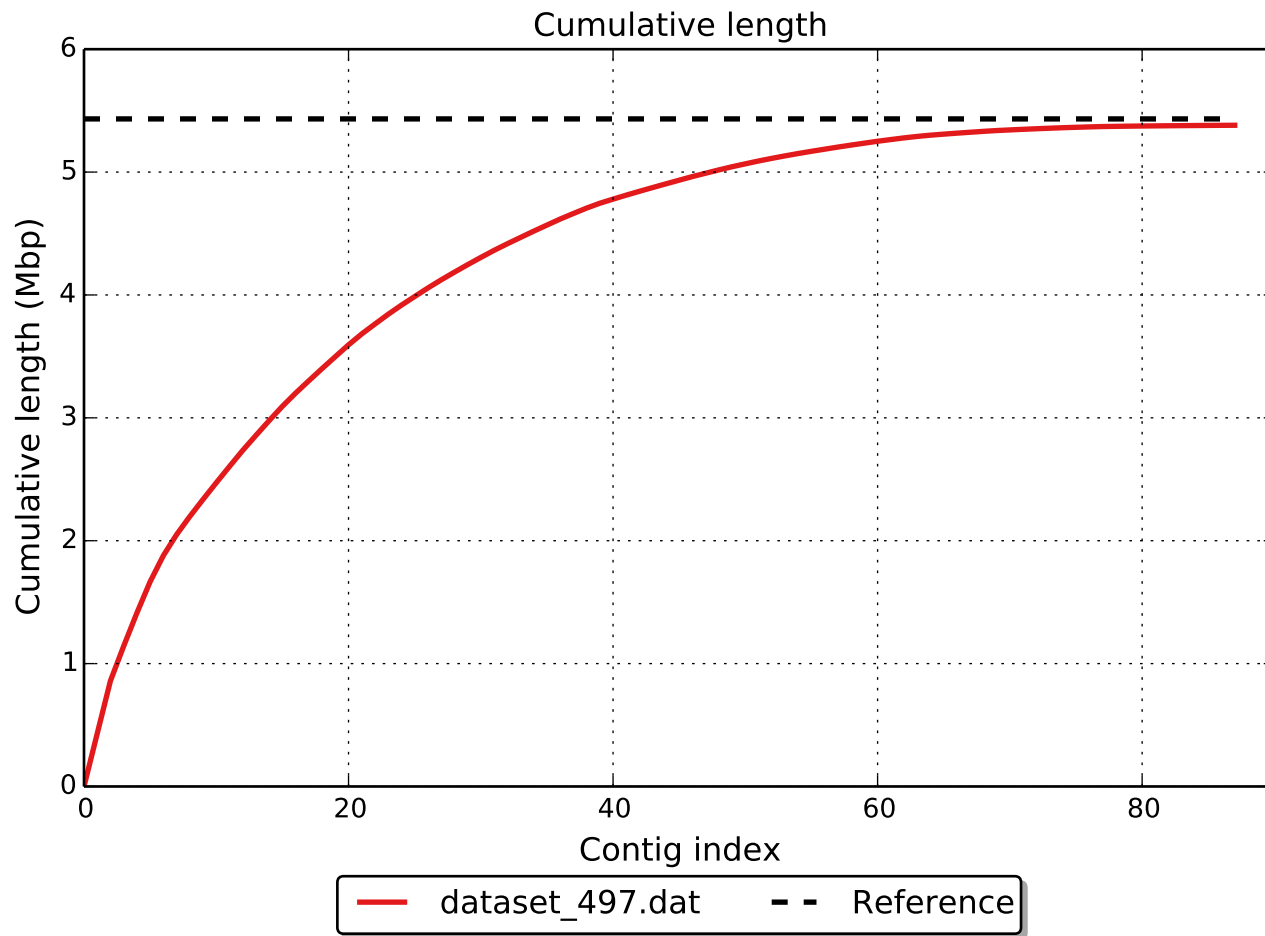
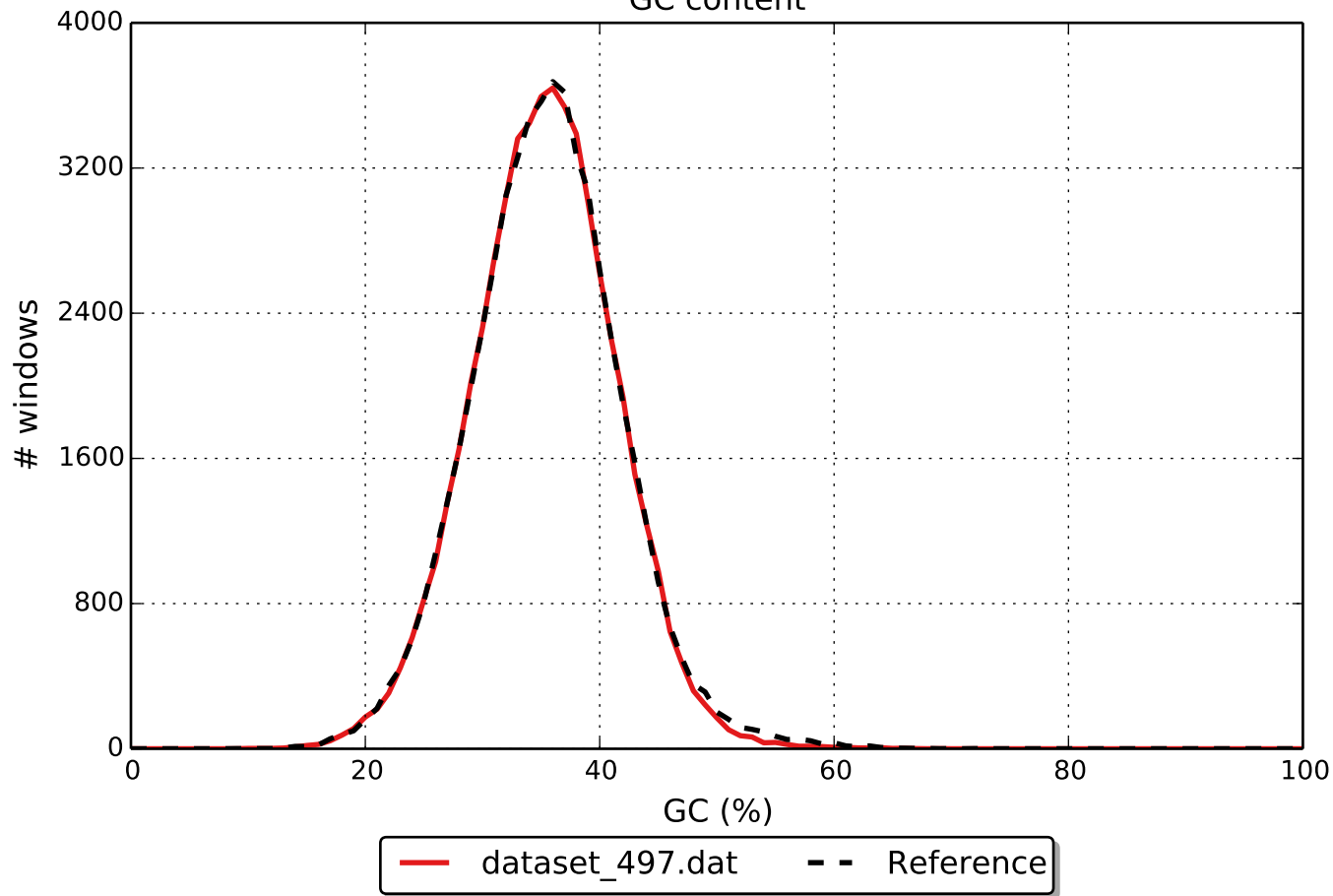


