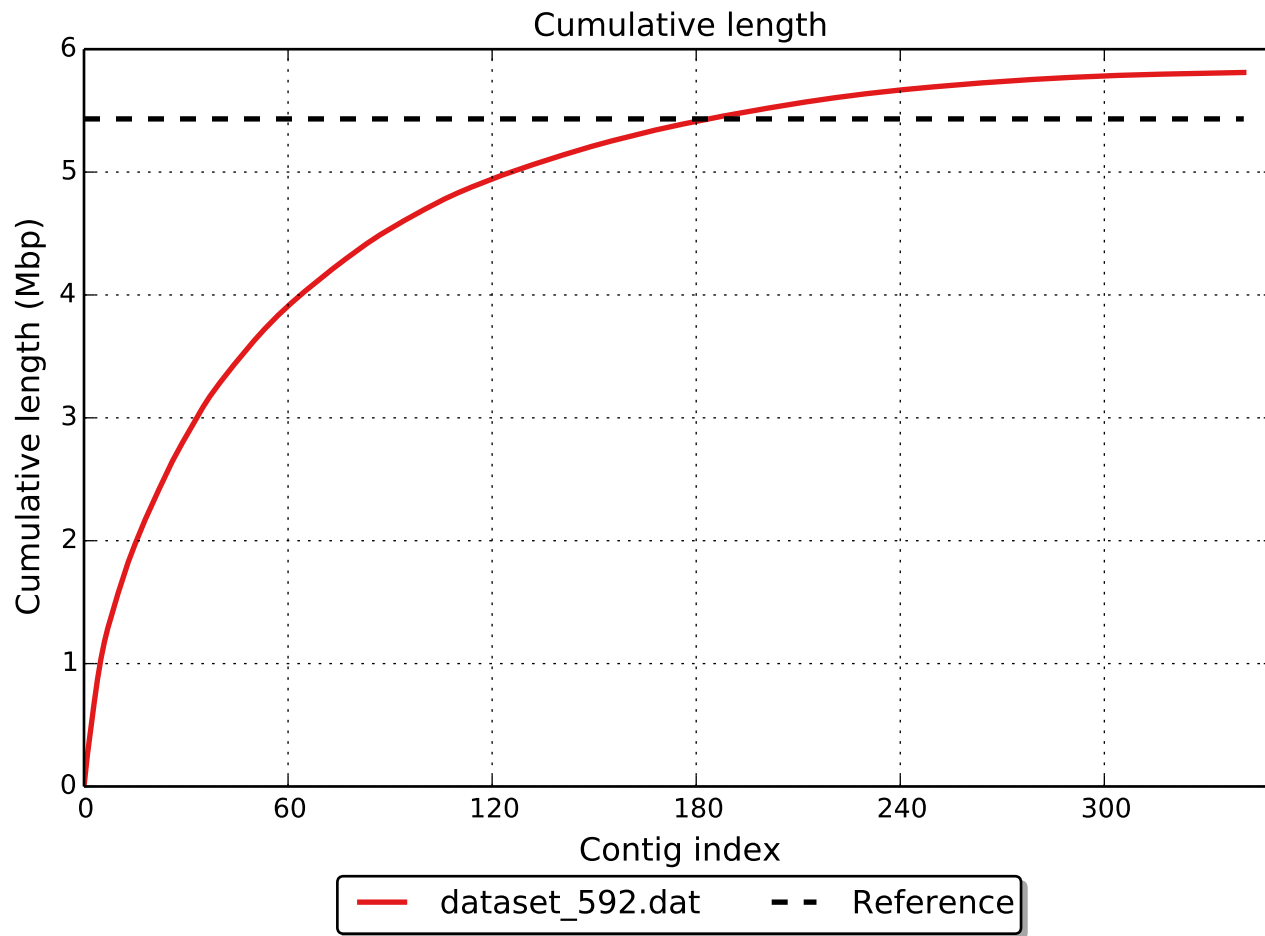
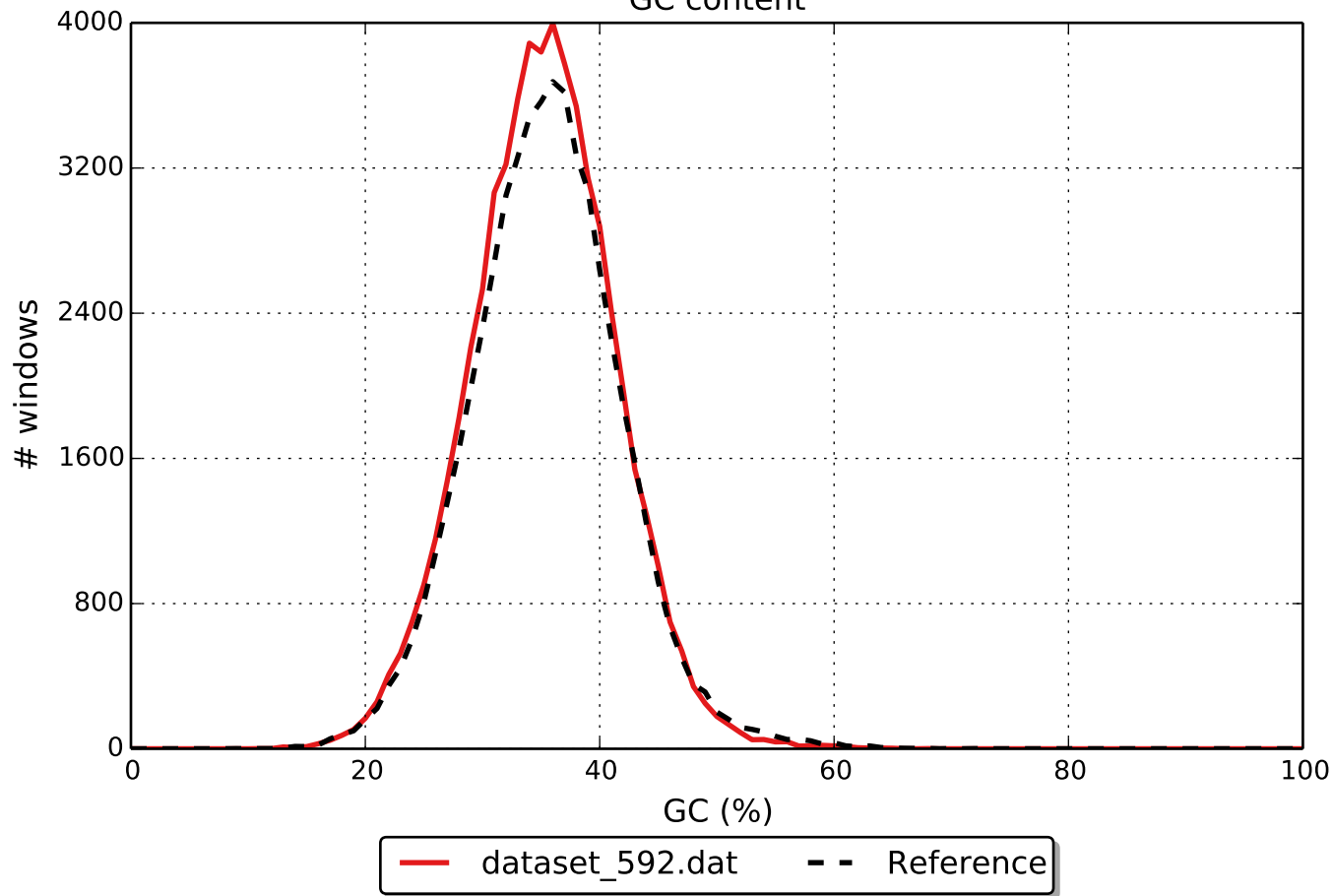


