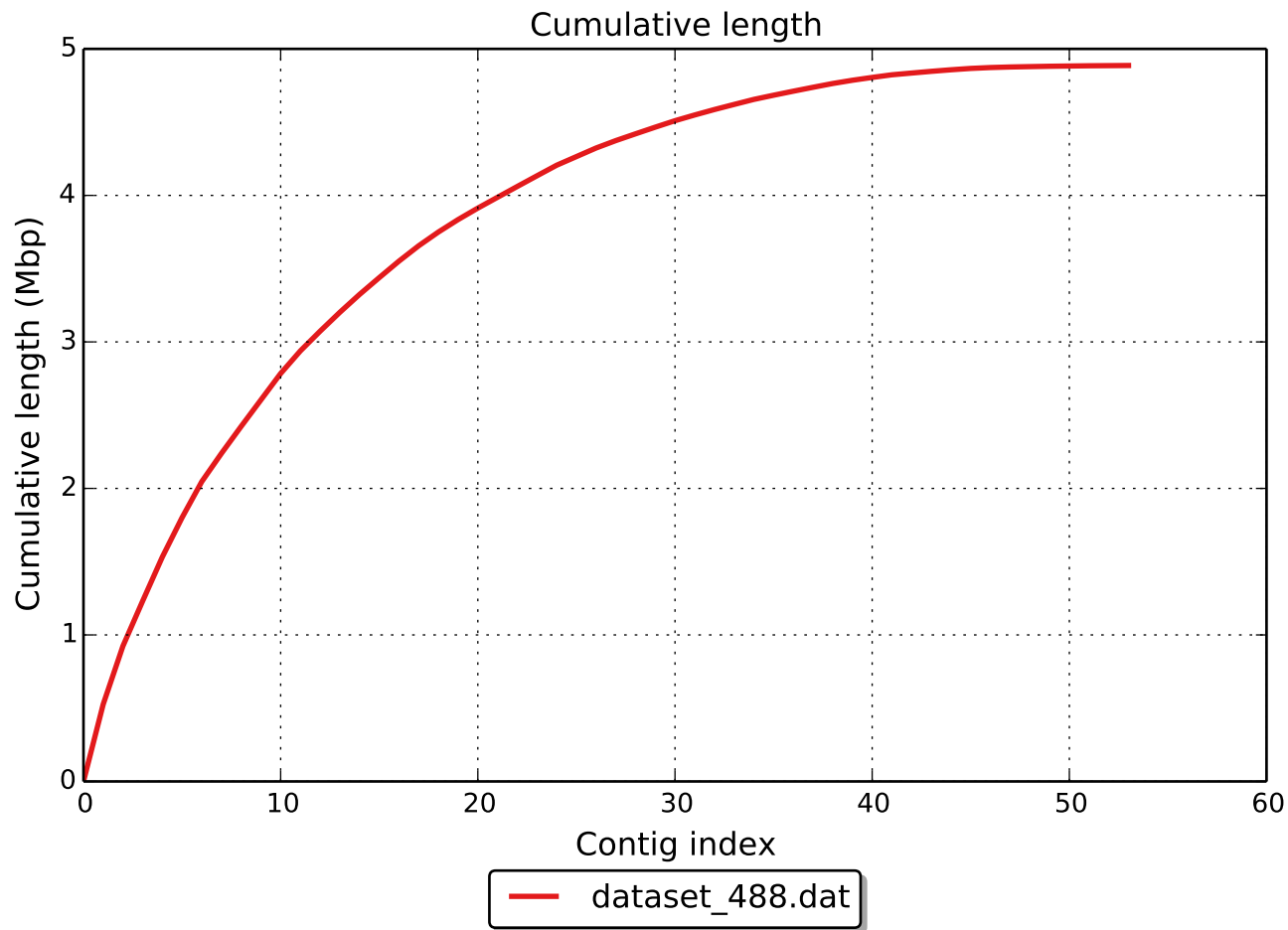
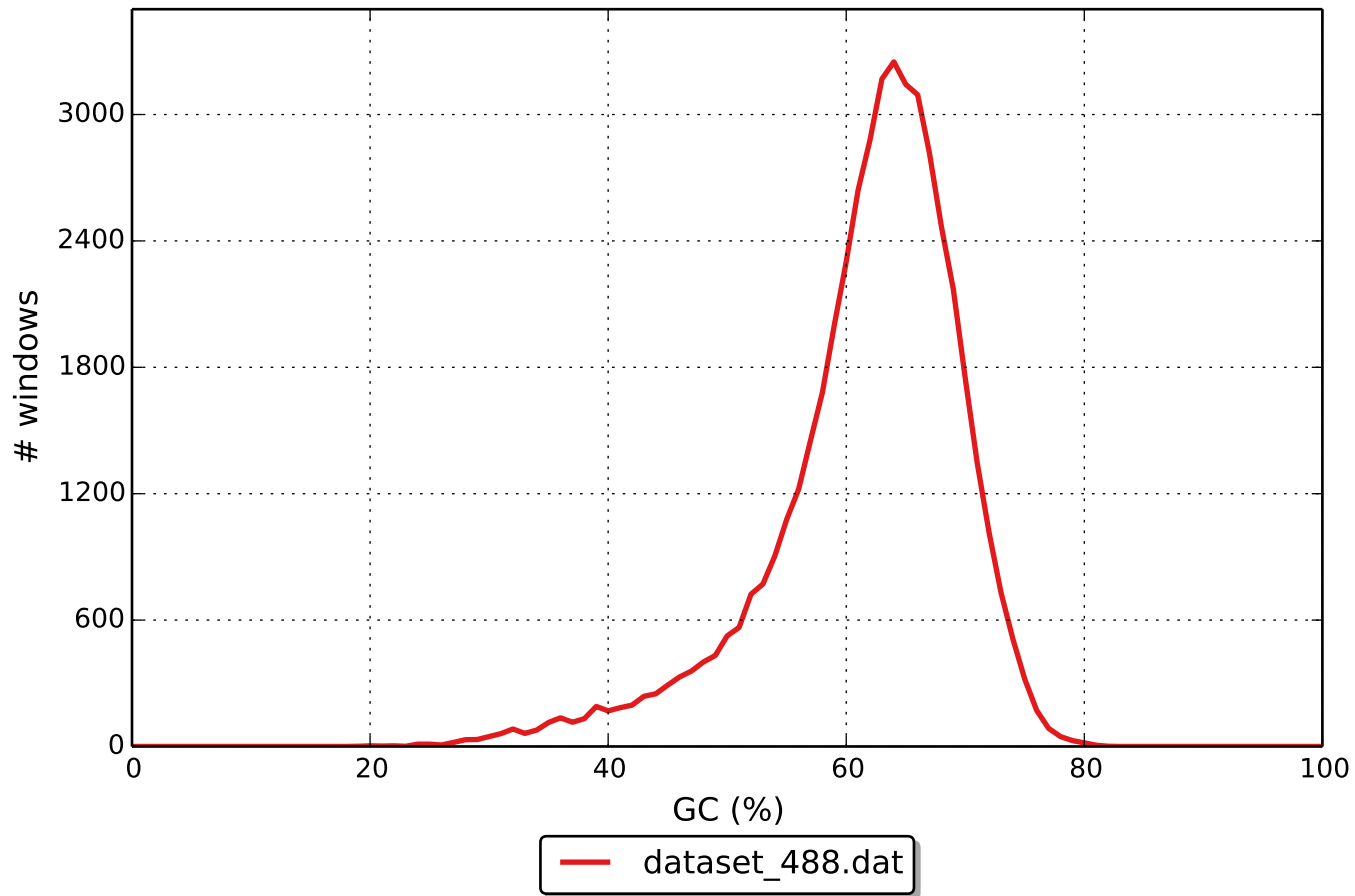