Combined\_reference

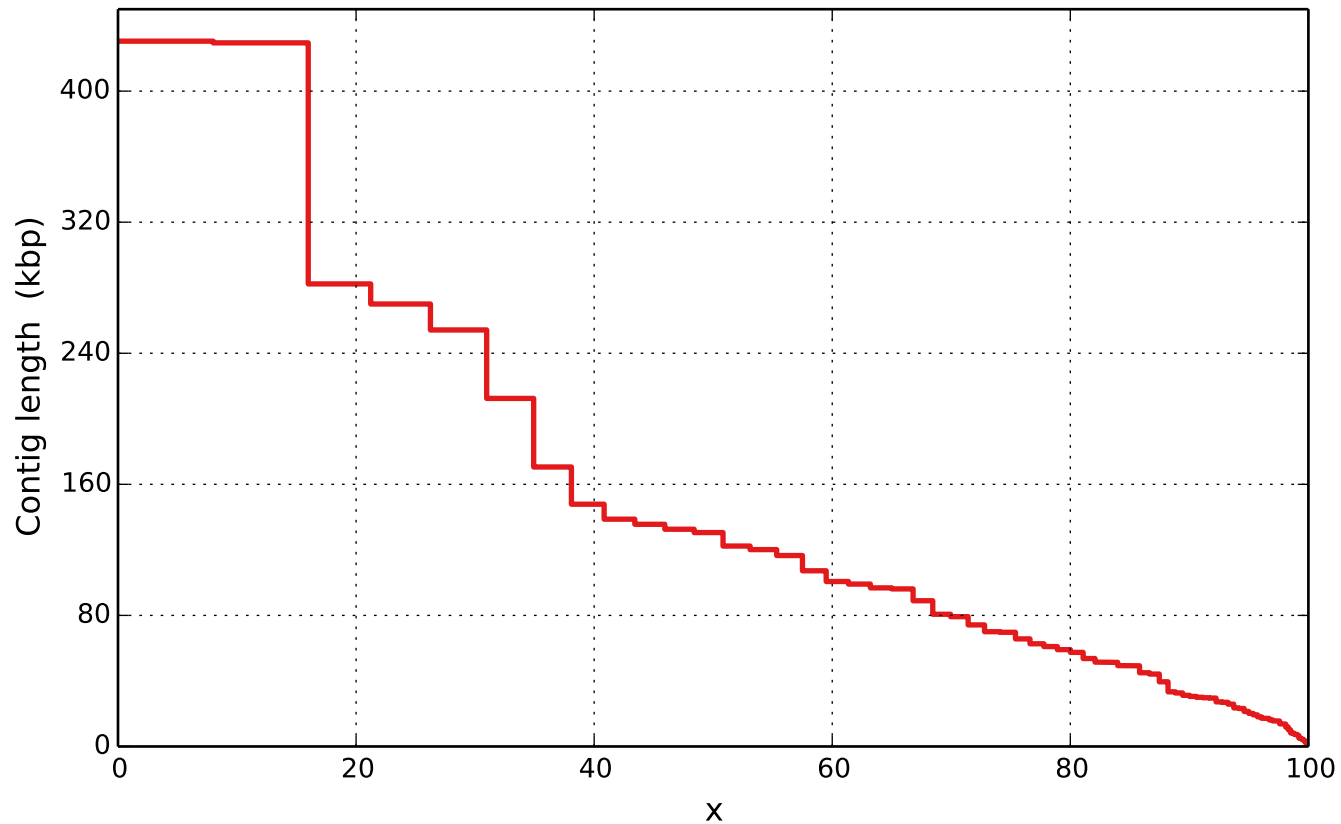
B\_cereus\_ATCC\_1098



GC content

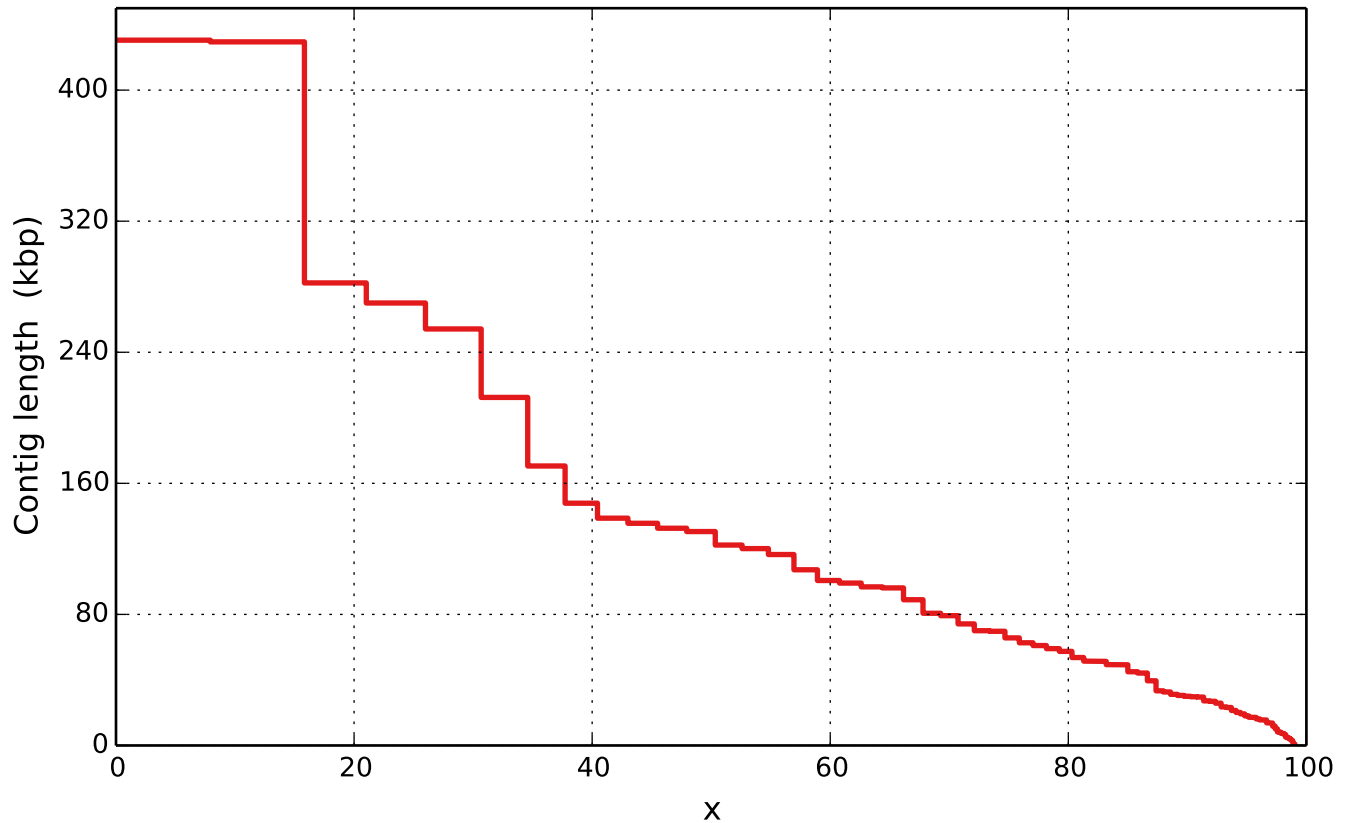