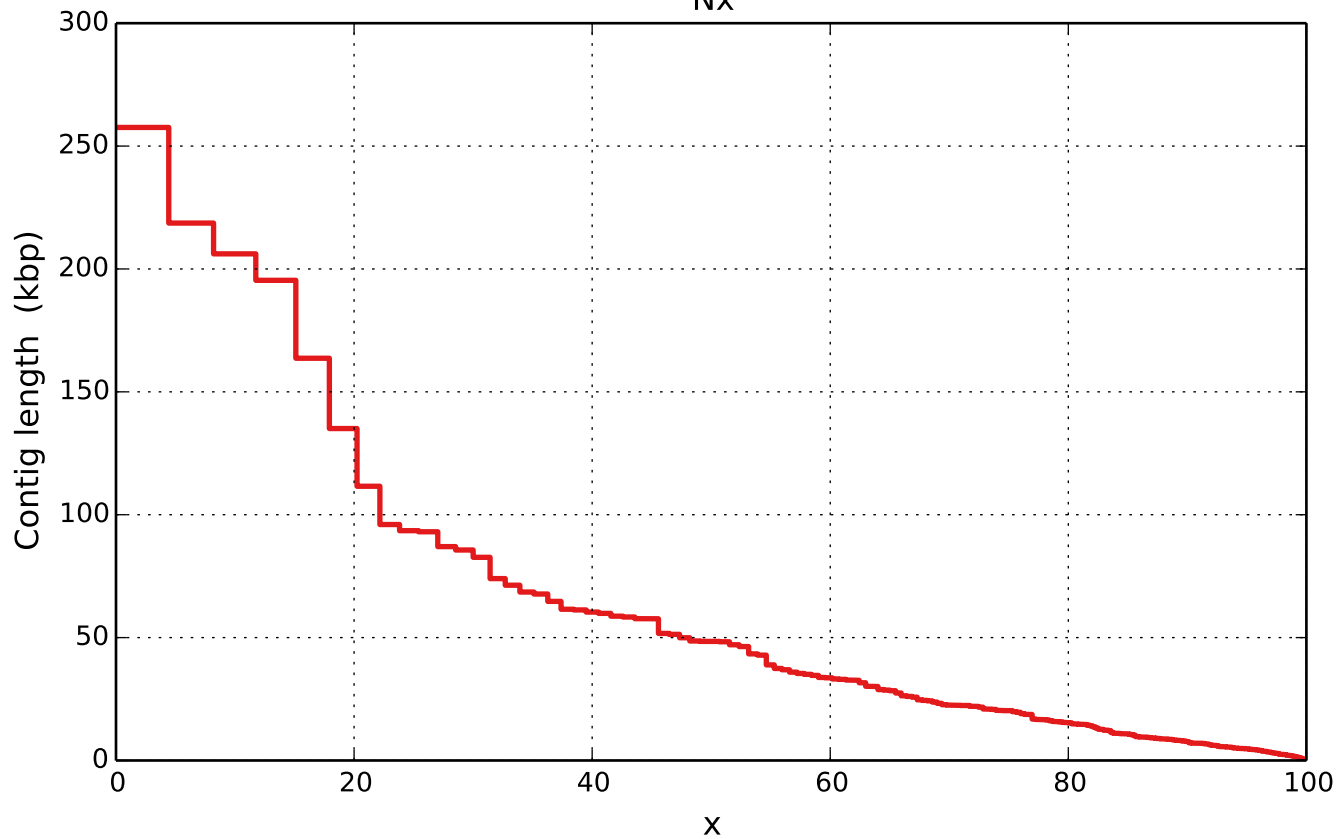
B_cereus_ATCC_1098



