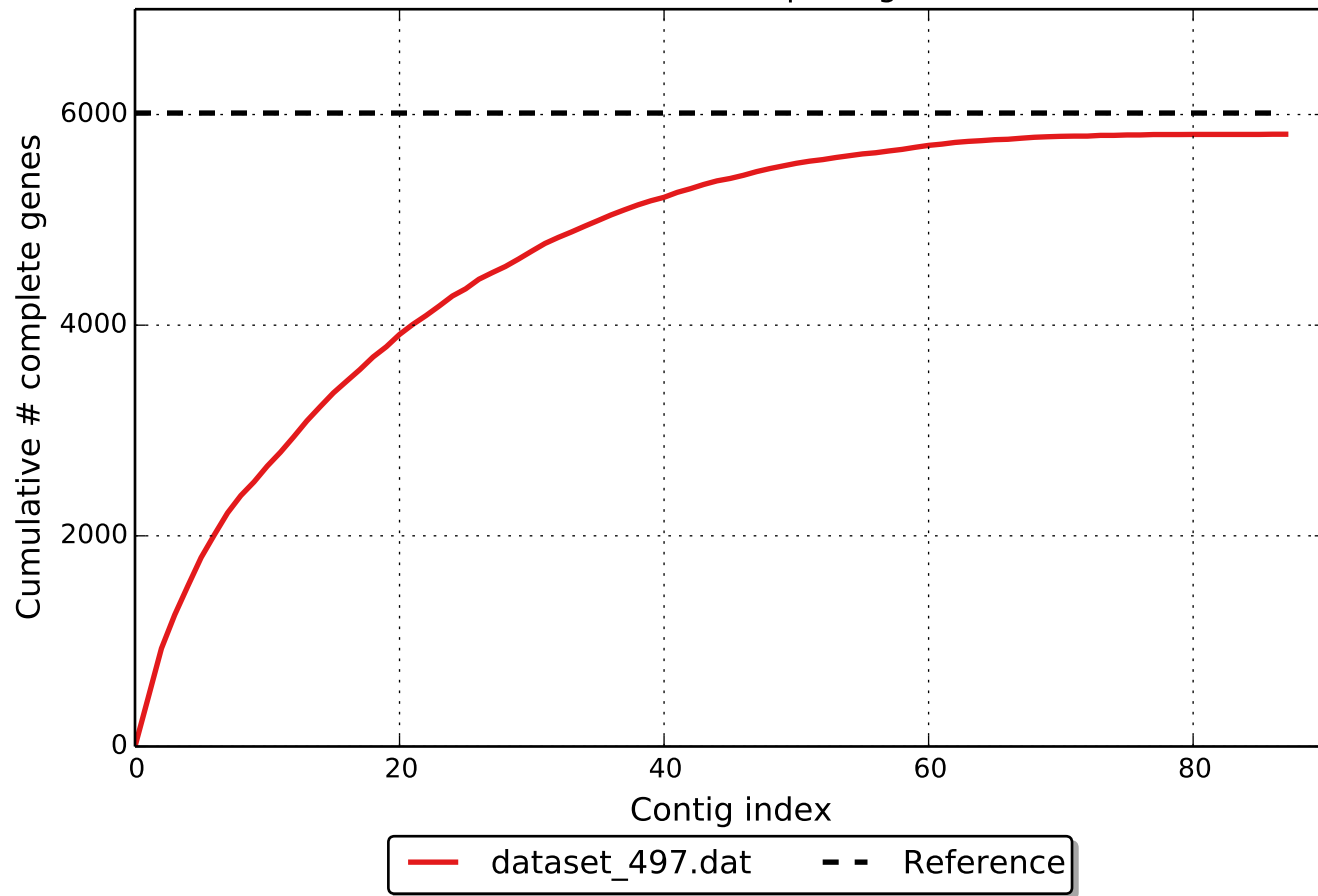
— dataset\_497.dat

NGAx