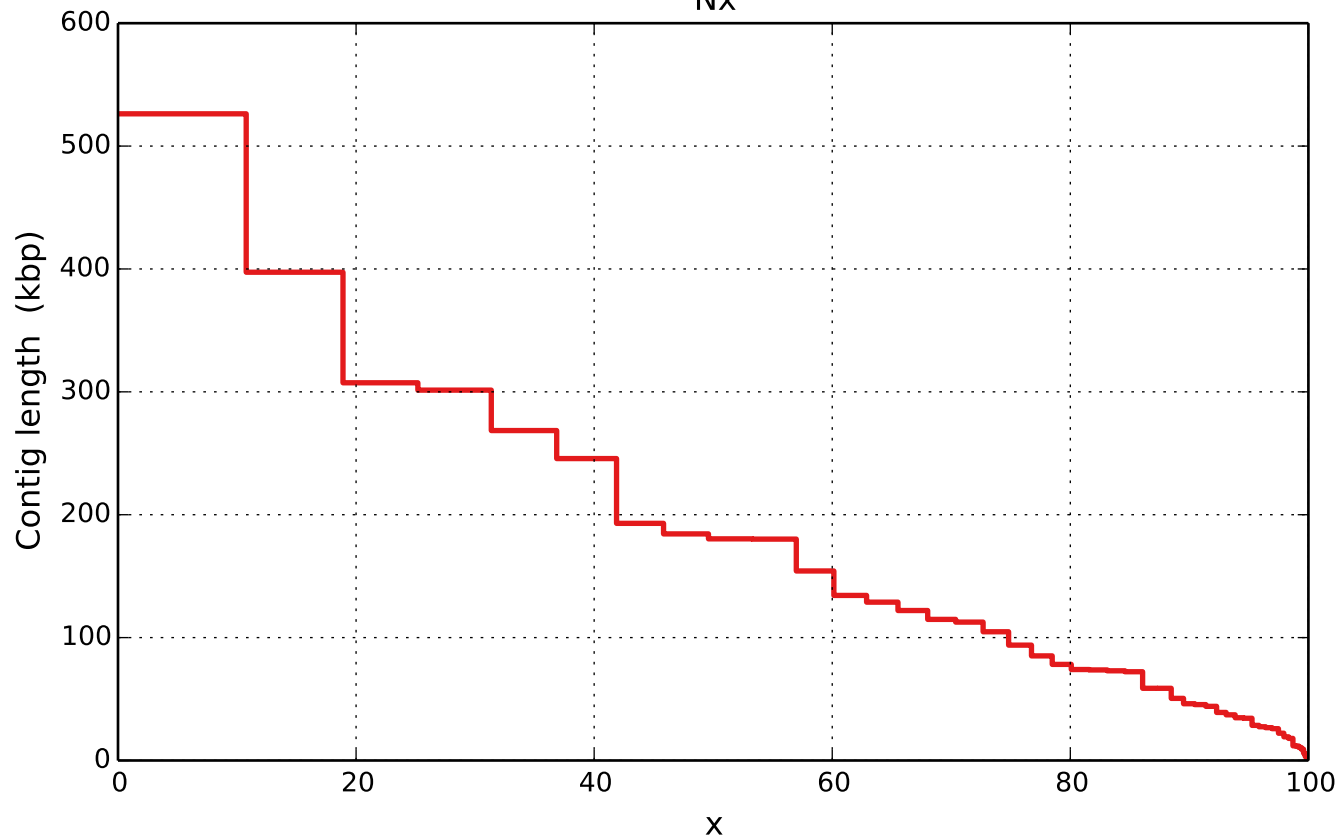
Contig



GC content



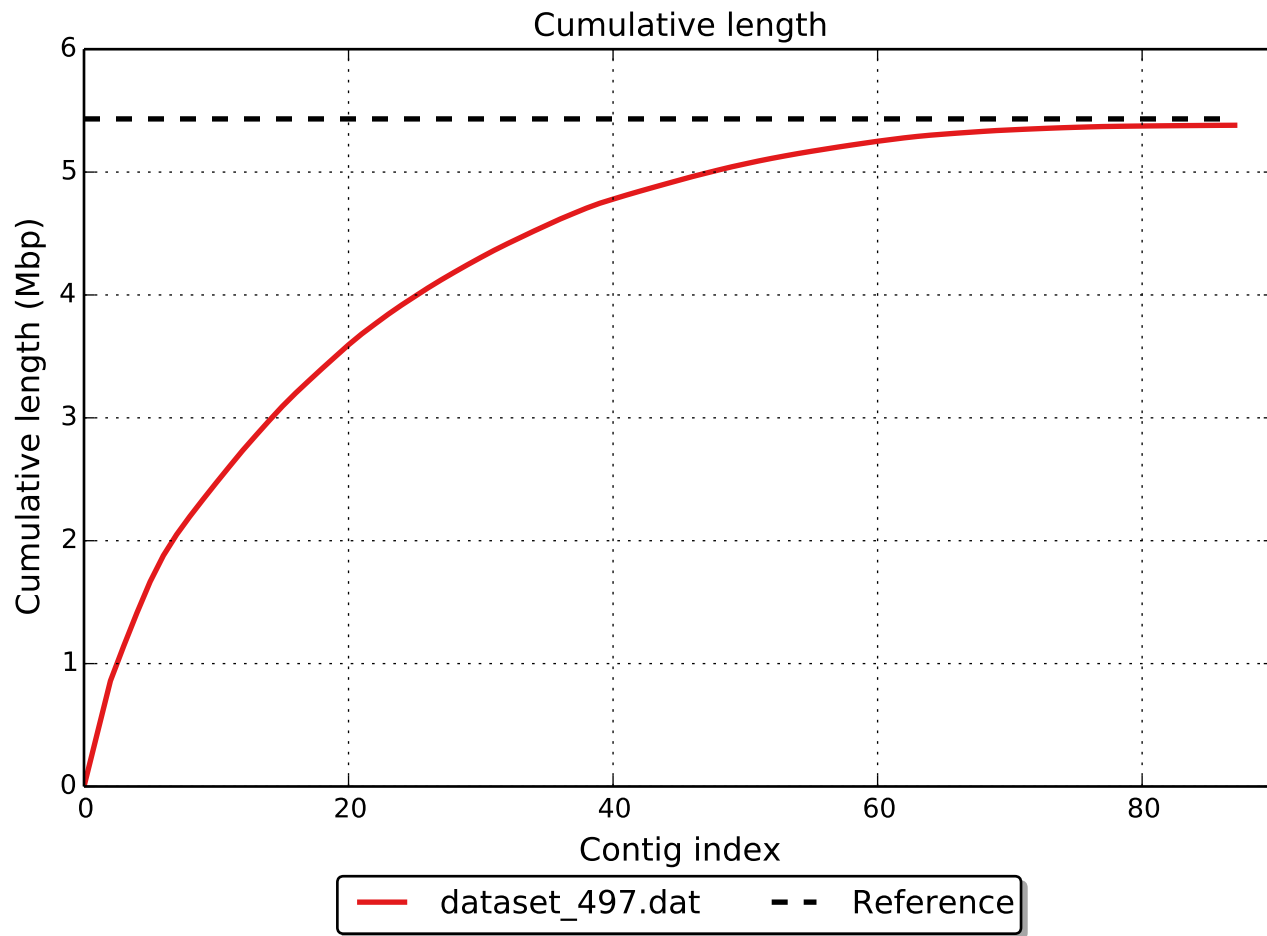
Nx



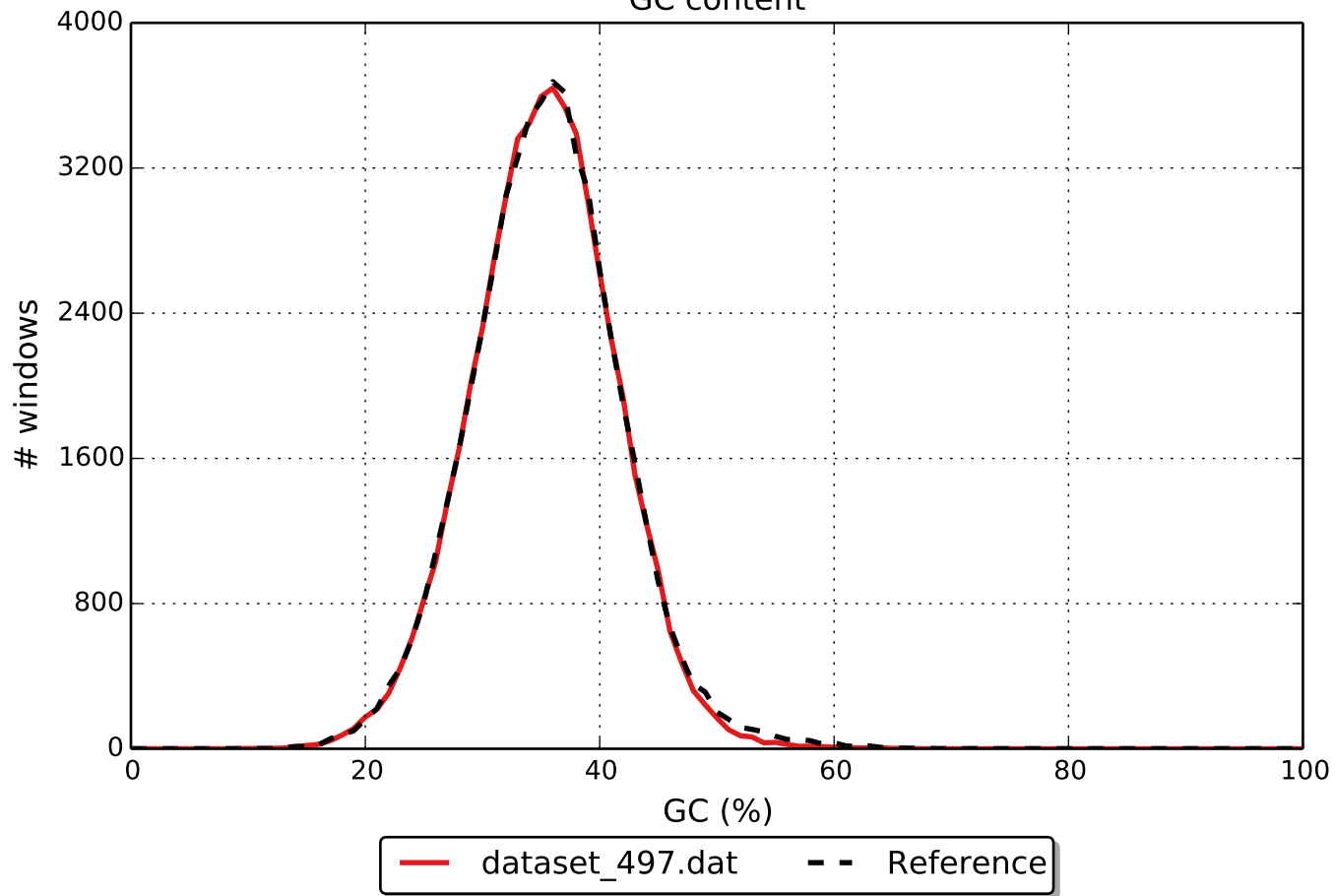
— dataset\_488.dat

Combined\_reference