GC content

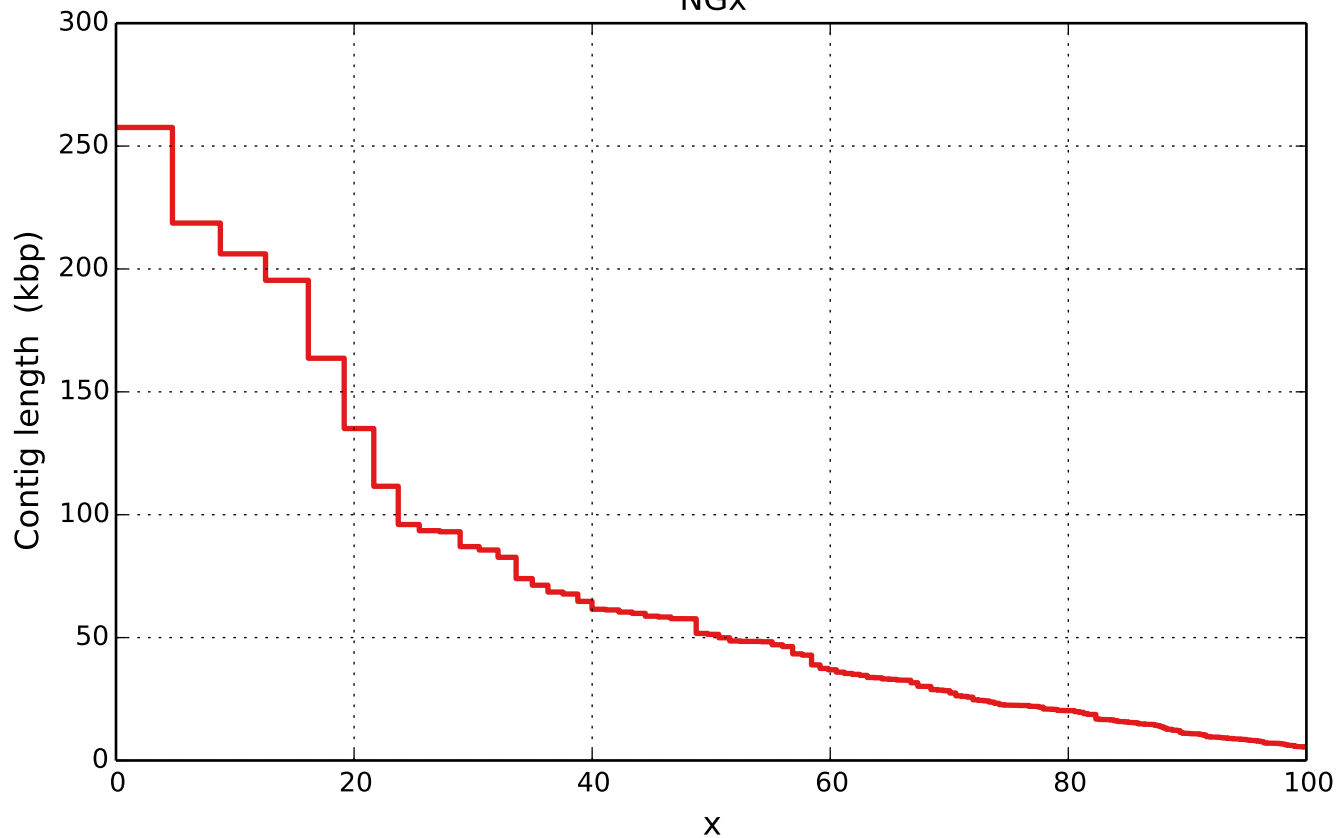


Nx