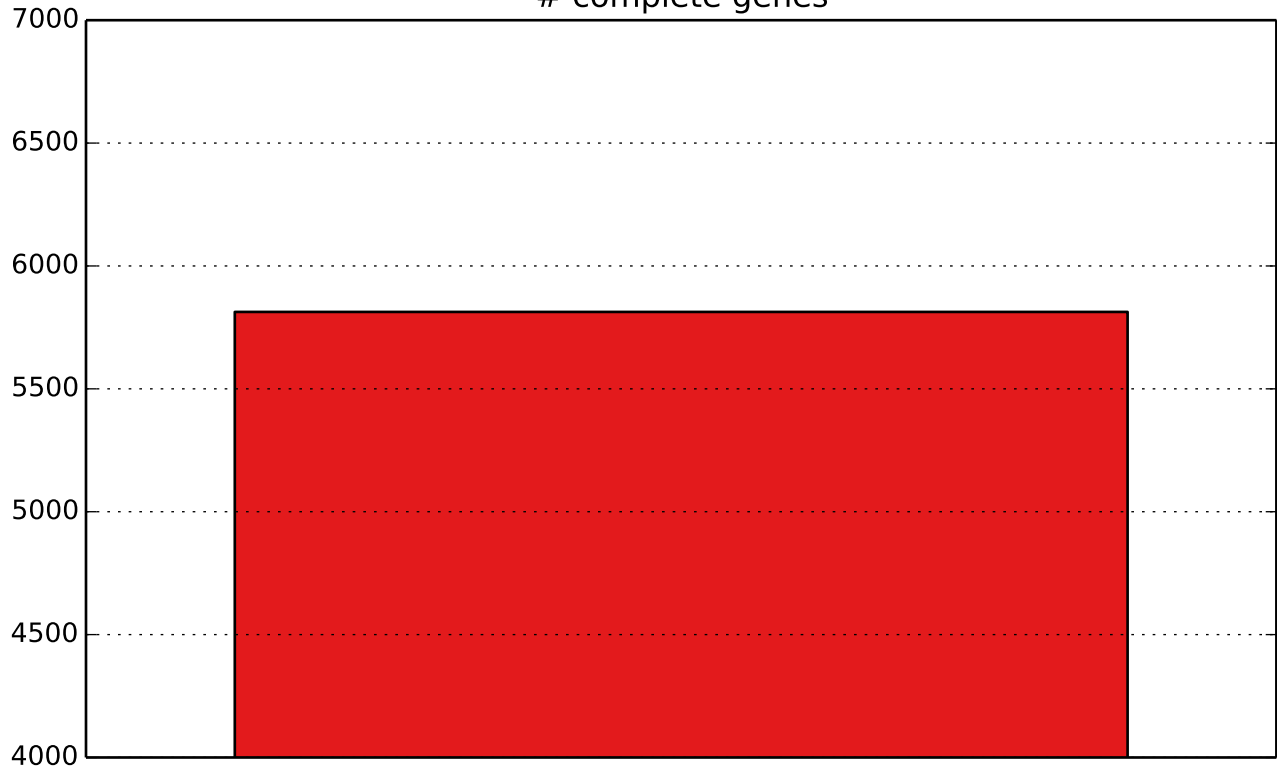




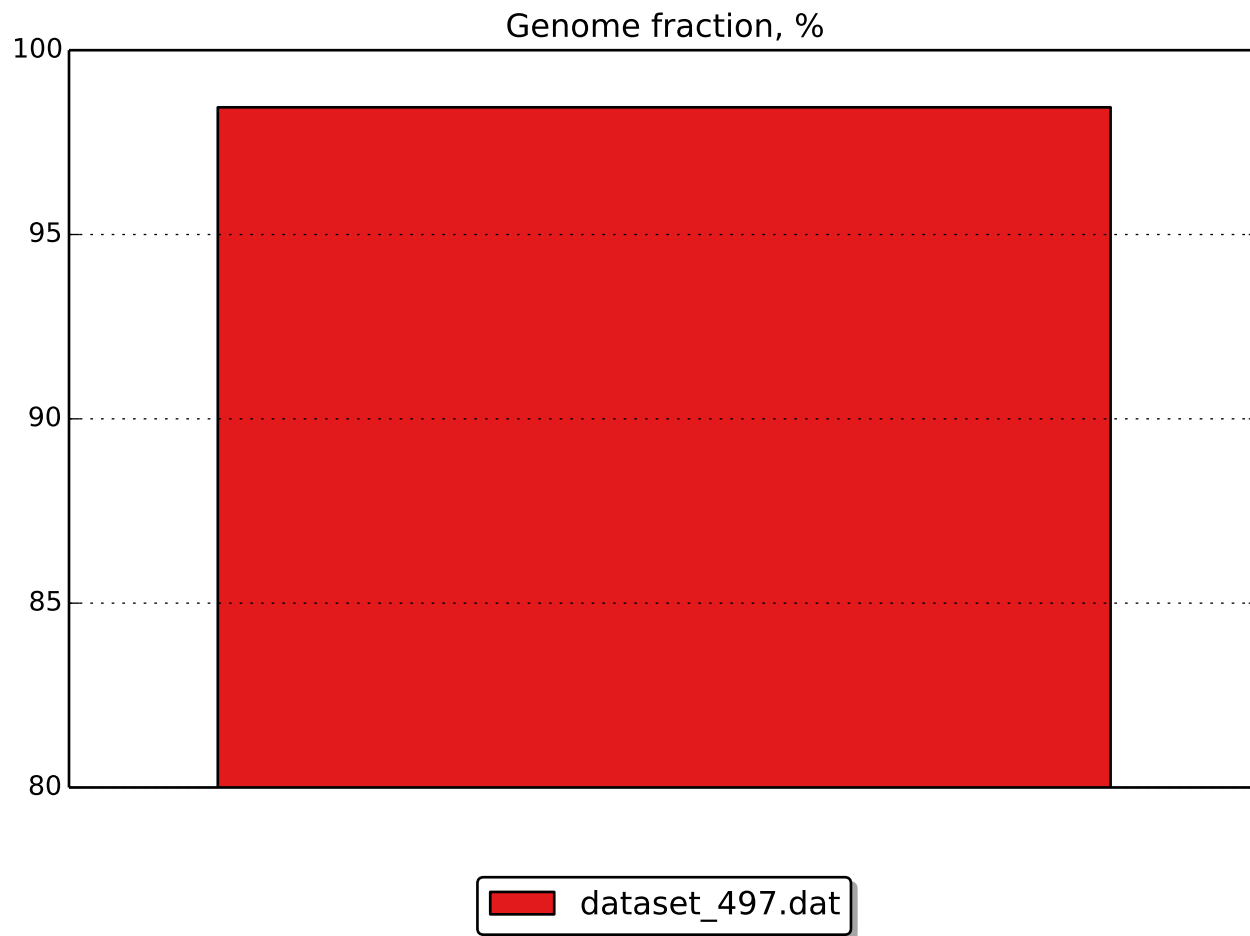