Nx

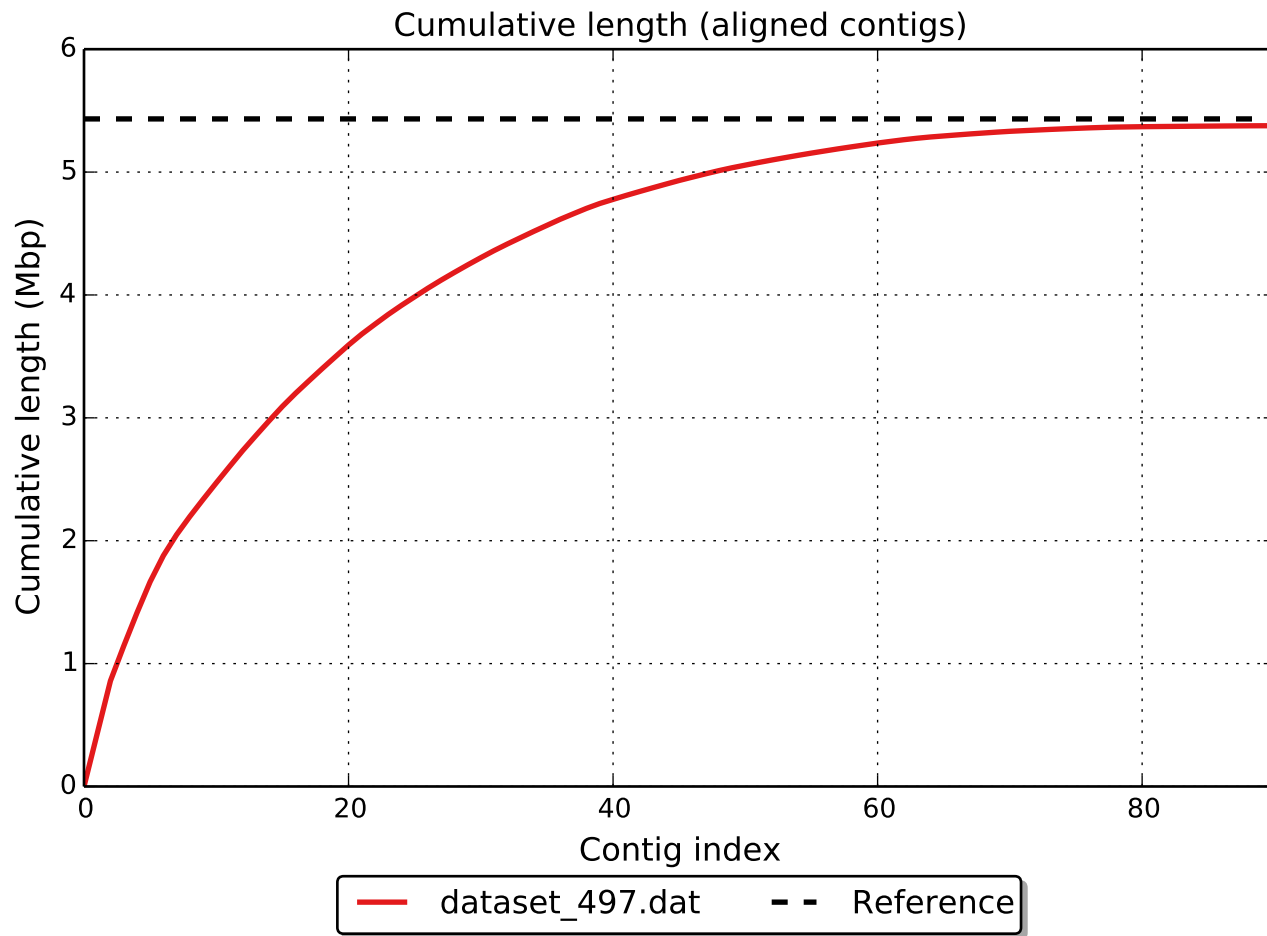


dataset\_497.dat

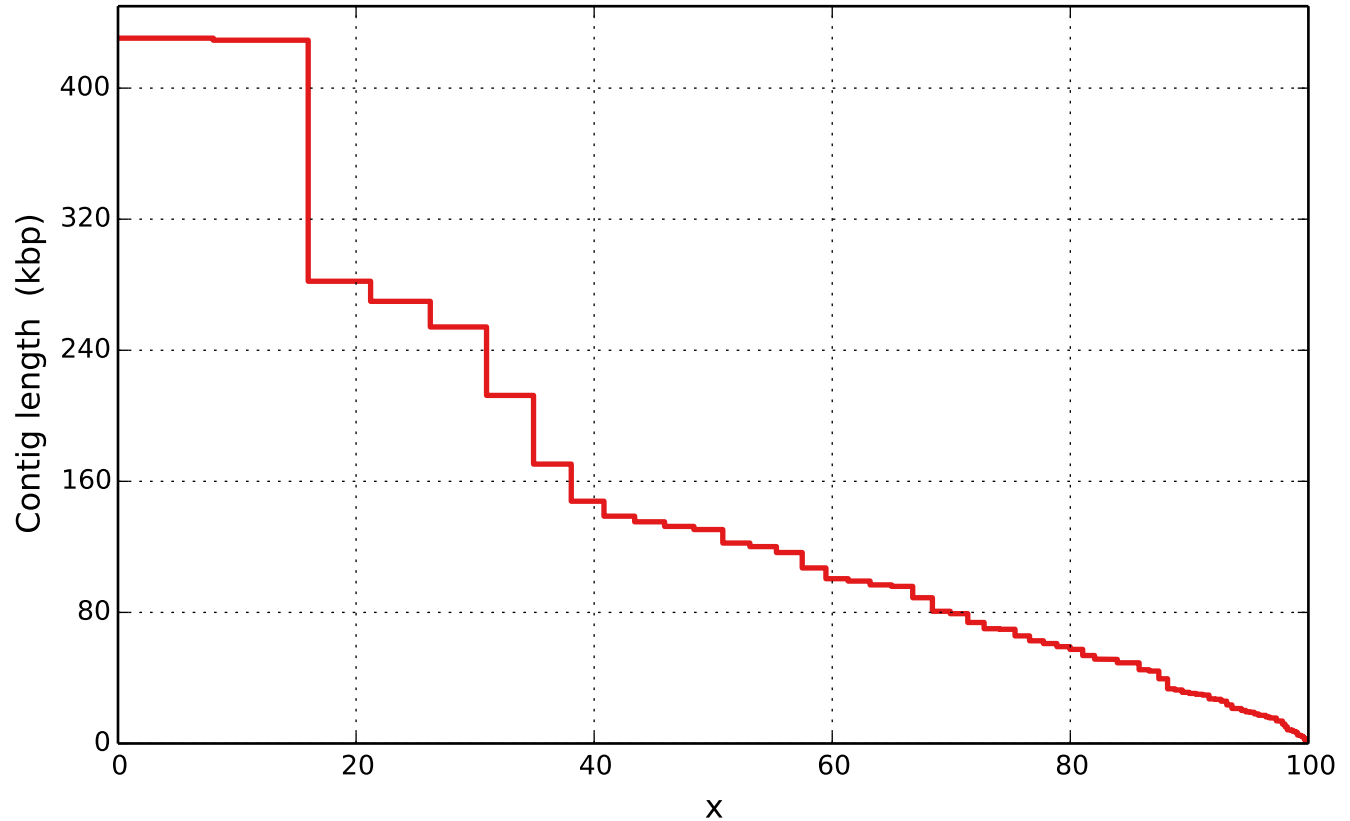
NGx



— dataset\_497.dat