B\_cereus\_ATCC\_1098



GC content



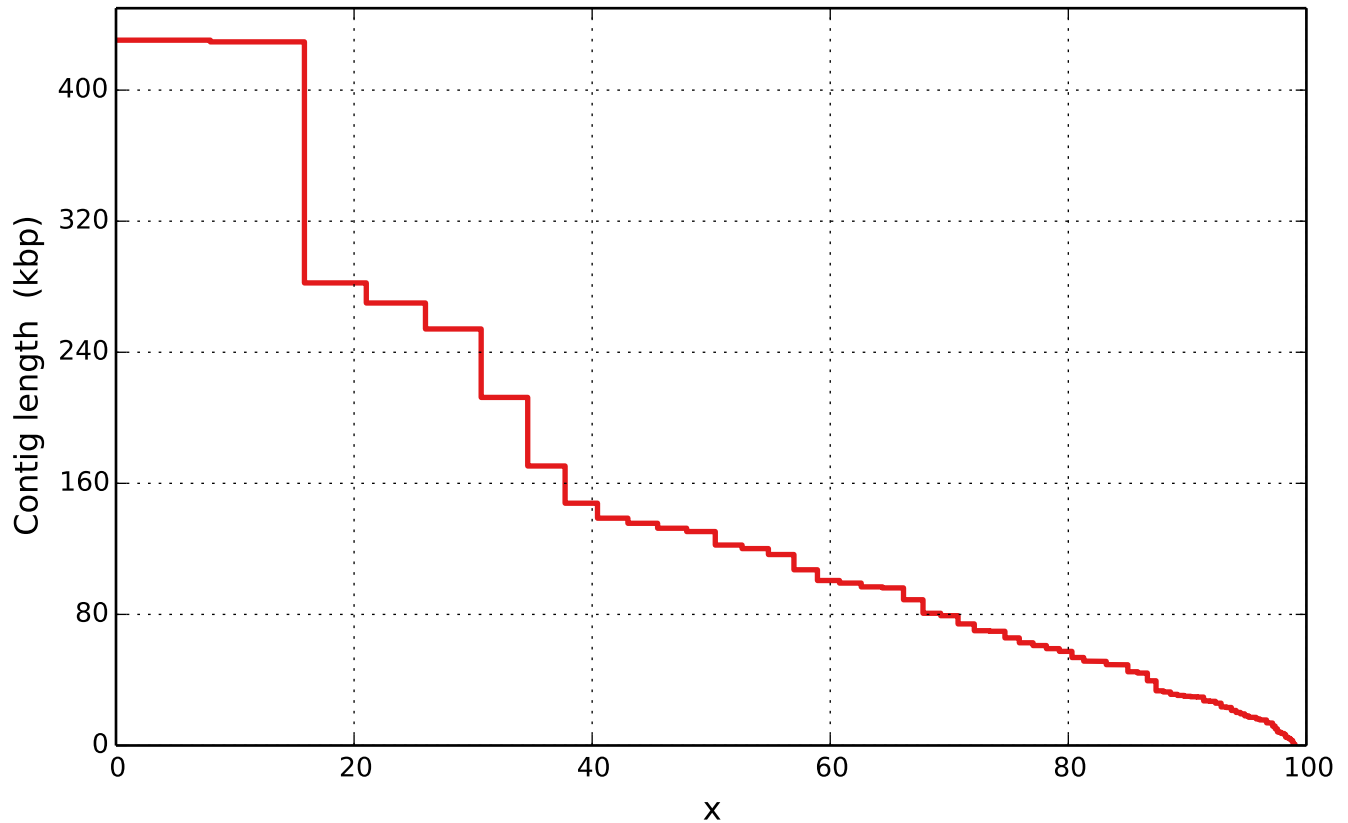


Nx

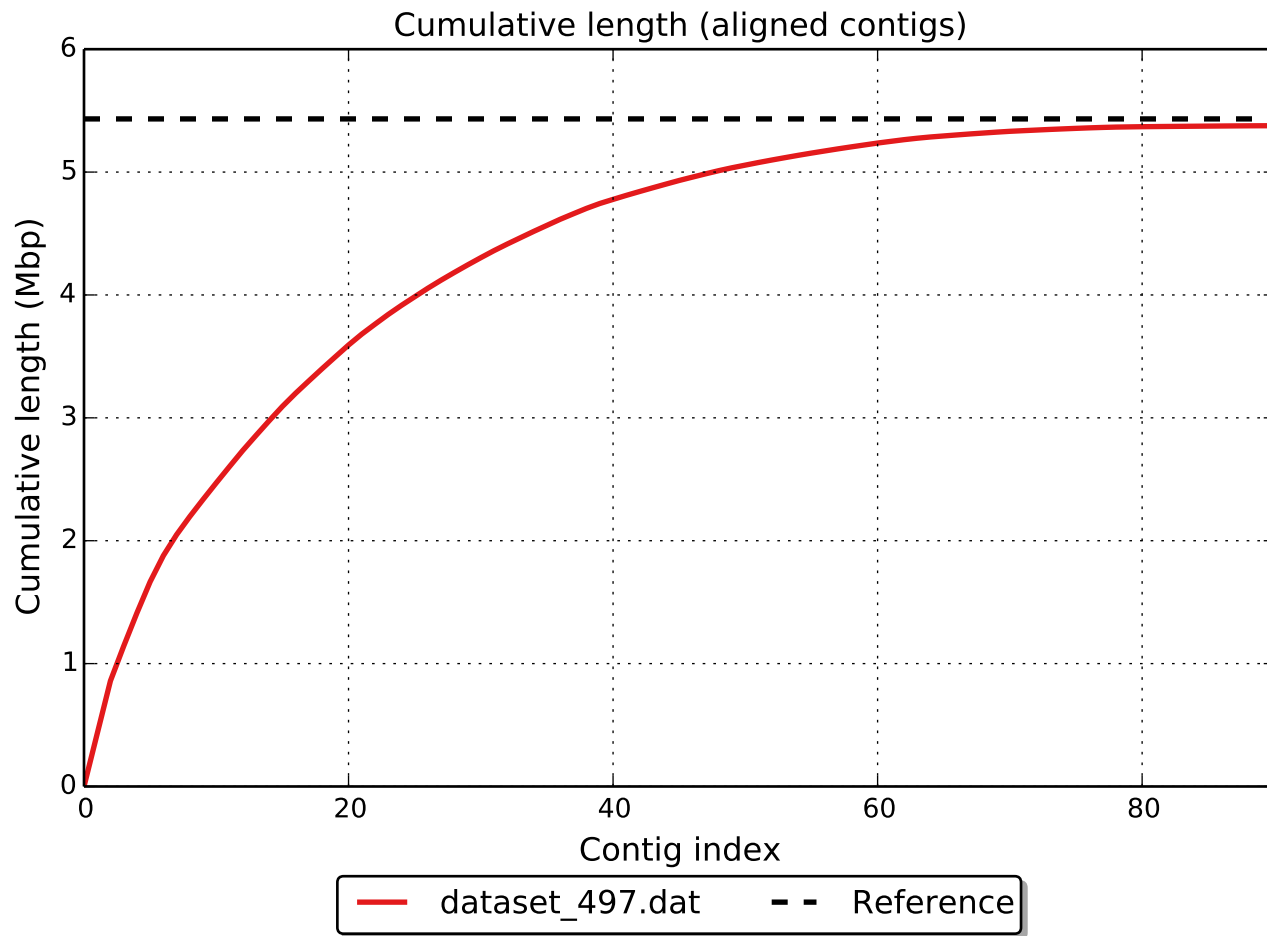


— dataset\_497.dat

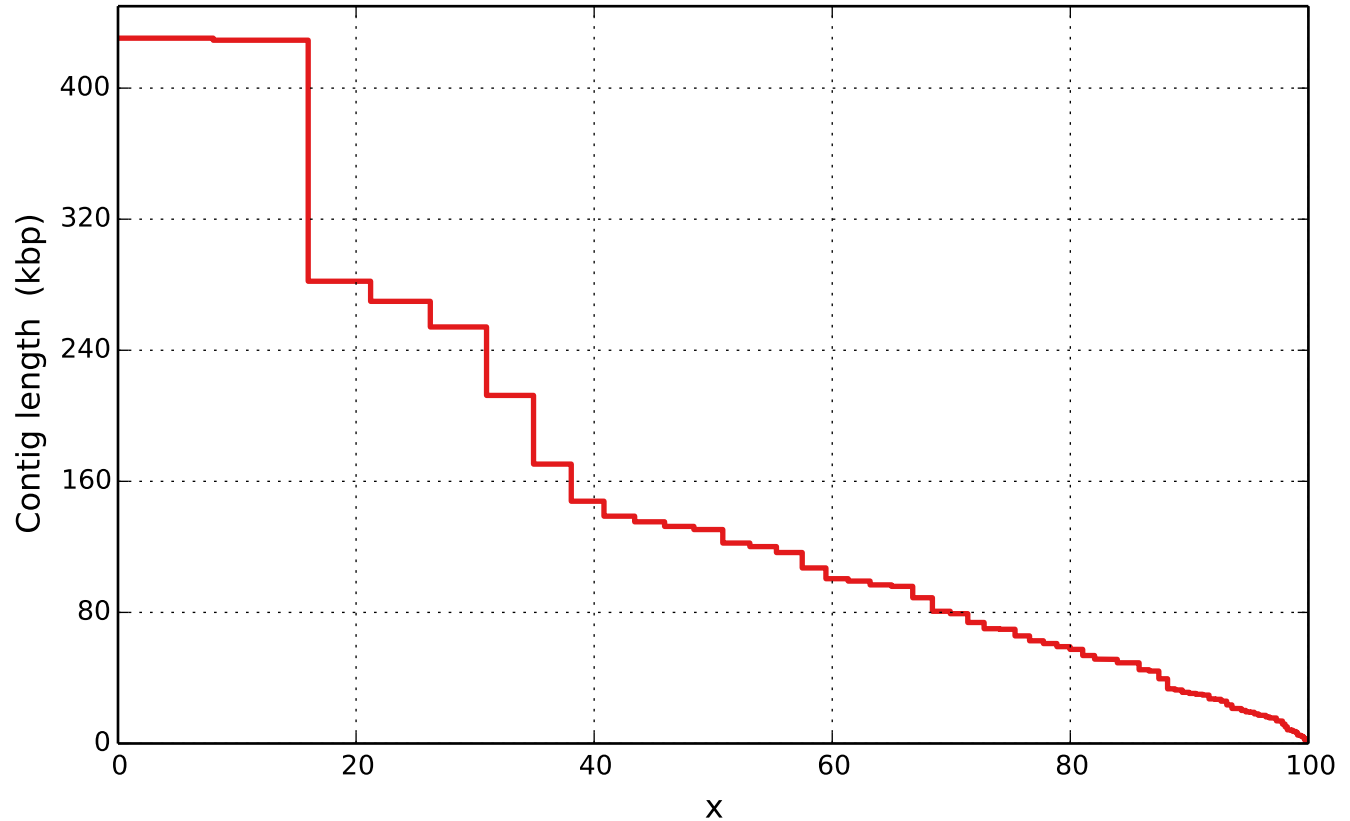
NGx



— dataset\_497.dat