Cumulative # complete genes



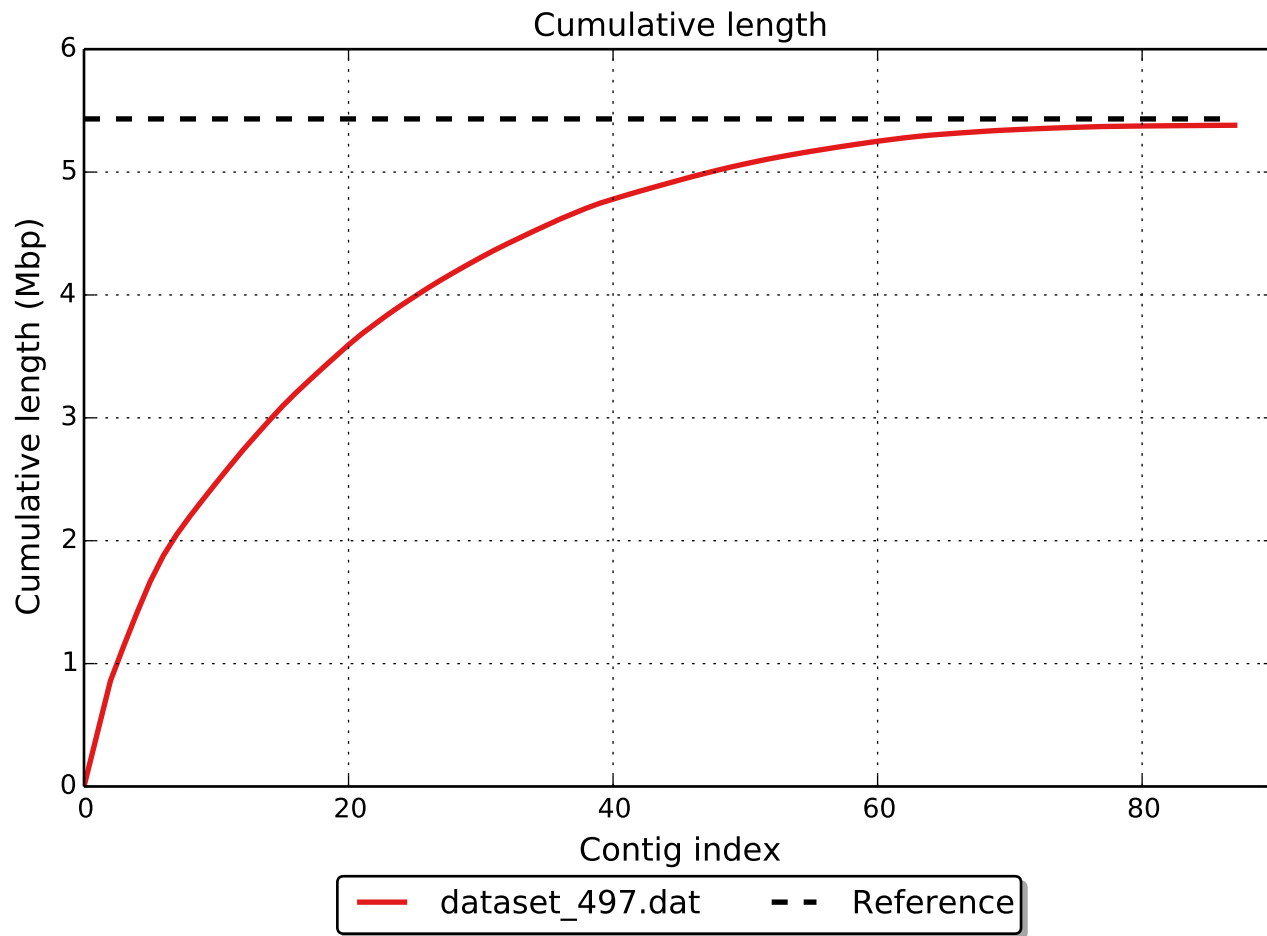