NAx



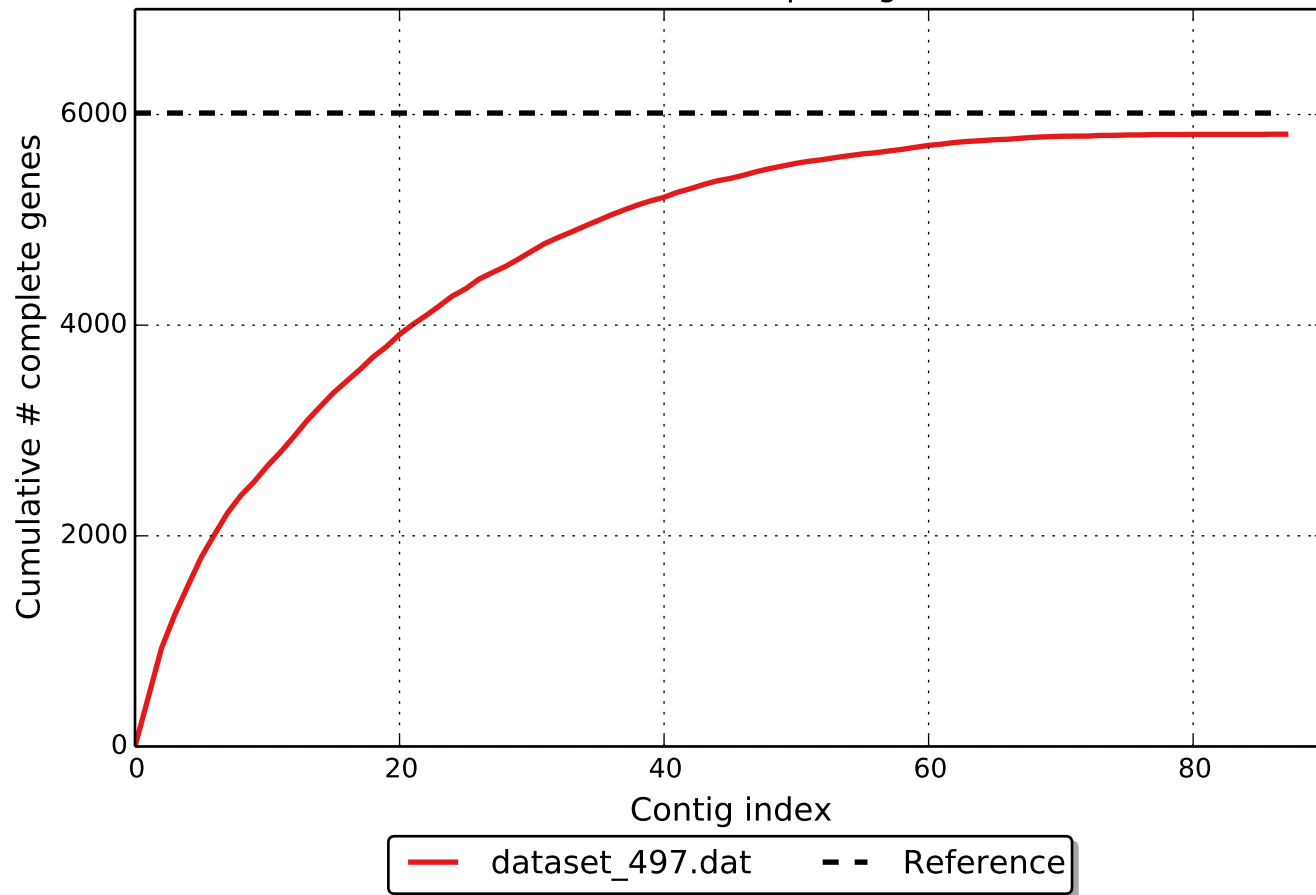
dataset\_497.dat



NGAx



Cumulative # complete genes



# complete genes

7000