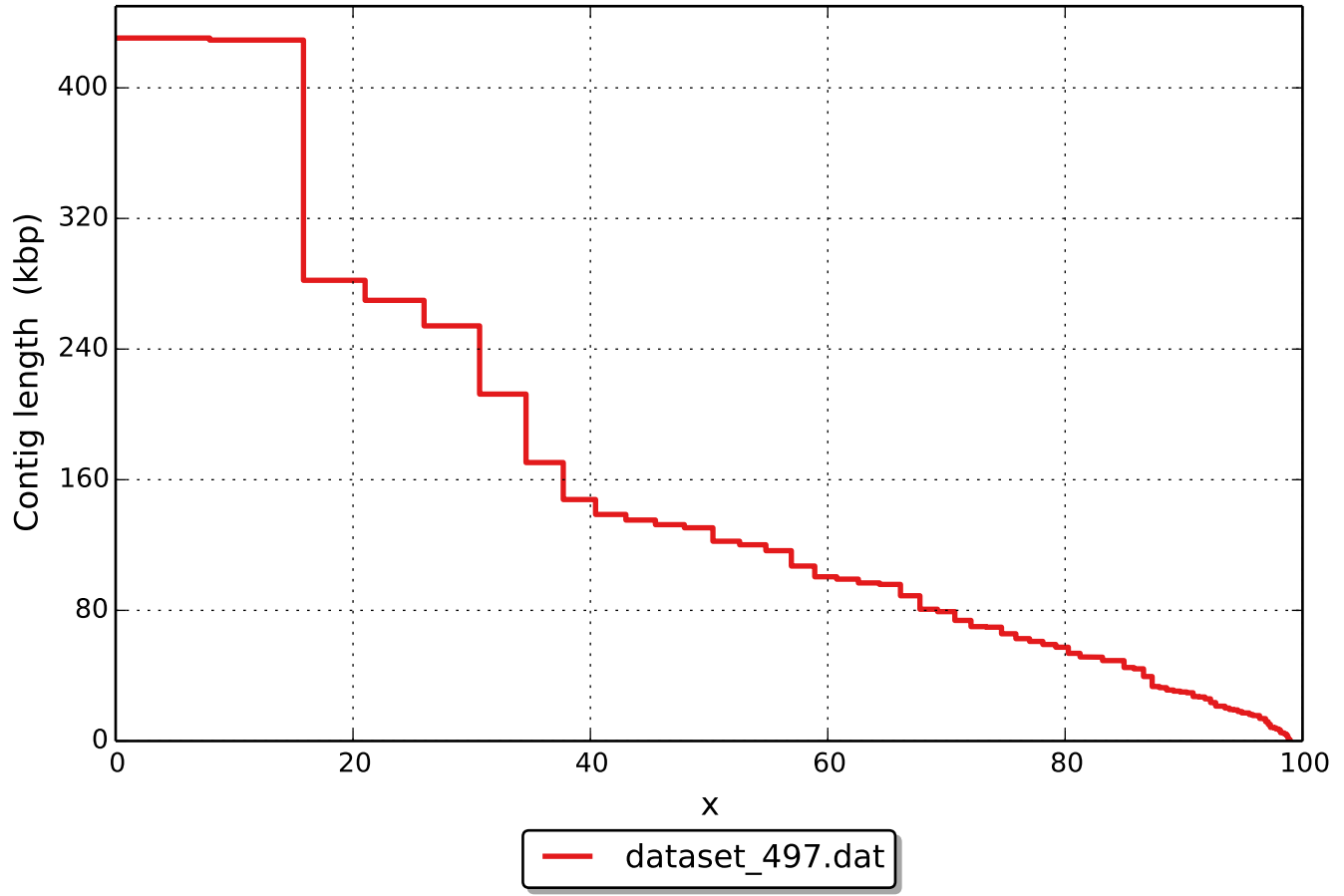


NAx

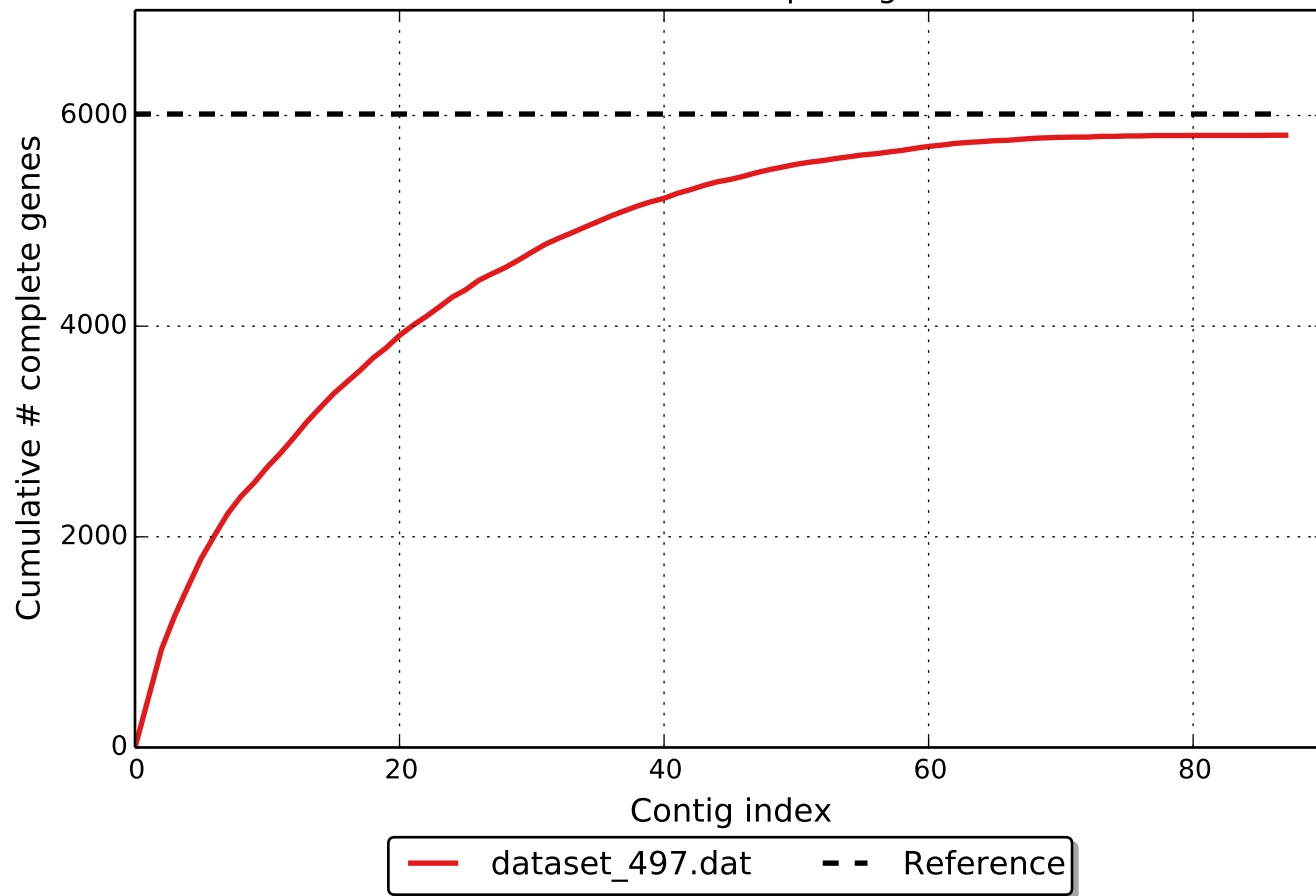


— dataset\_497.dat

NGAx



Cumulative # complete genes



# complete genes

7000

6500