# complete genes



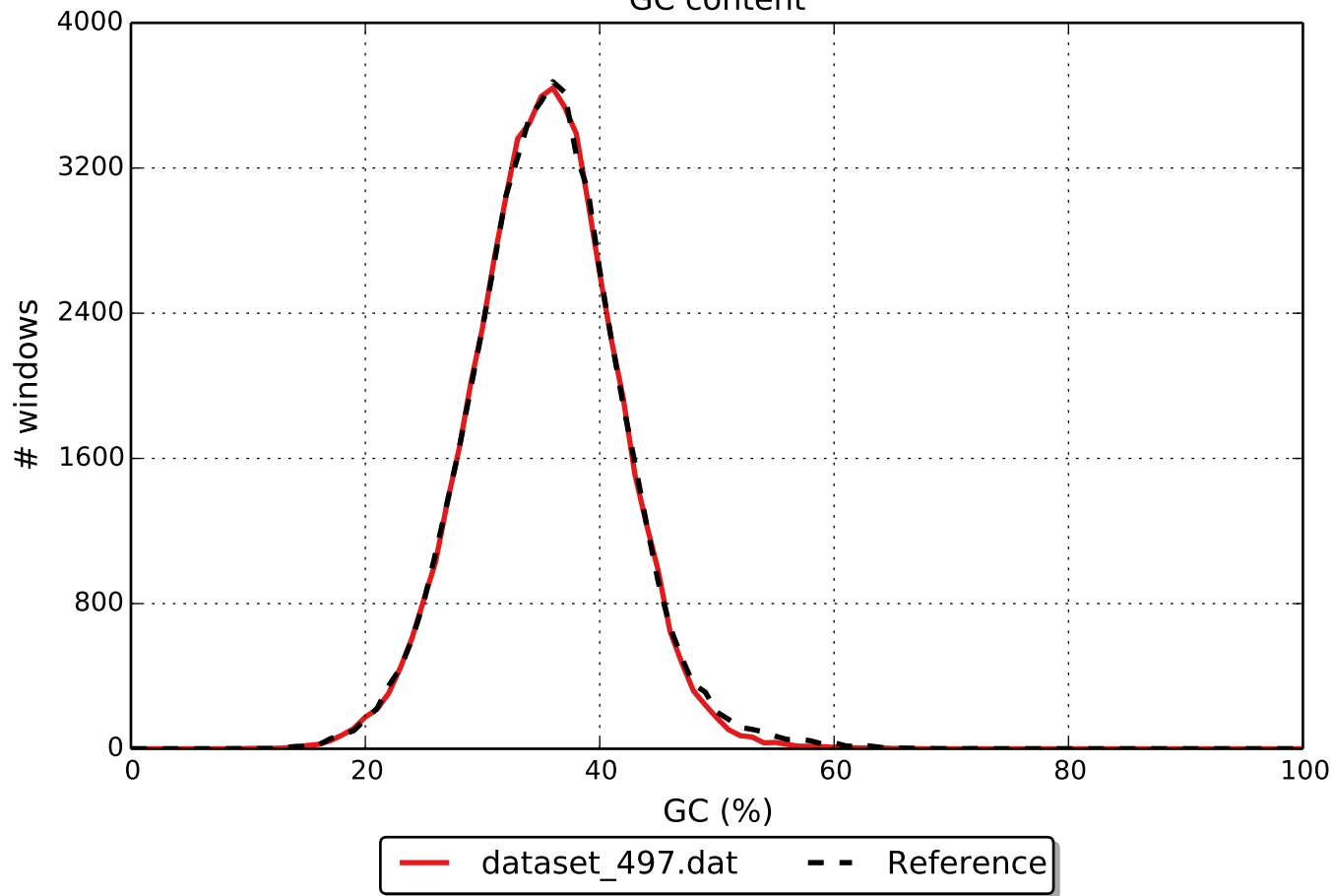
dataset\_497.dat