6500

6000

5500

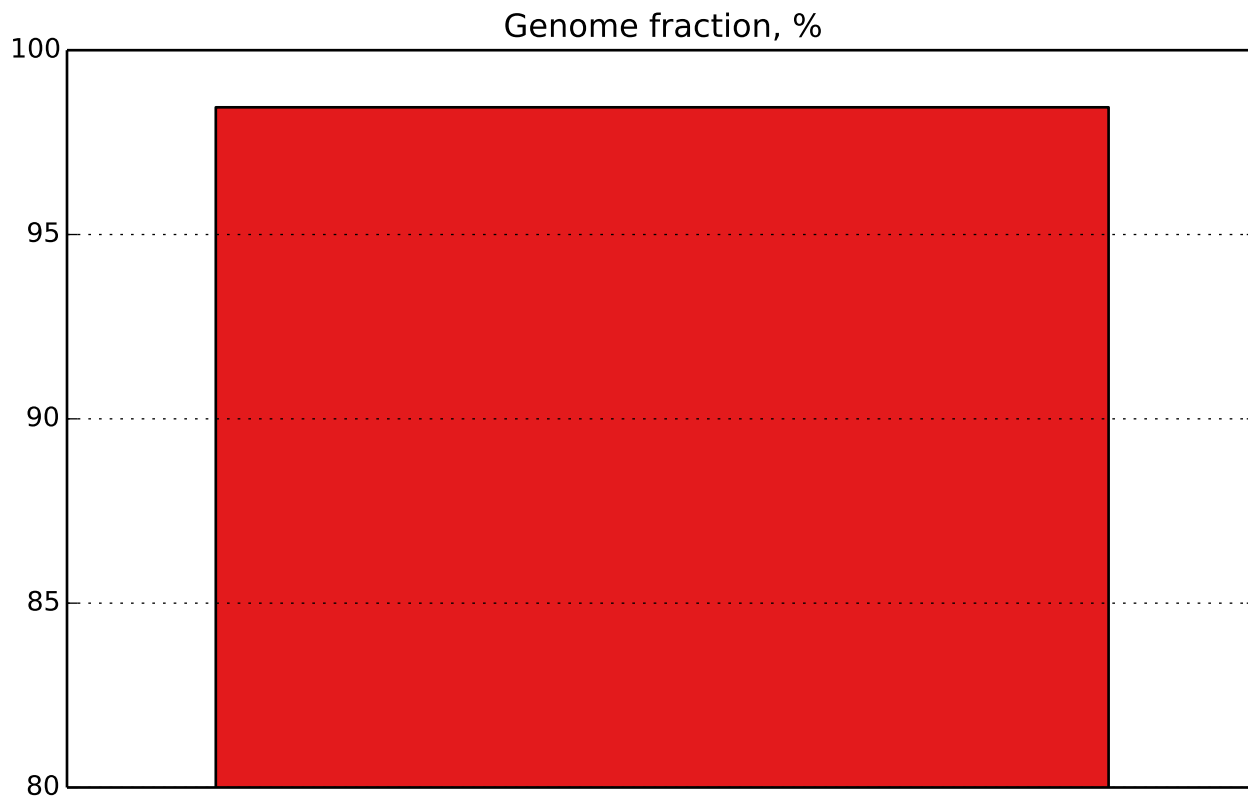
5000

4500

4000

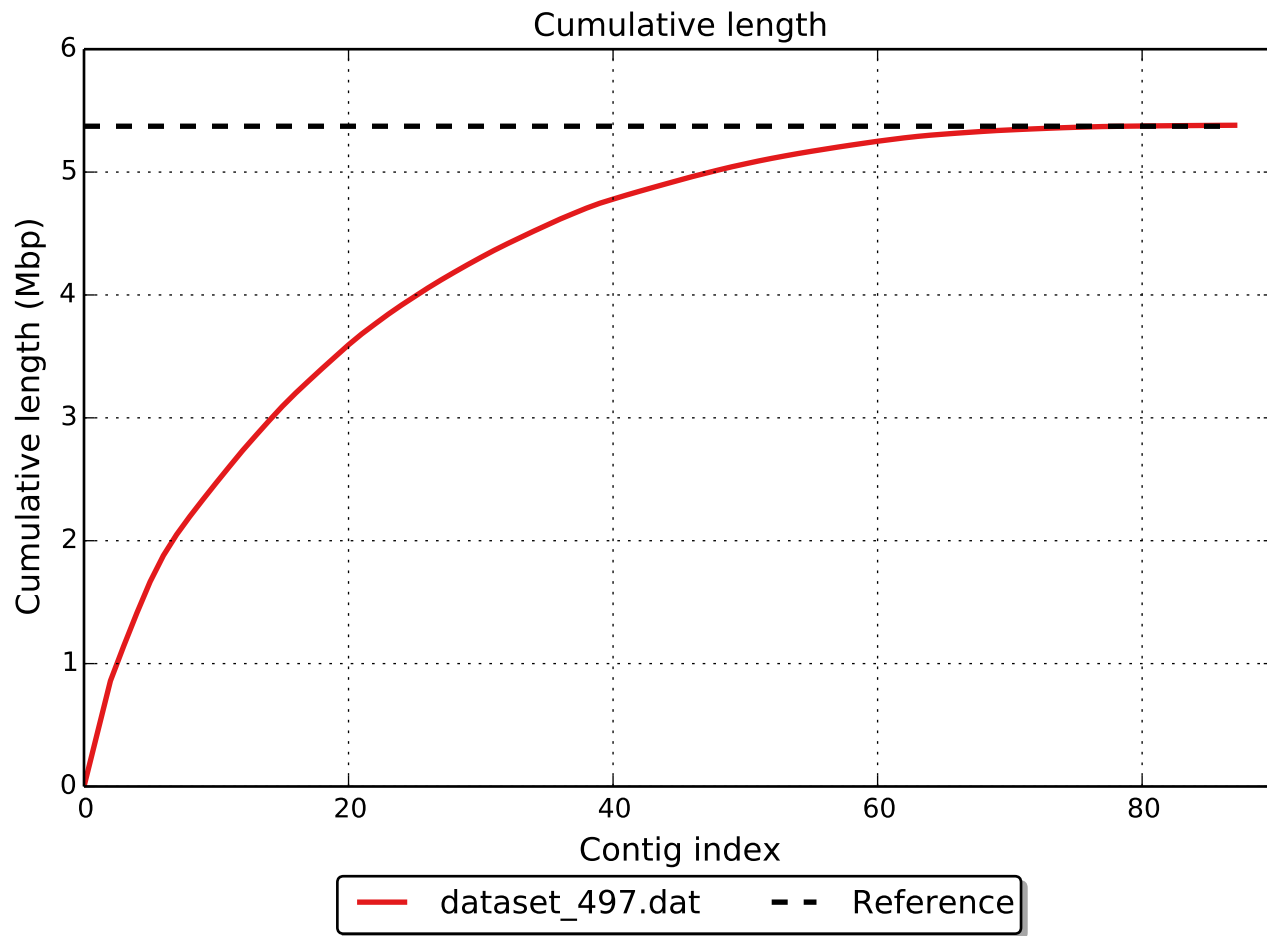


dataset\_497.dat