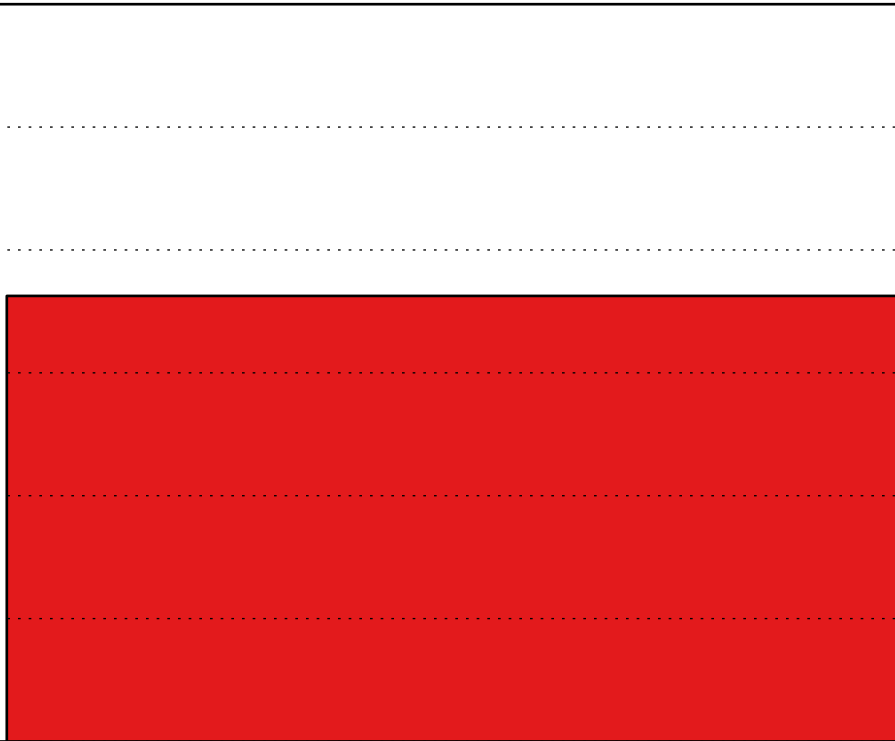
6000

5500

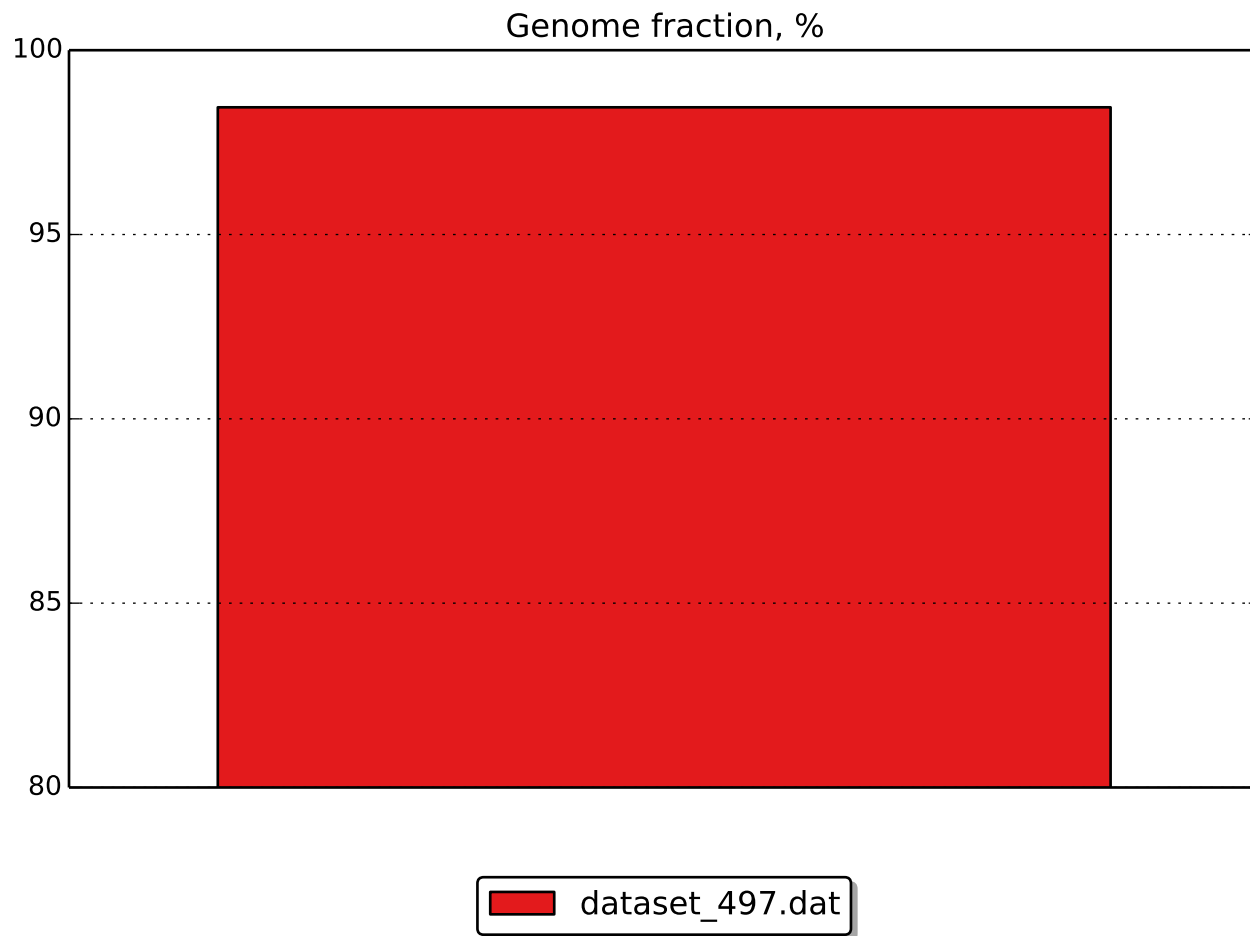
5000

4500

4000

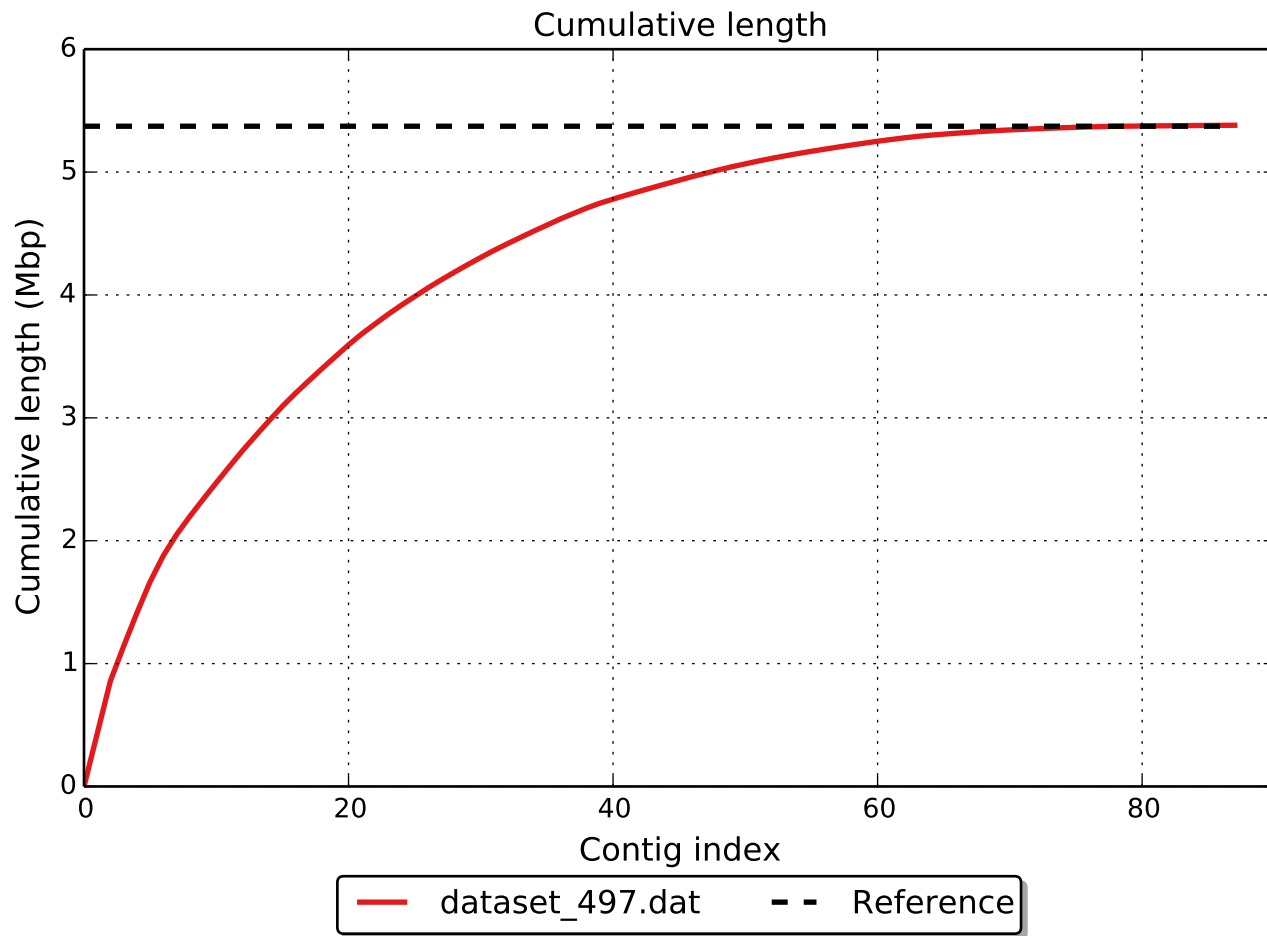


dataset\_497.dat

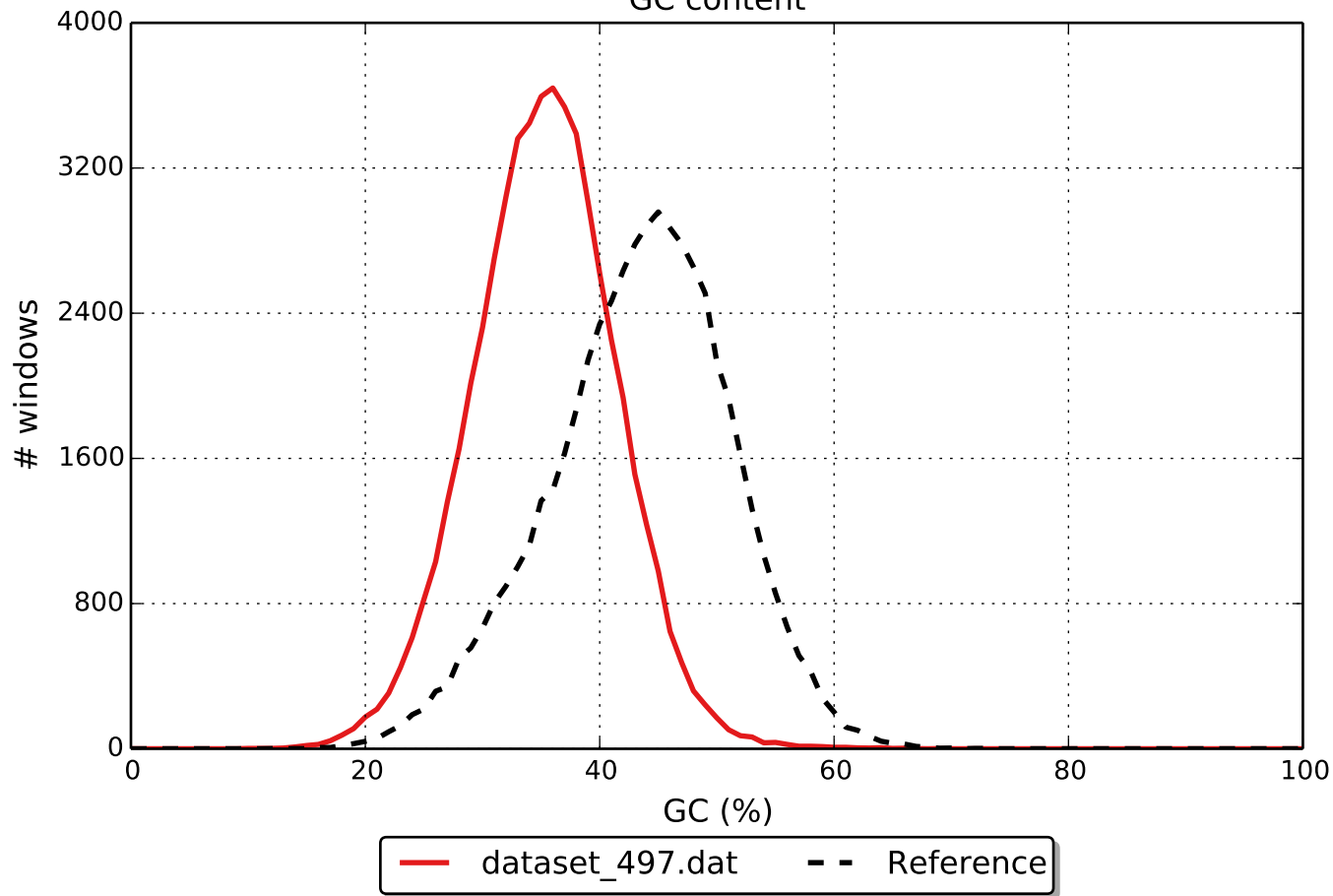