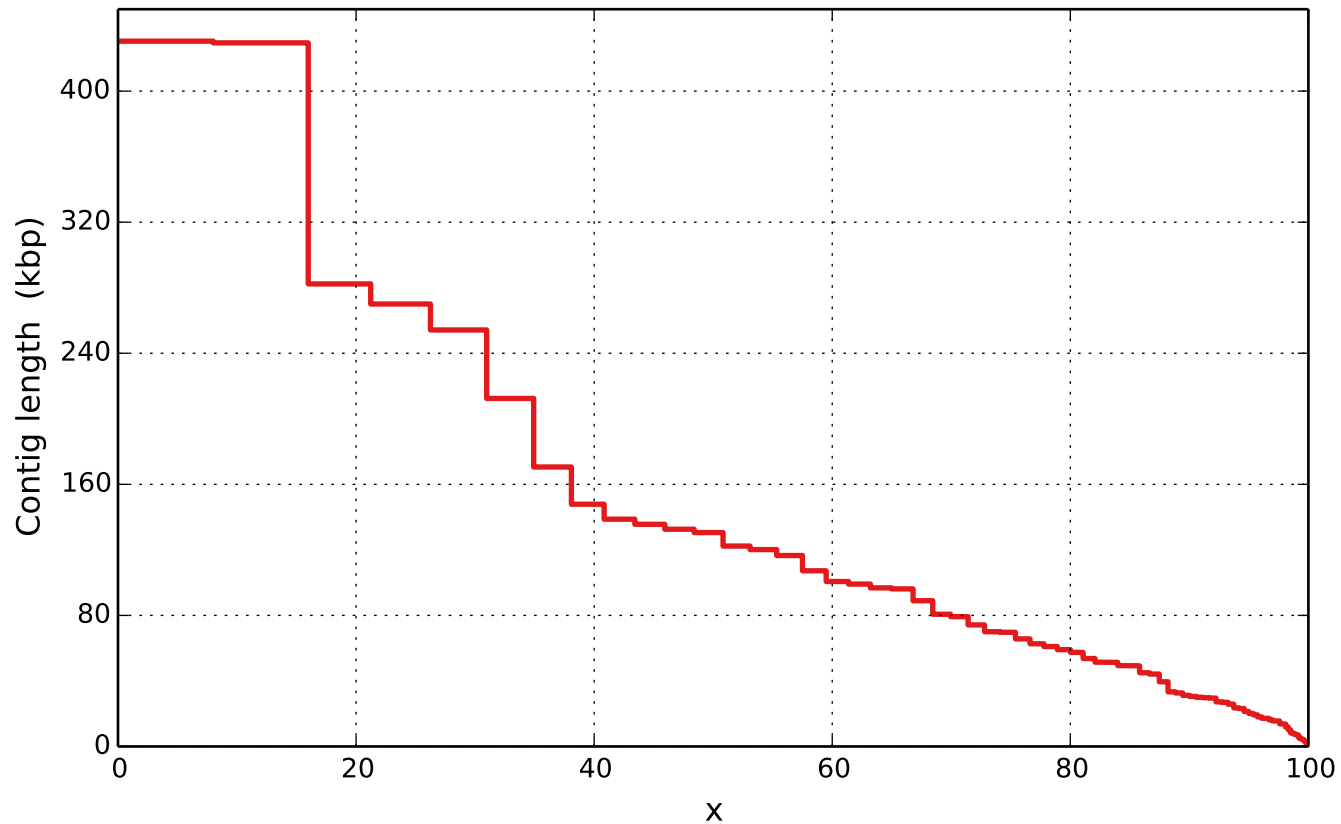
Contig



GC content