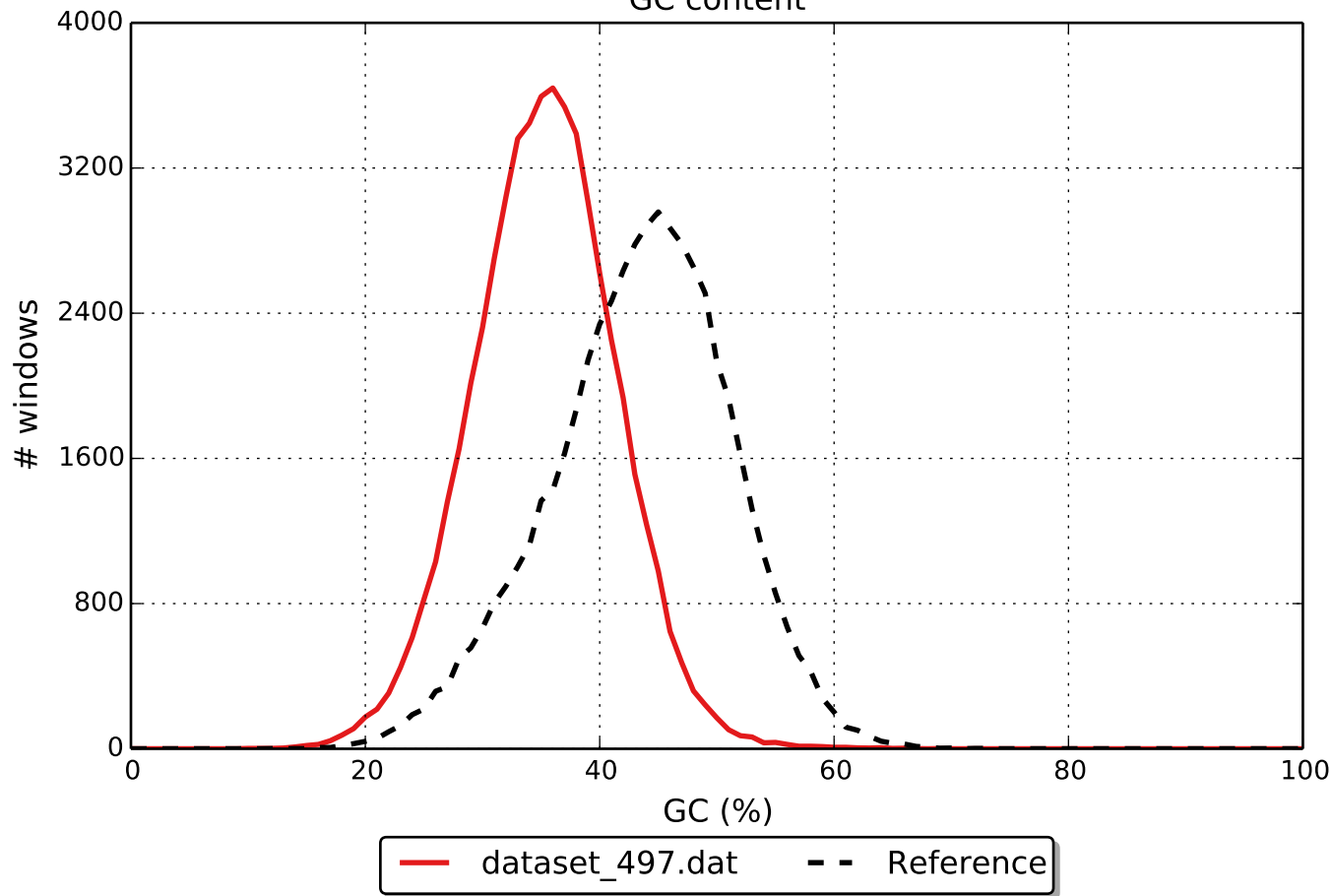


dataset\_497.dat

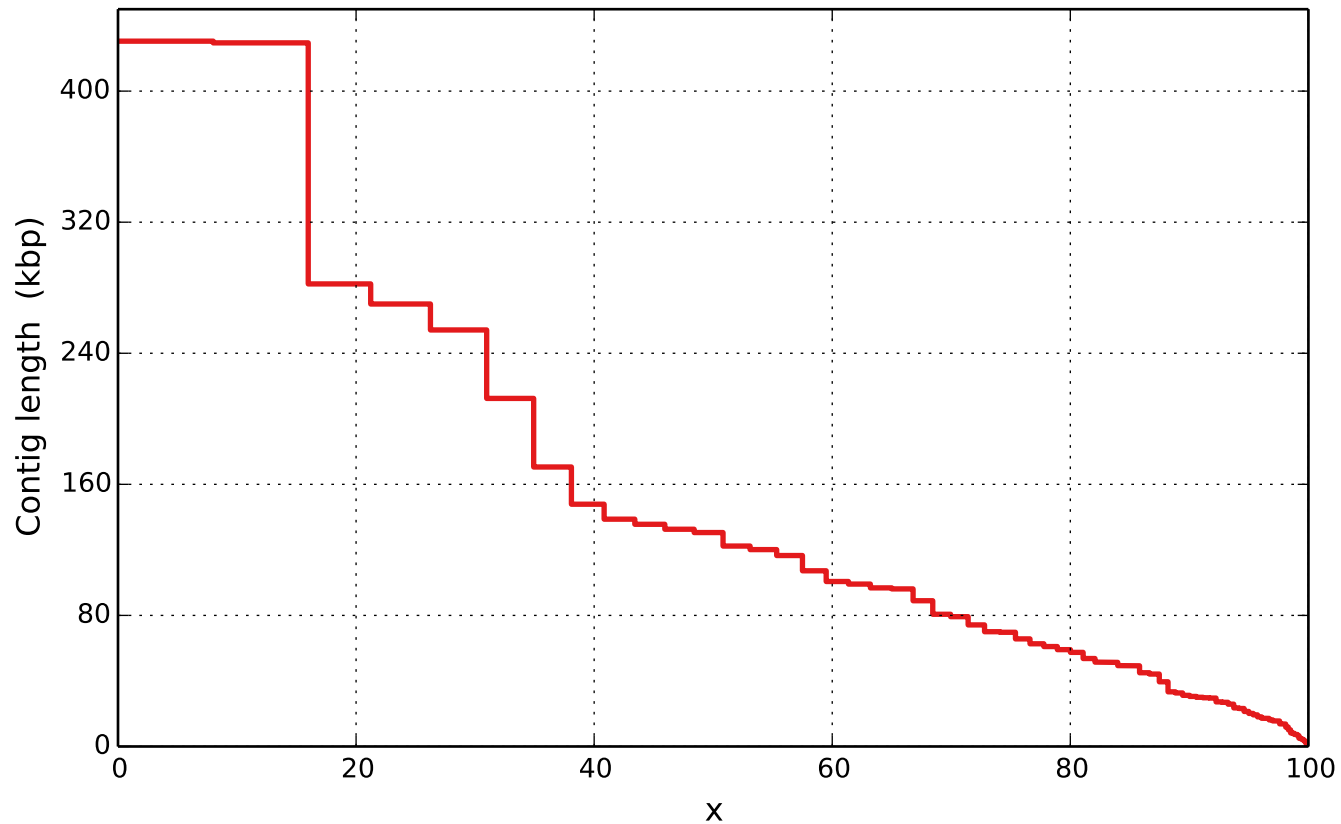
B\_fragilis\_638R



GC content