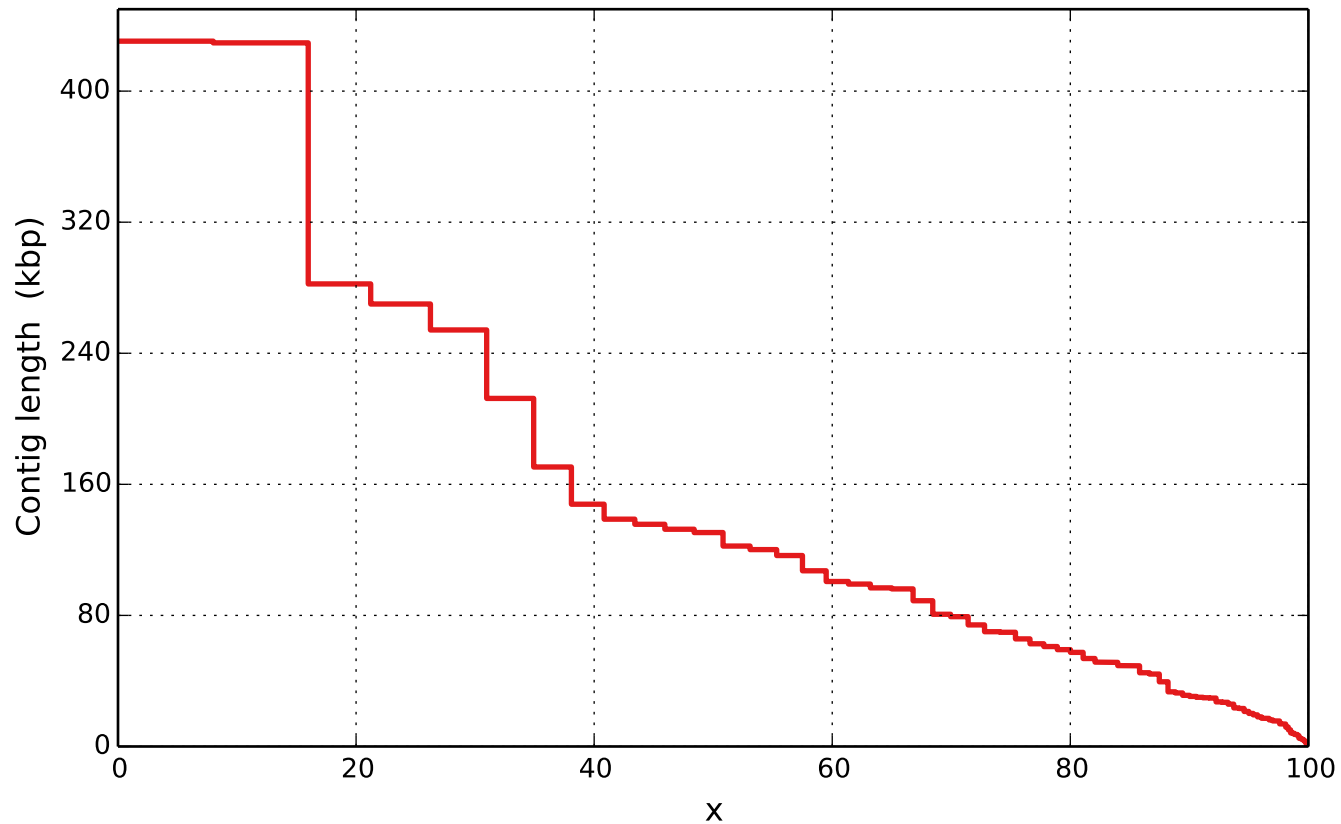
B\_fragilis\_638R



GC content

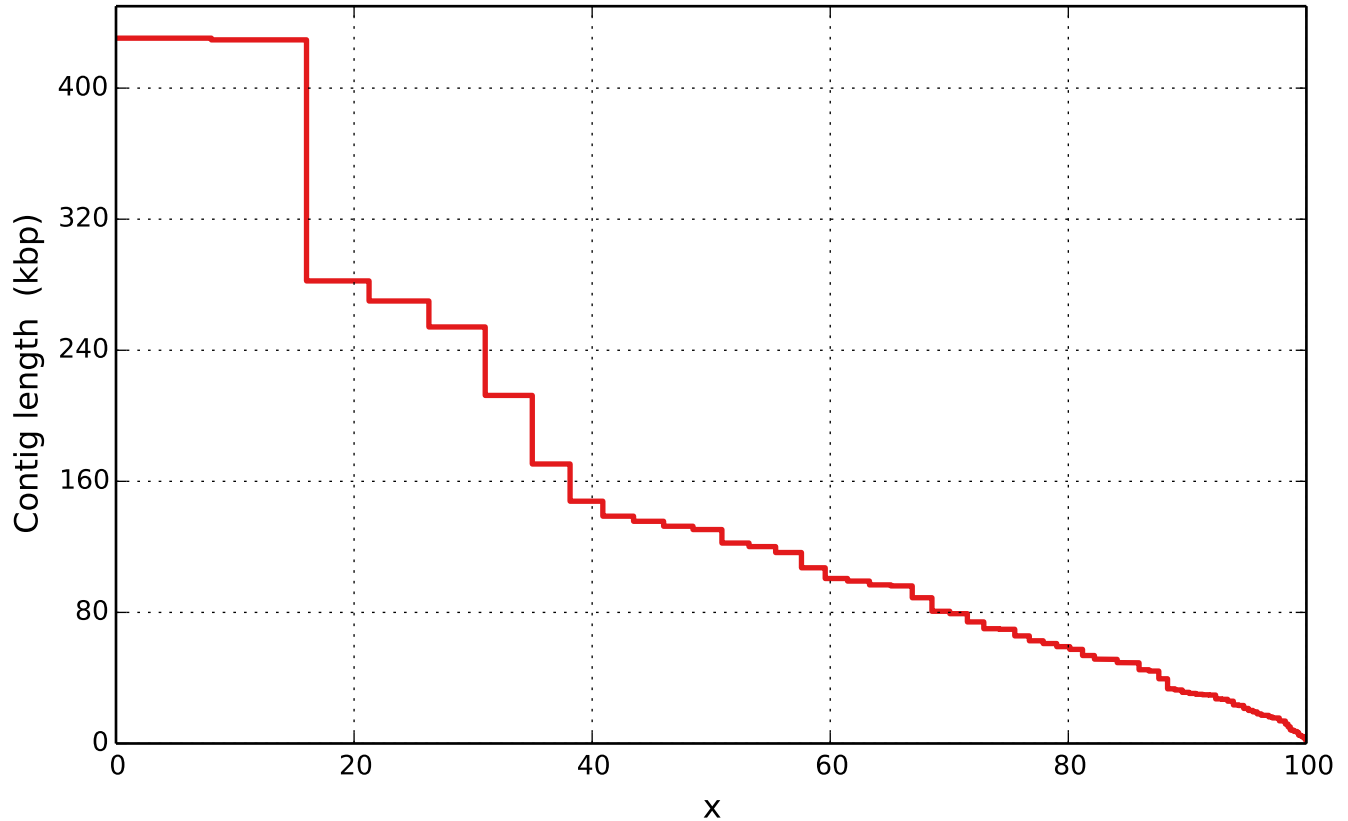


Nx



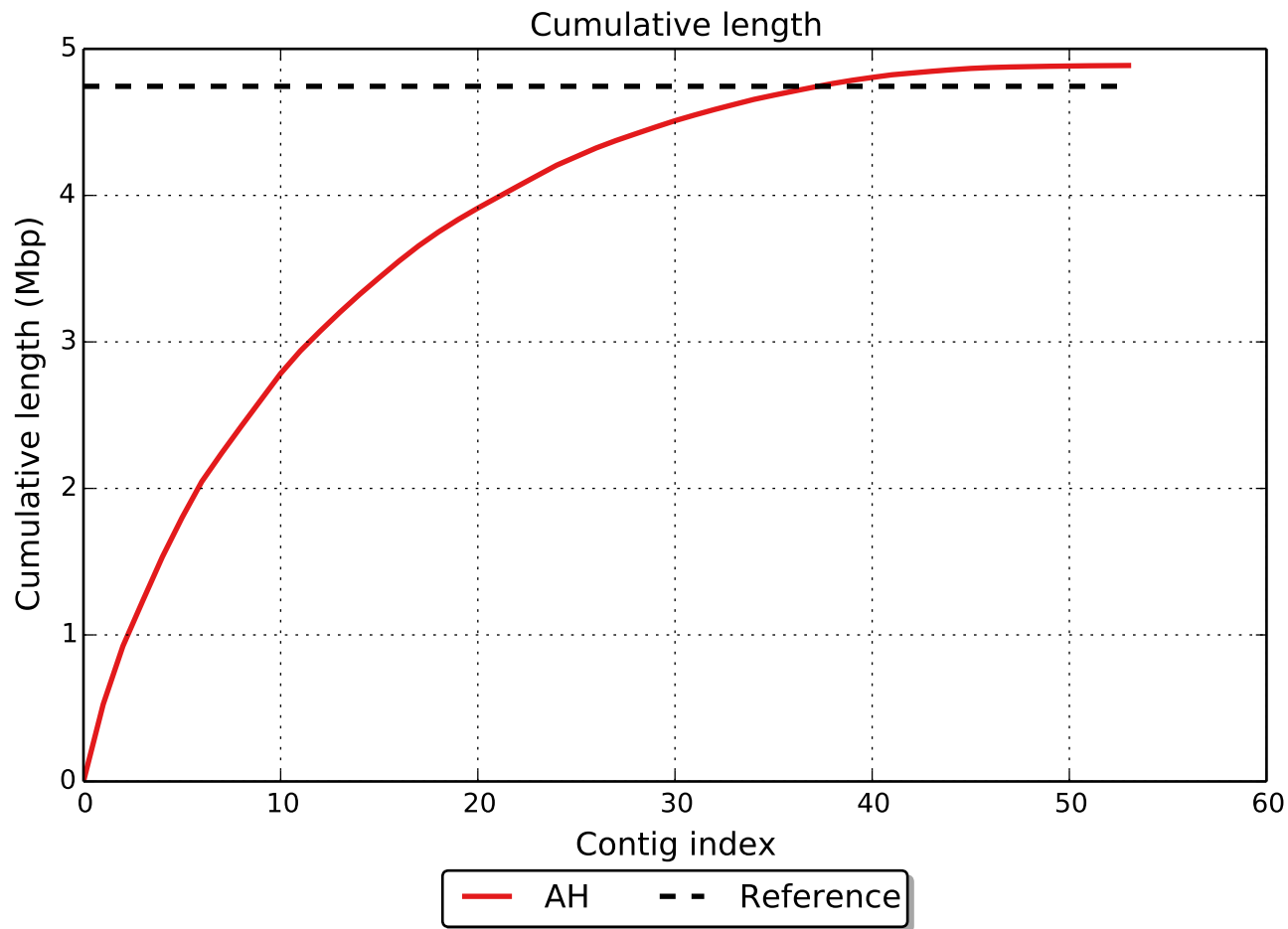
dataset\_497.dat

NGx