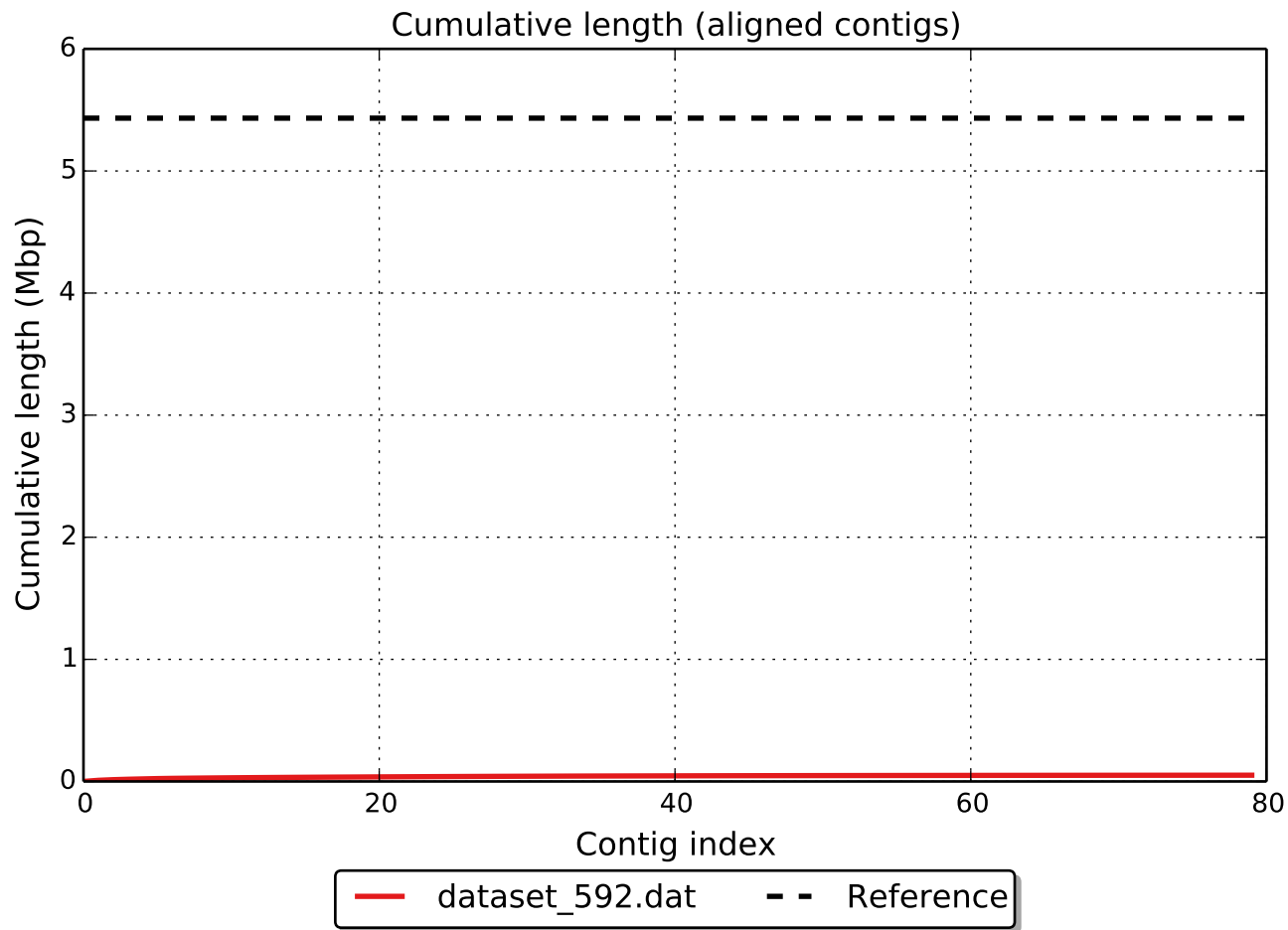


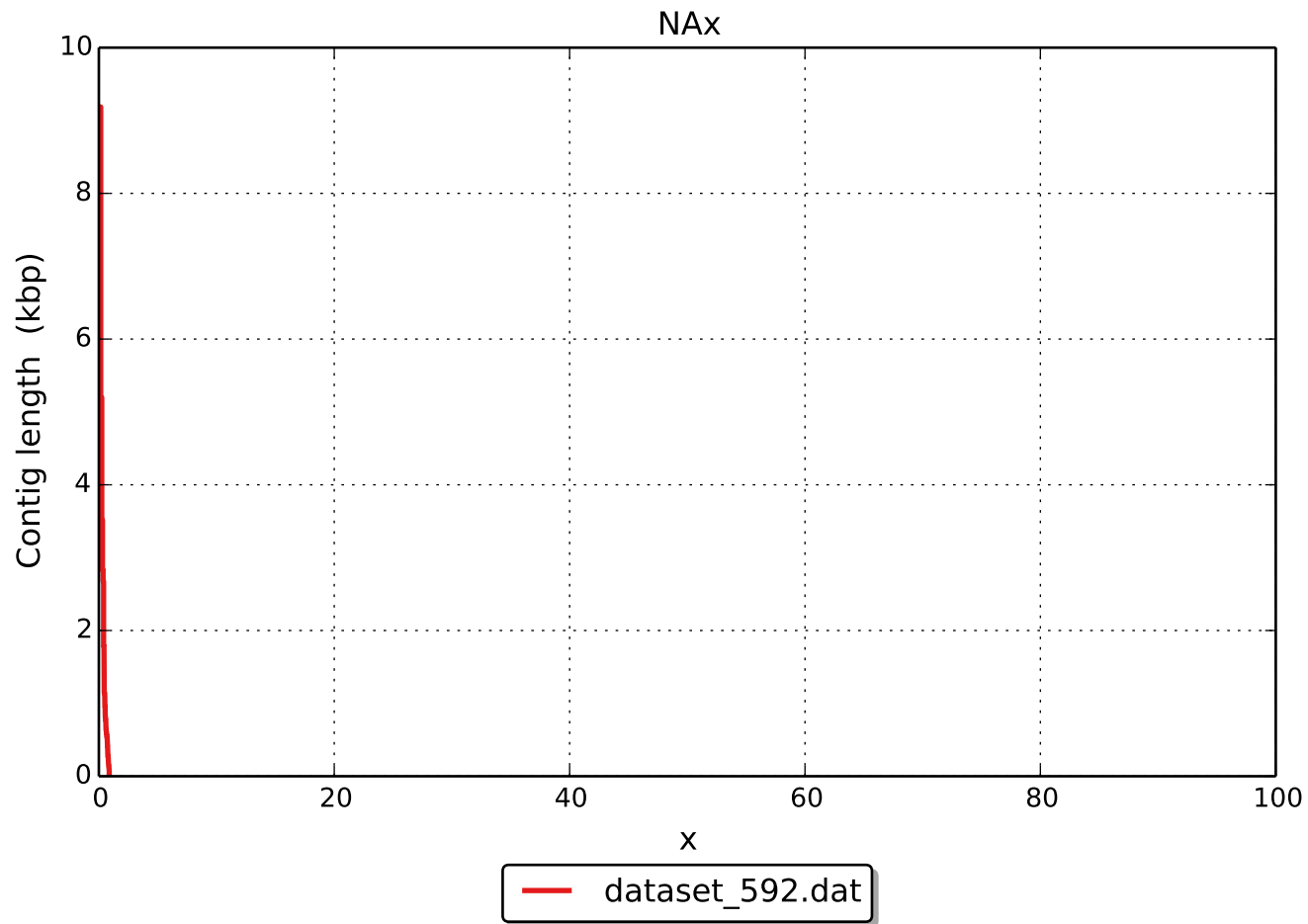