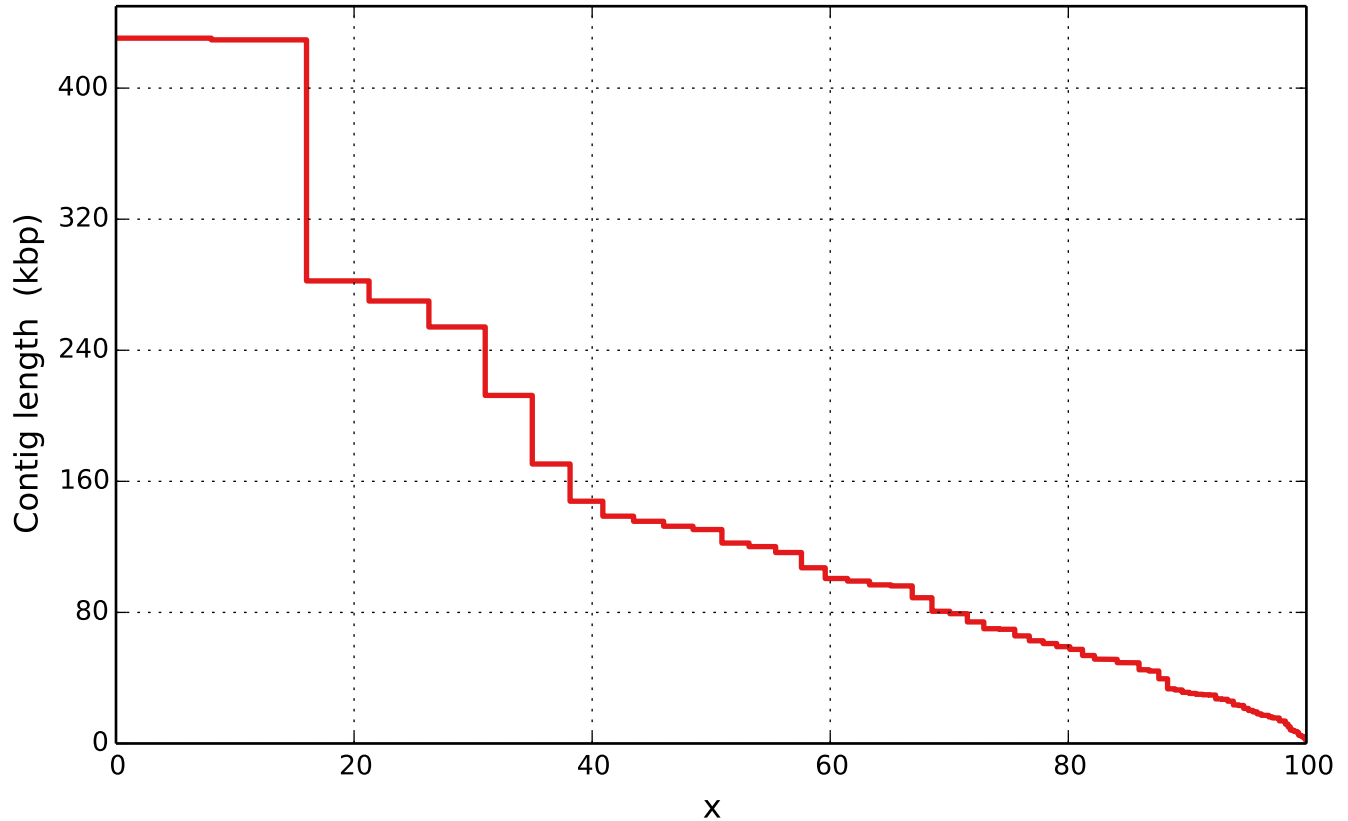
Nx



dataset\_497.dat

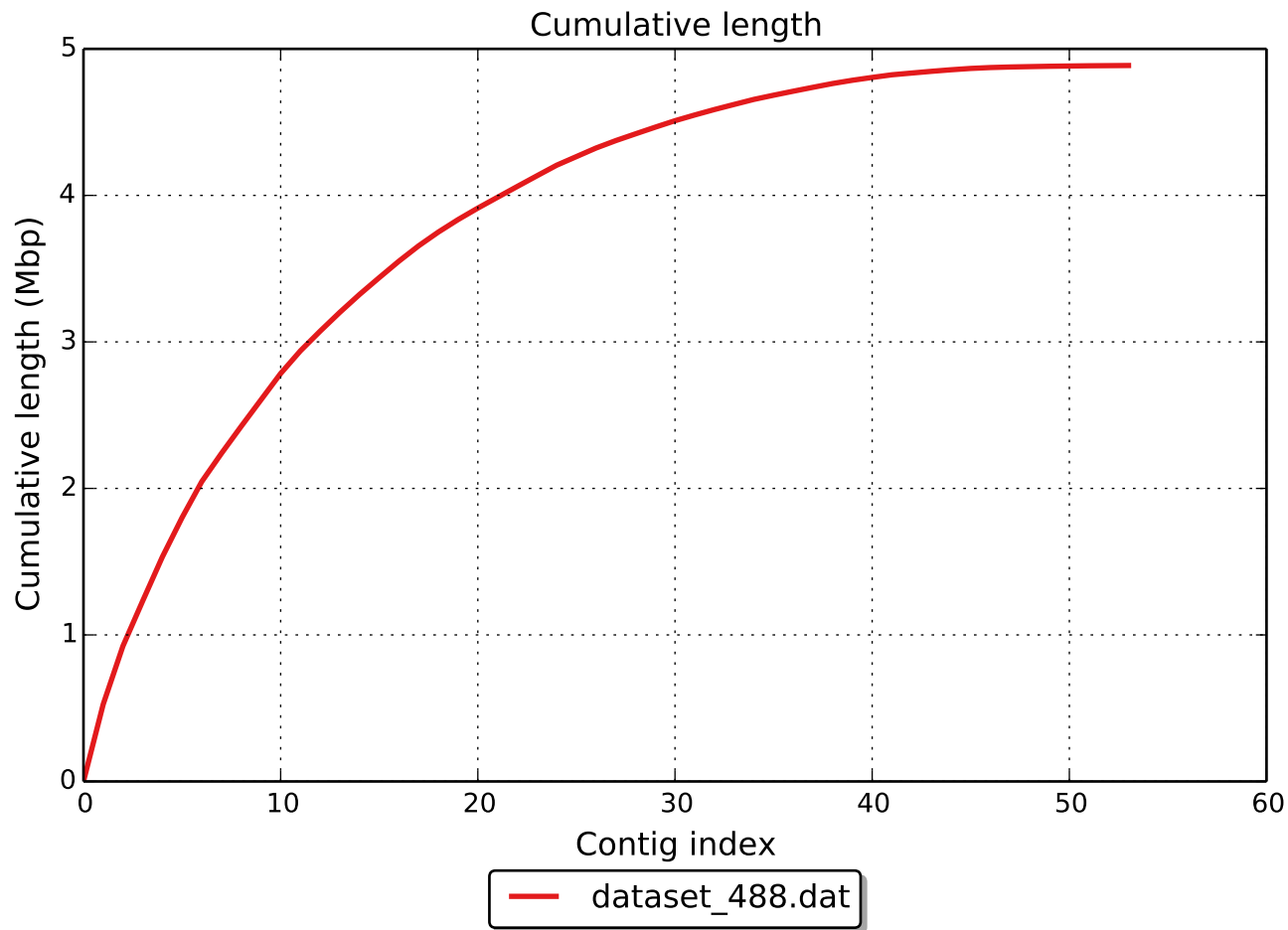


NGx