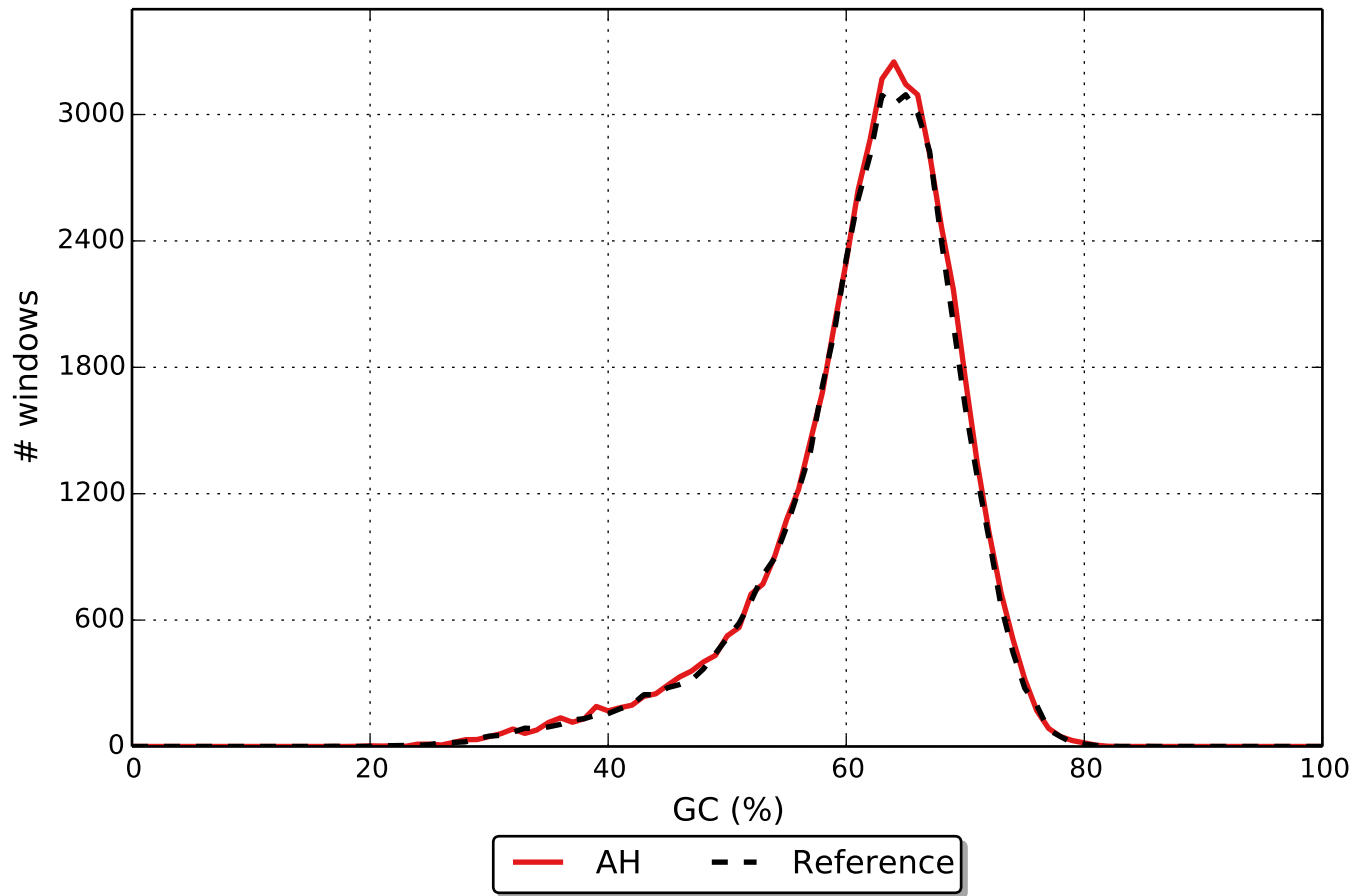


dataset\_497.dat

Contig

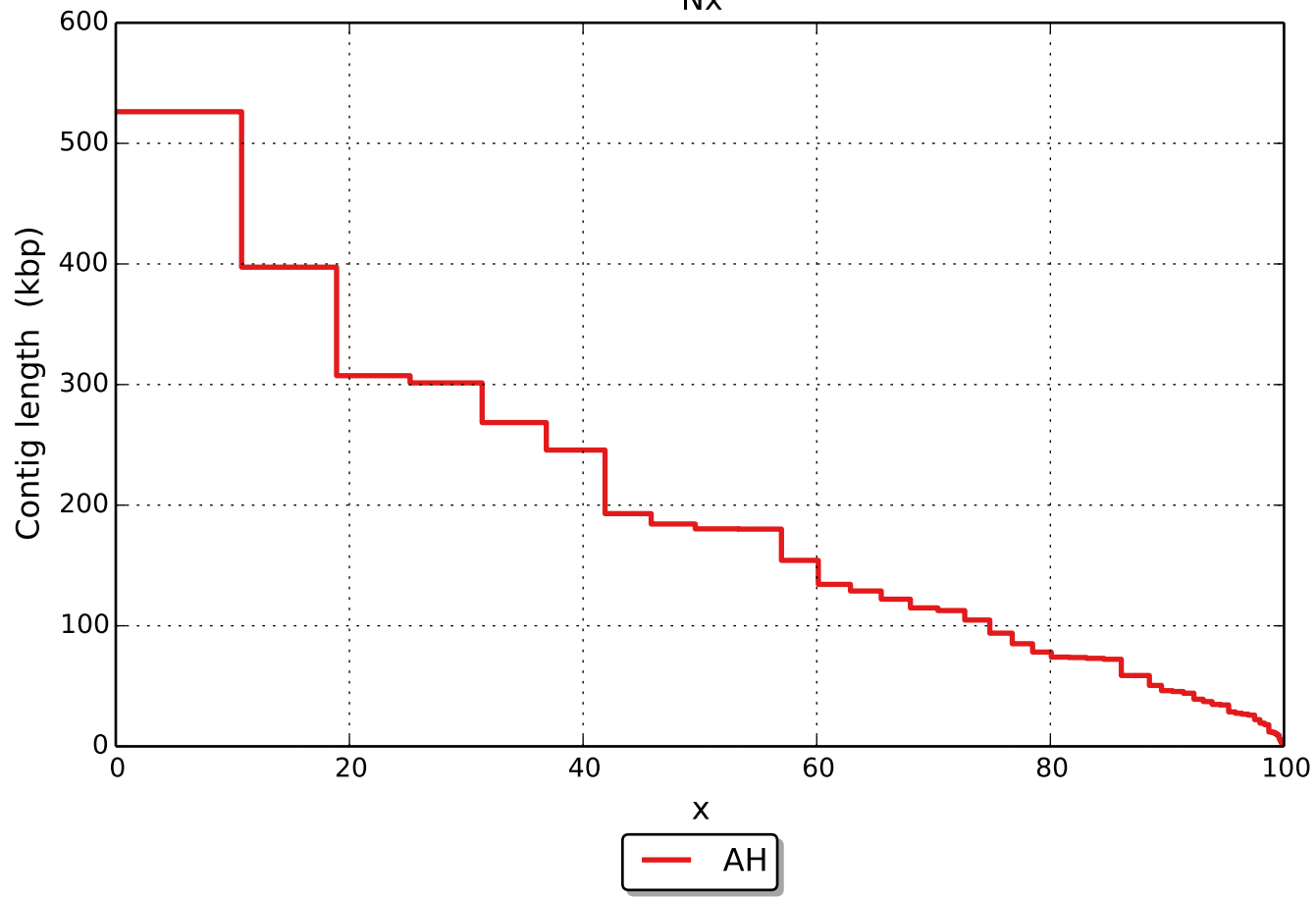


# GC content

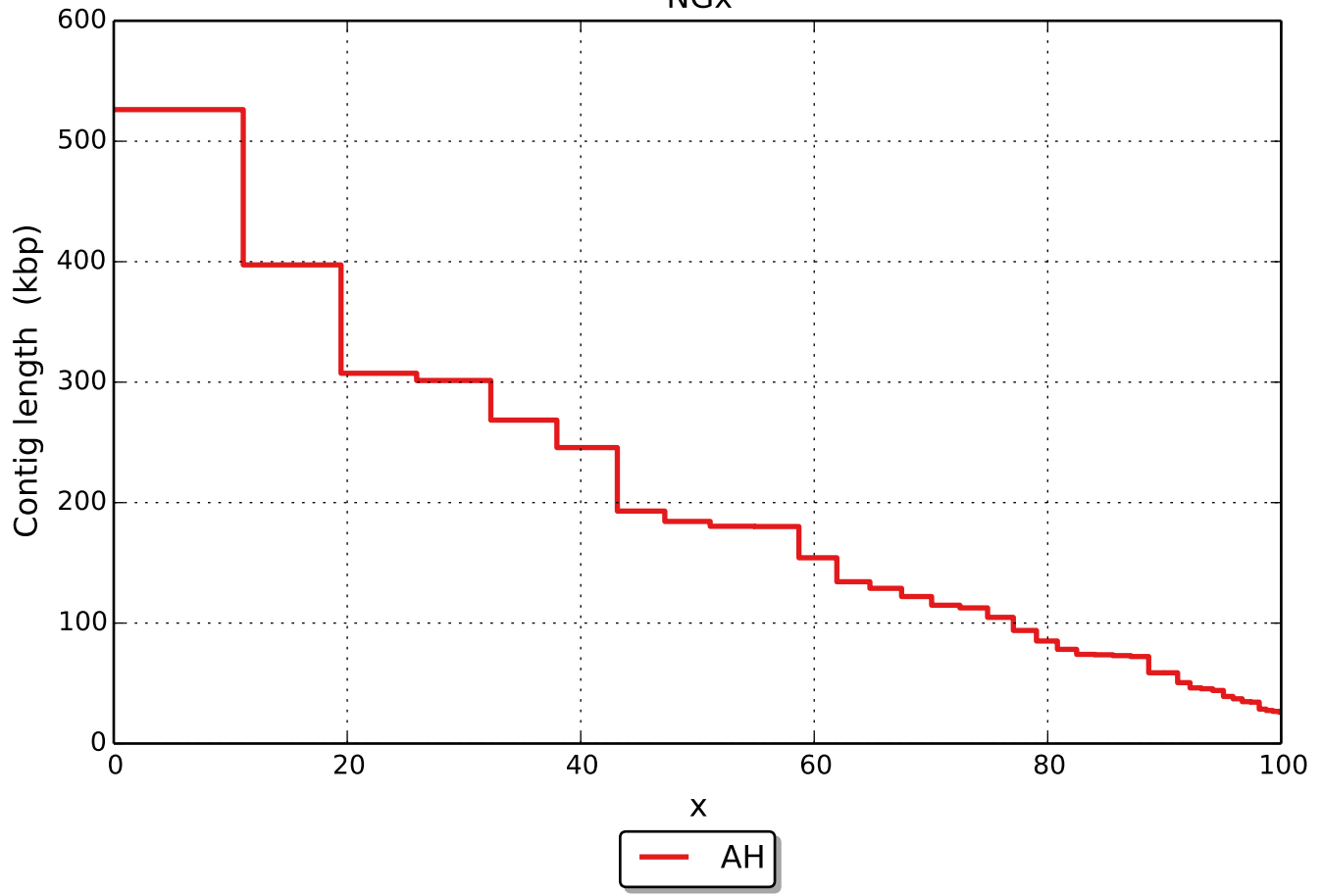


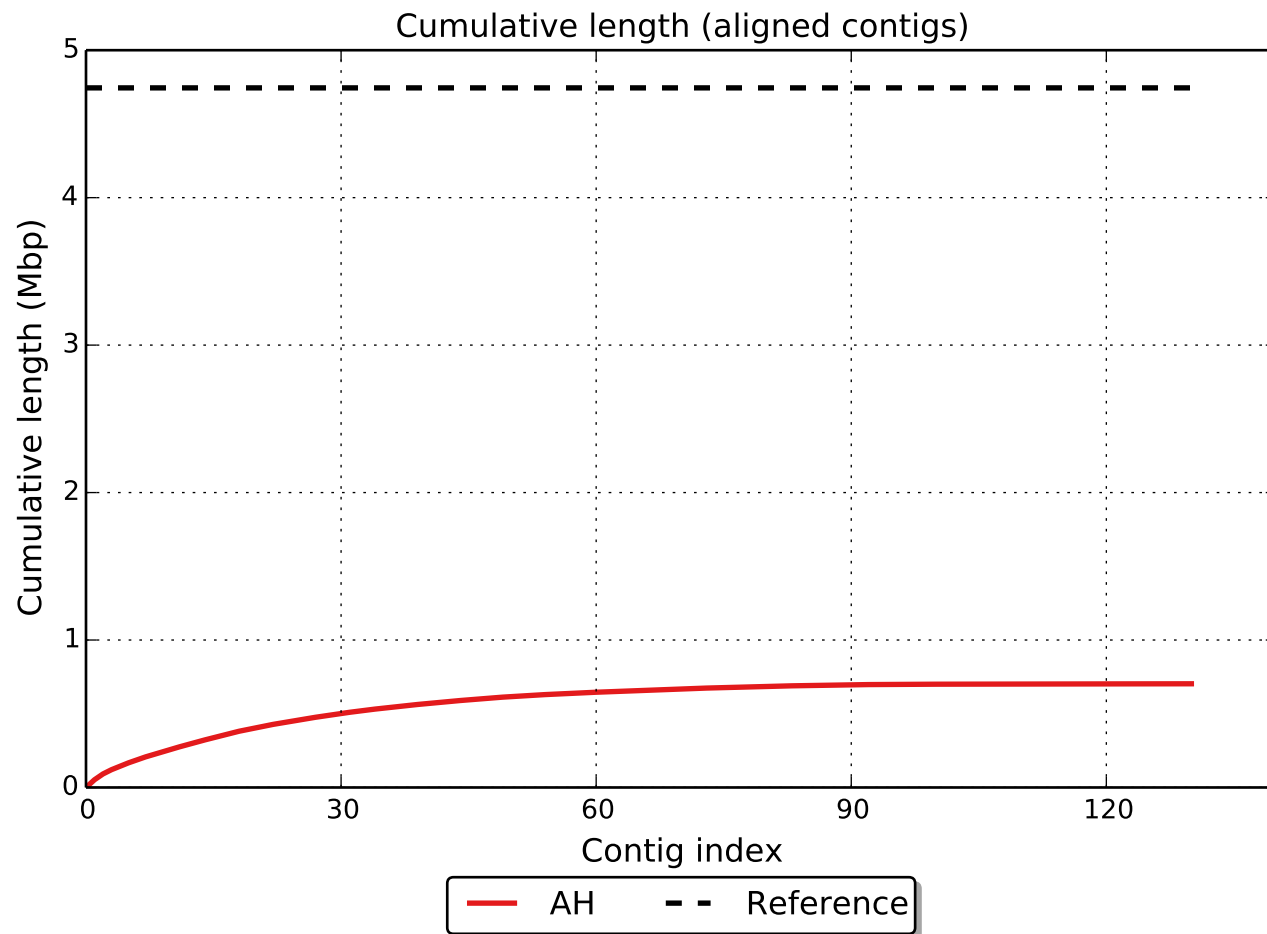