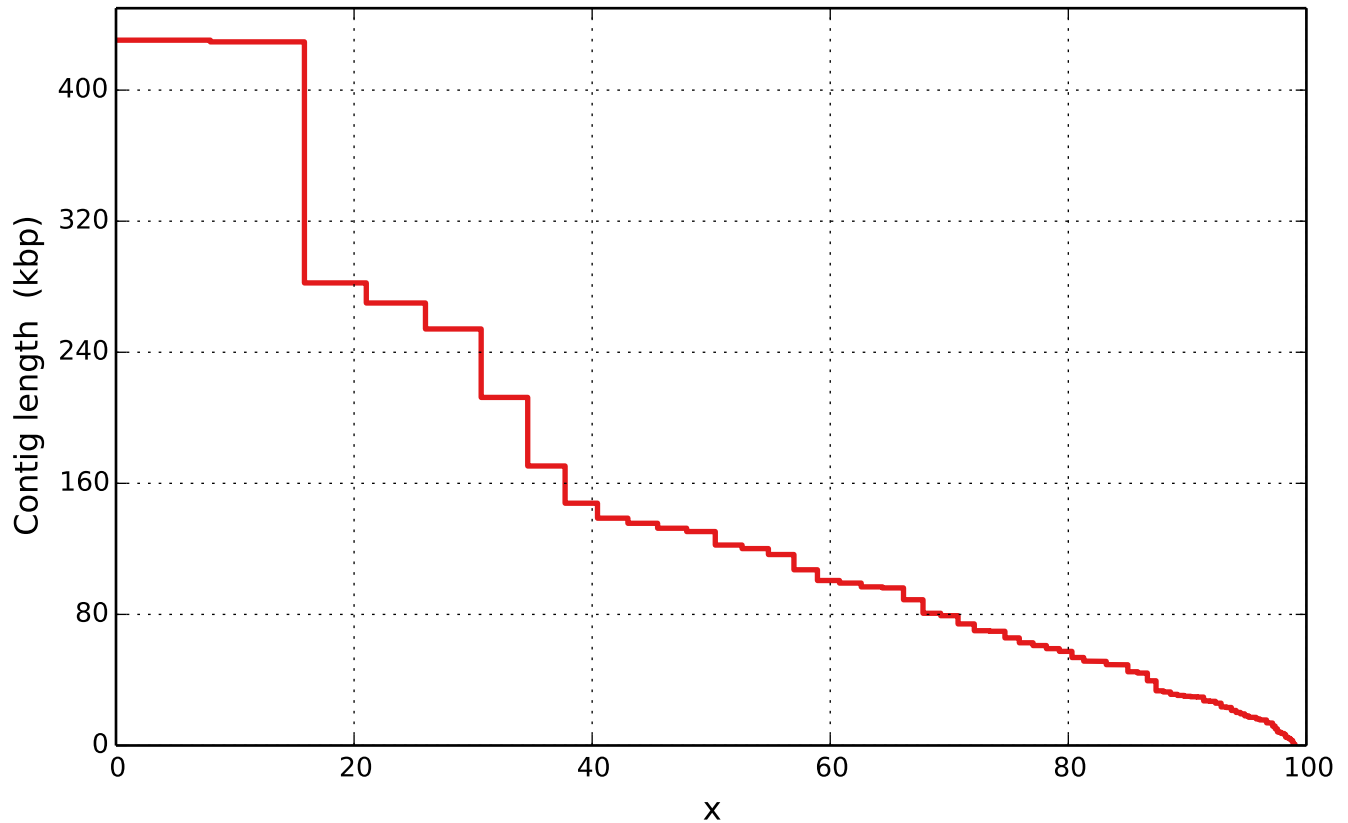


Nx



— dataset\_497.dat

NGx