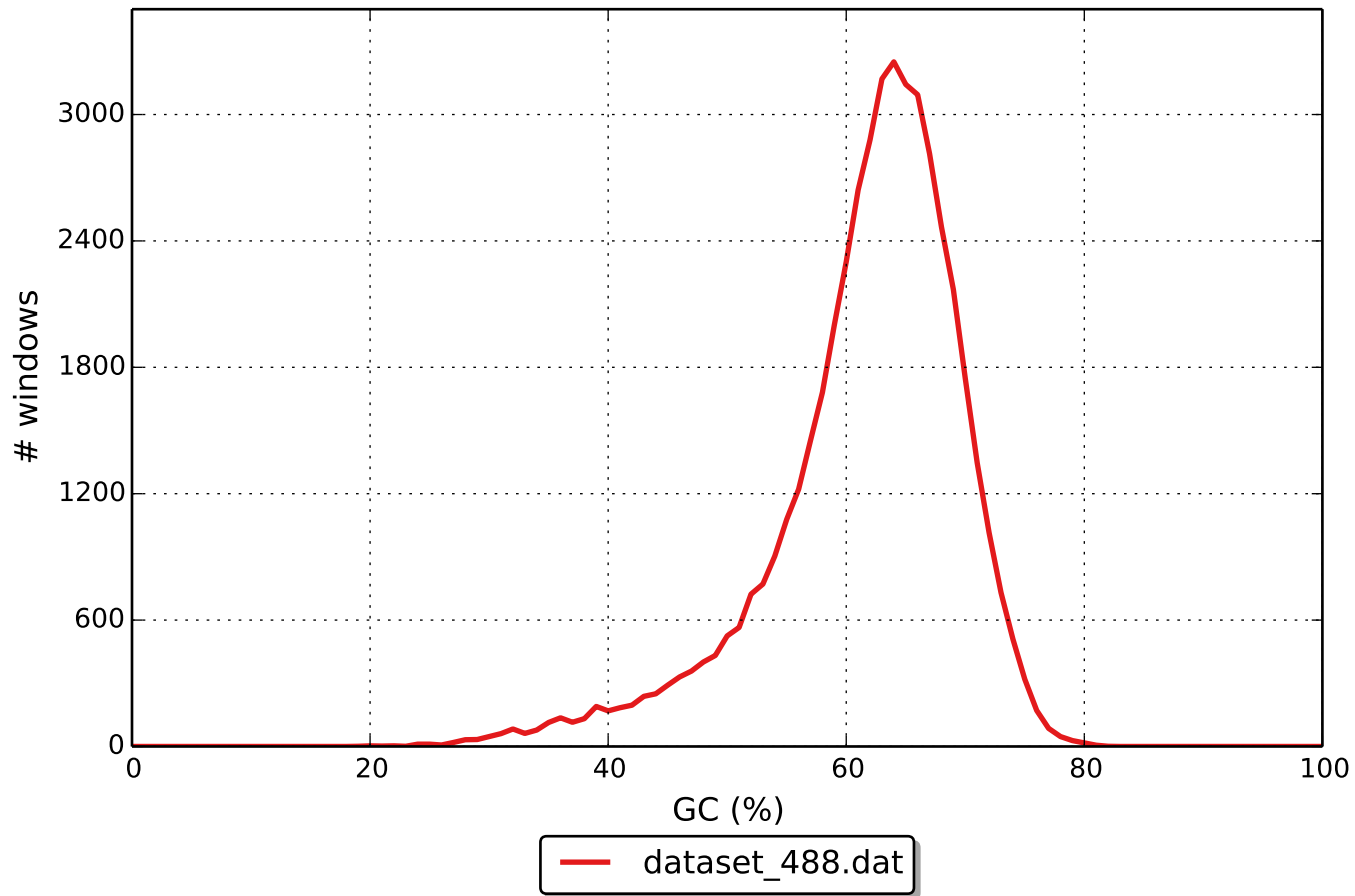


— dataset\_497.dat

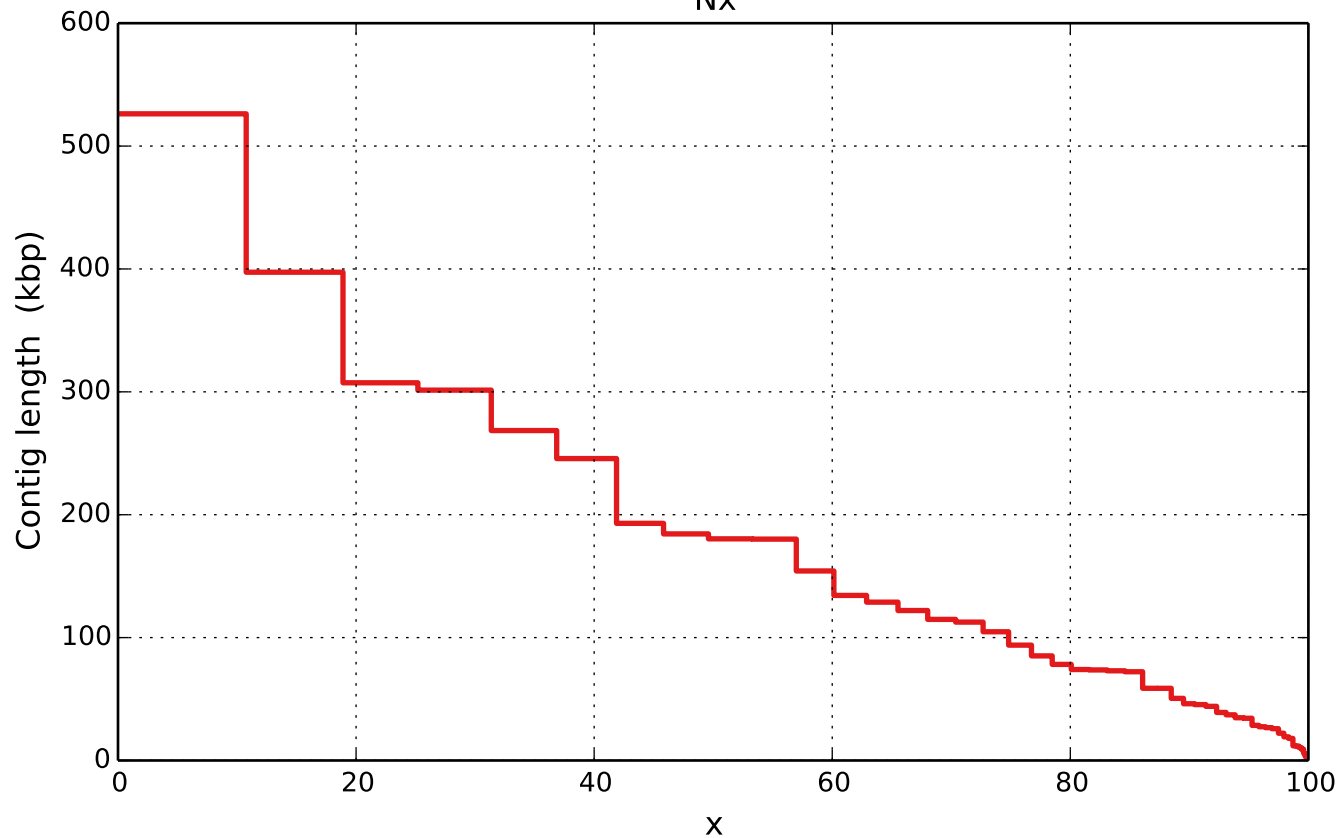
Contig



GC content



Nx



— dataset\_488.dat