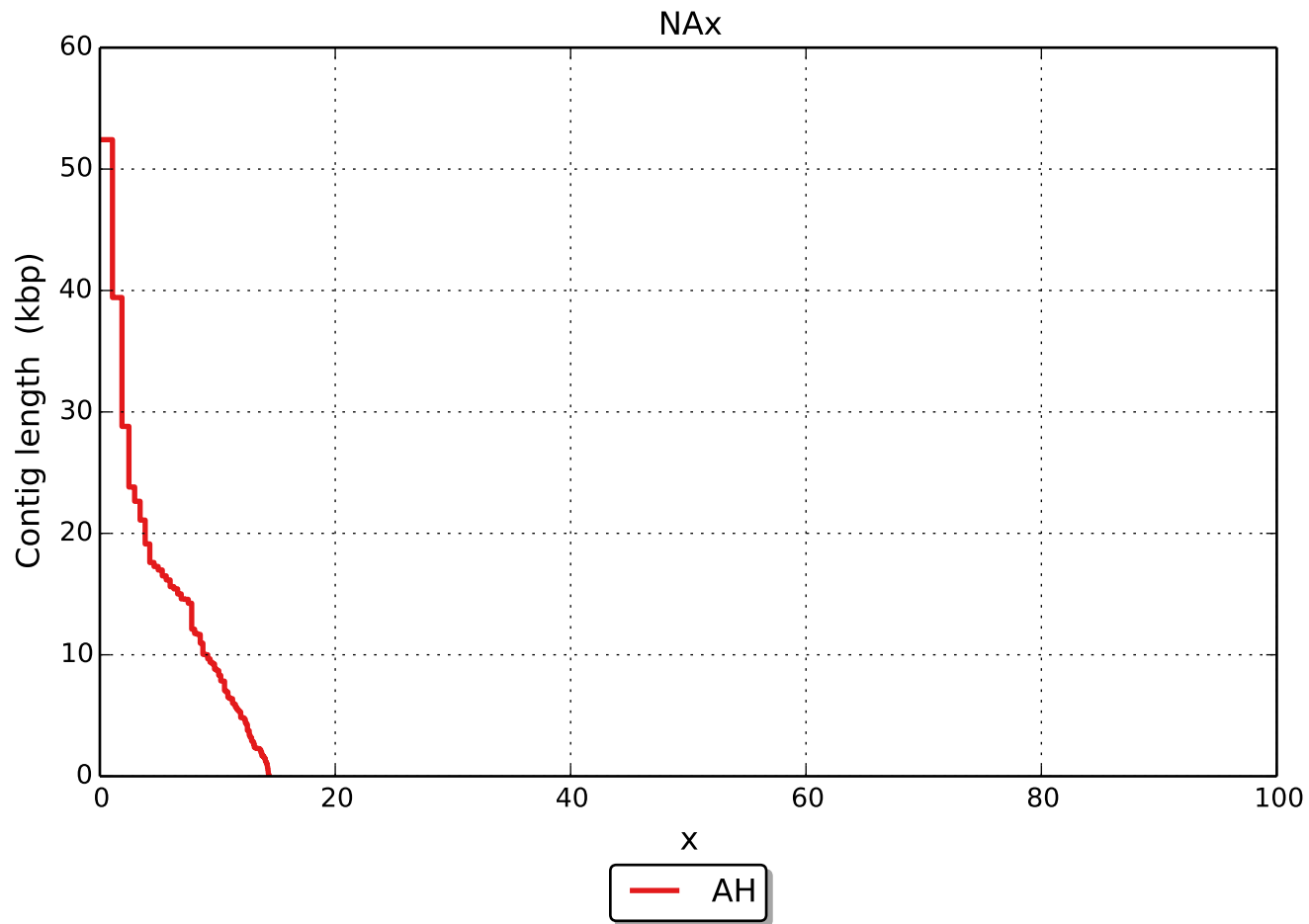
Nx



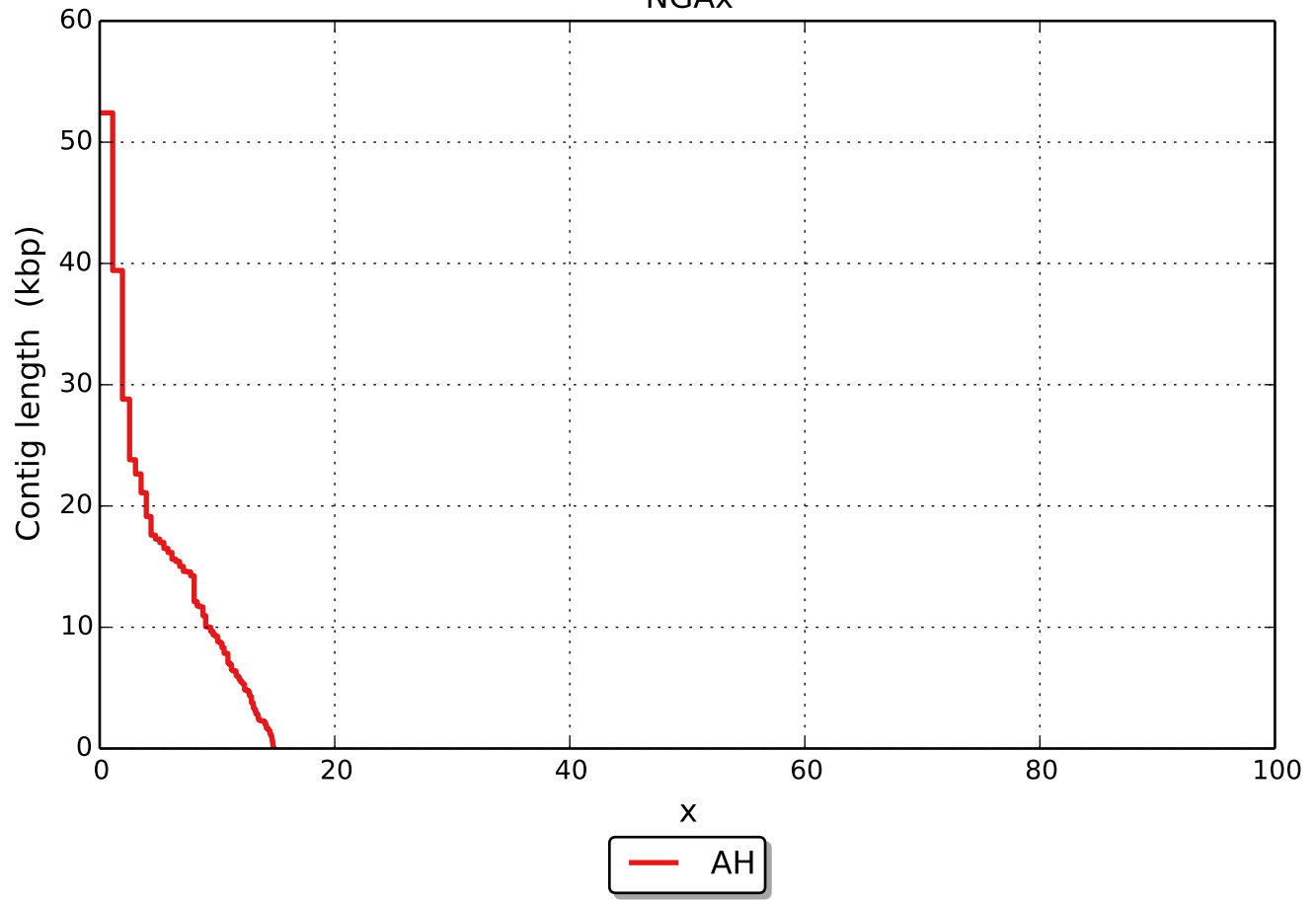
NGx



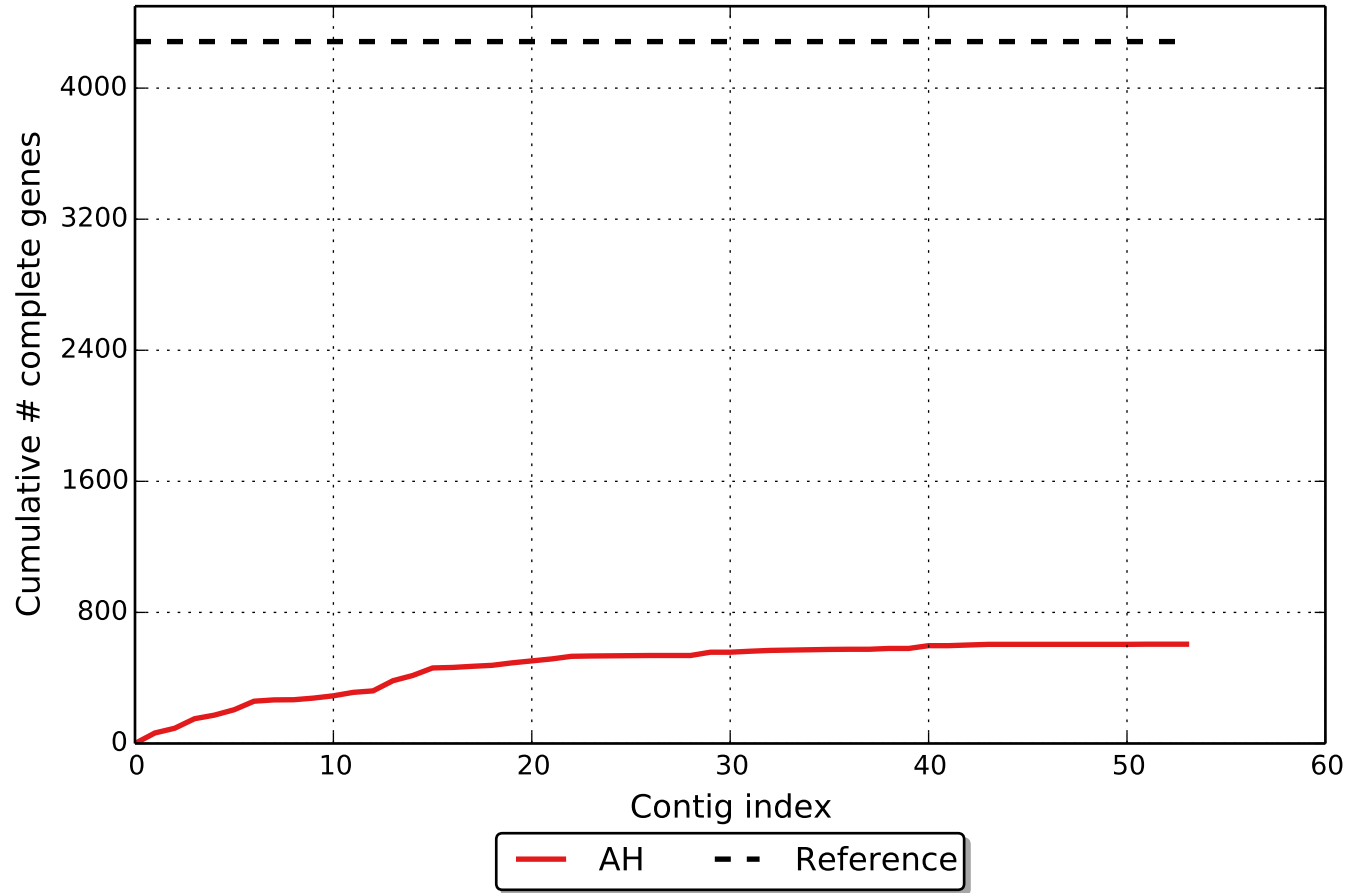




NGAx



Cumulative # complete genes



# complete genes