dataset_592.dat

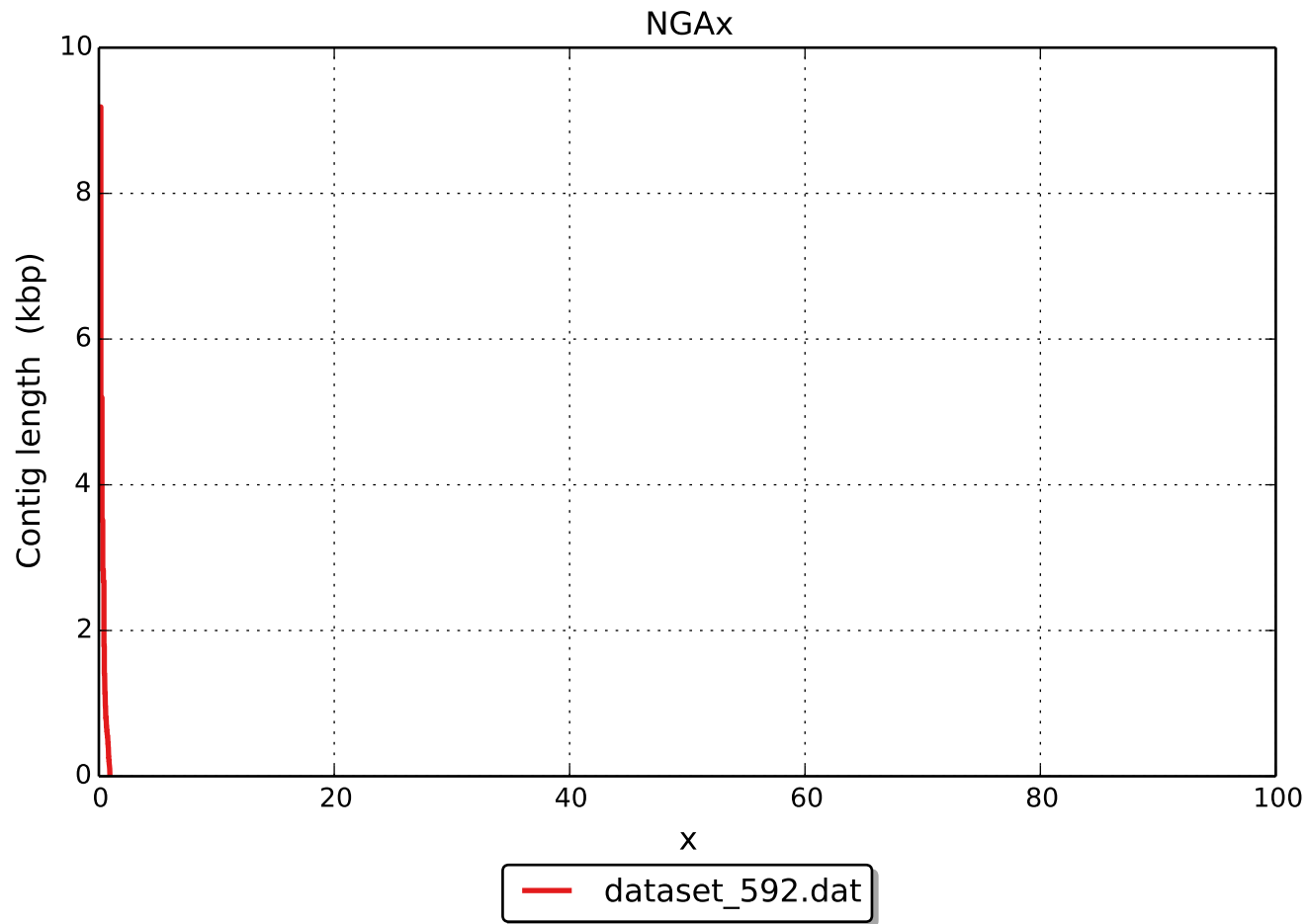