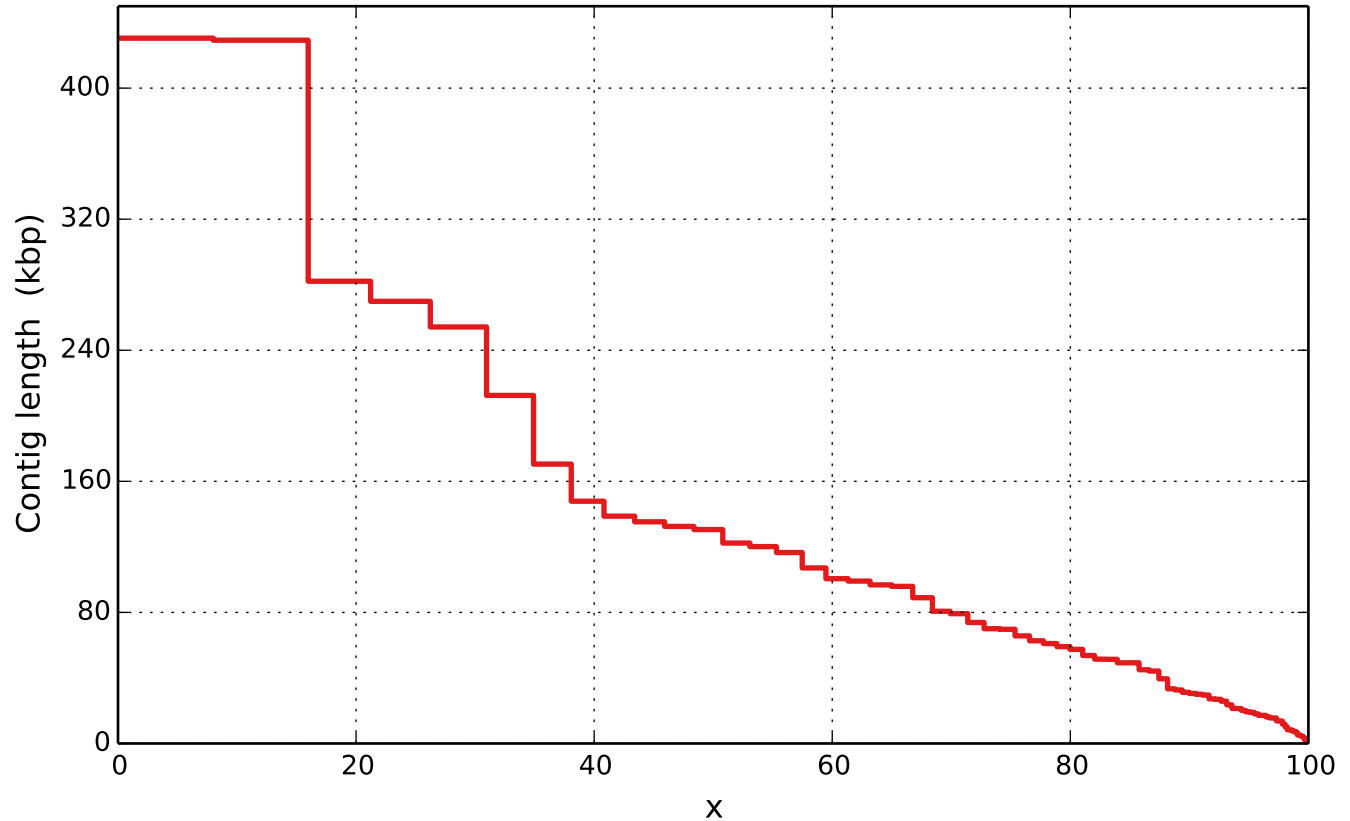


— dataset\_497.dat



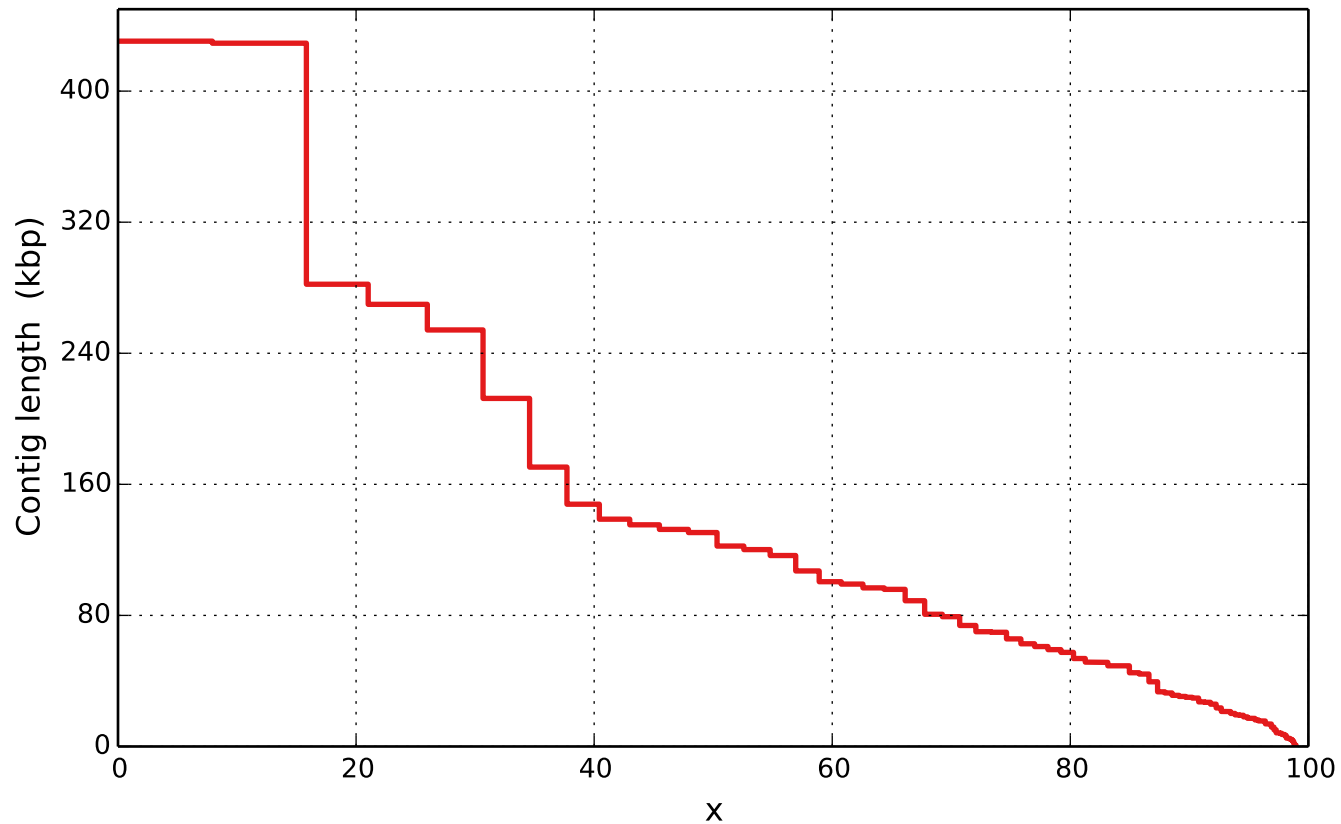


NAx



— dataset\_497.dat

NGAx



— dataset\_497.dat