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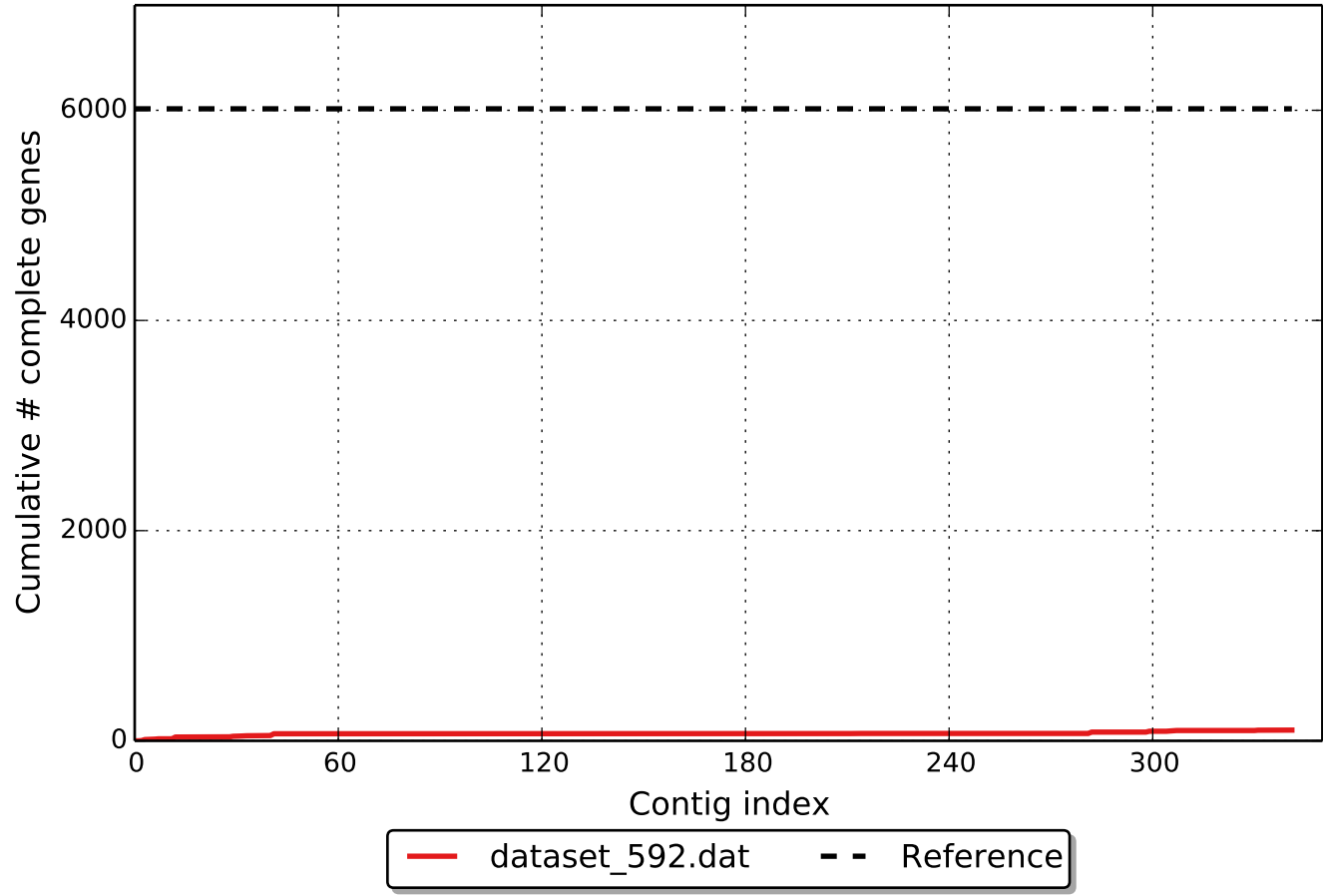
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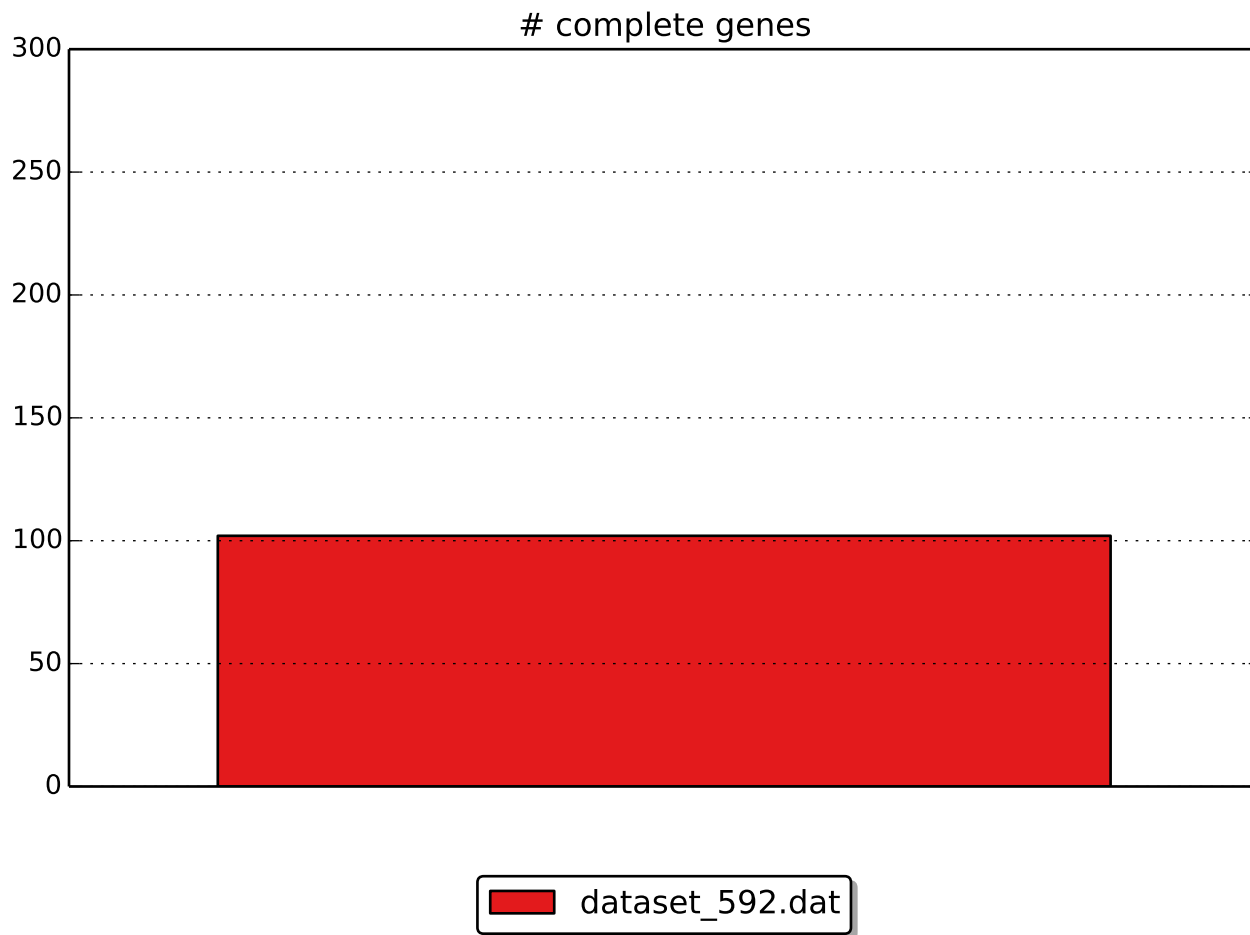


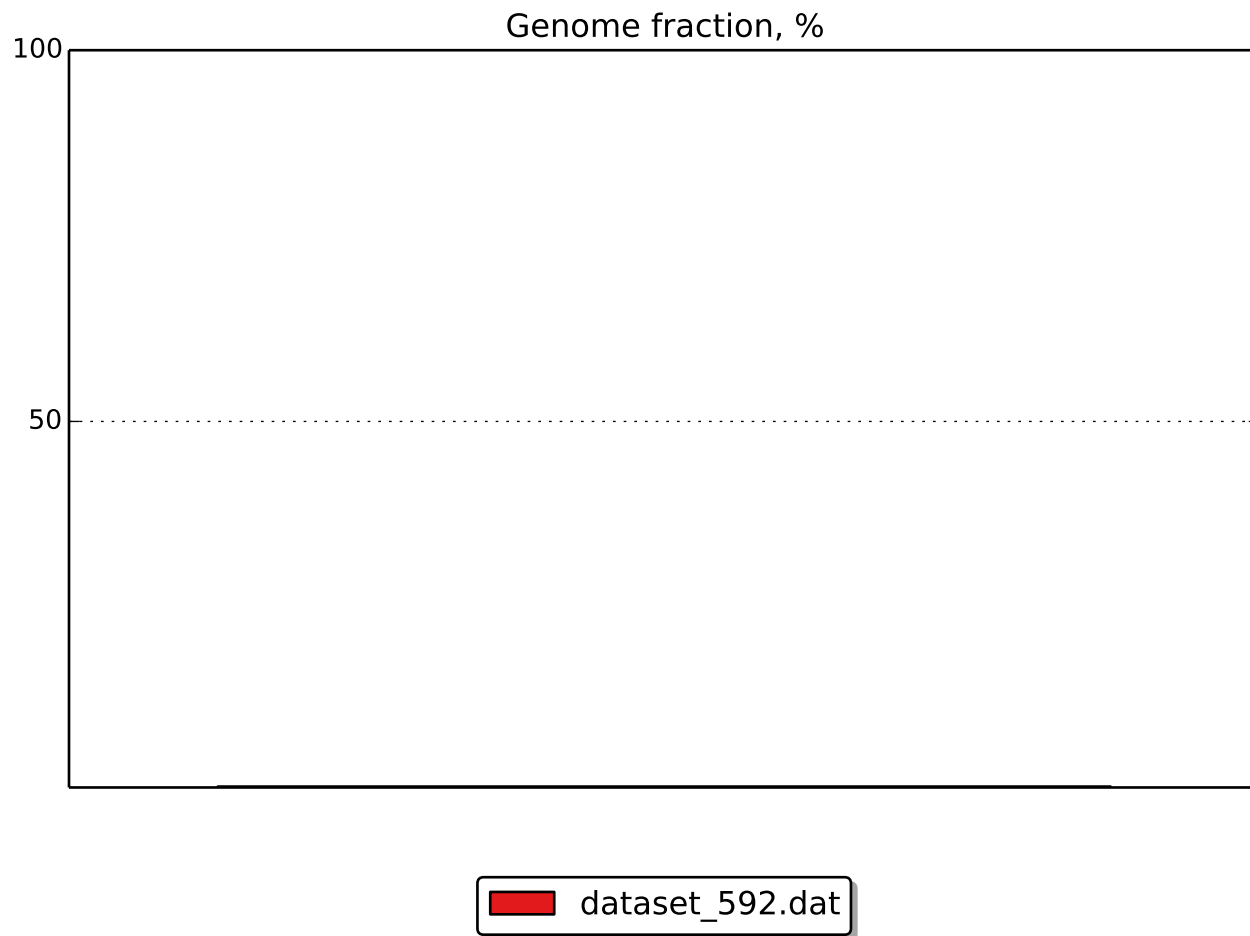




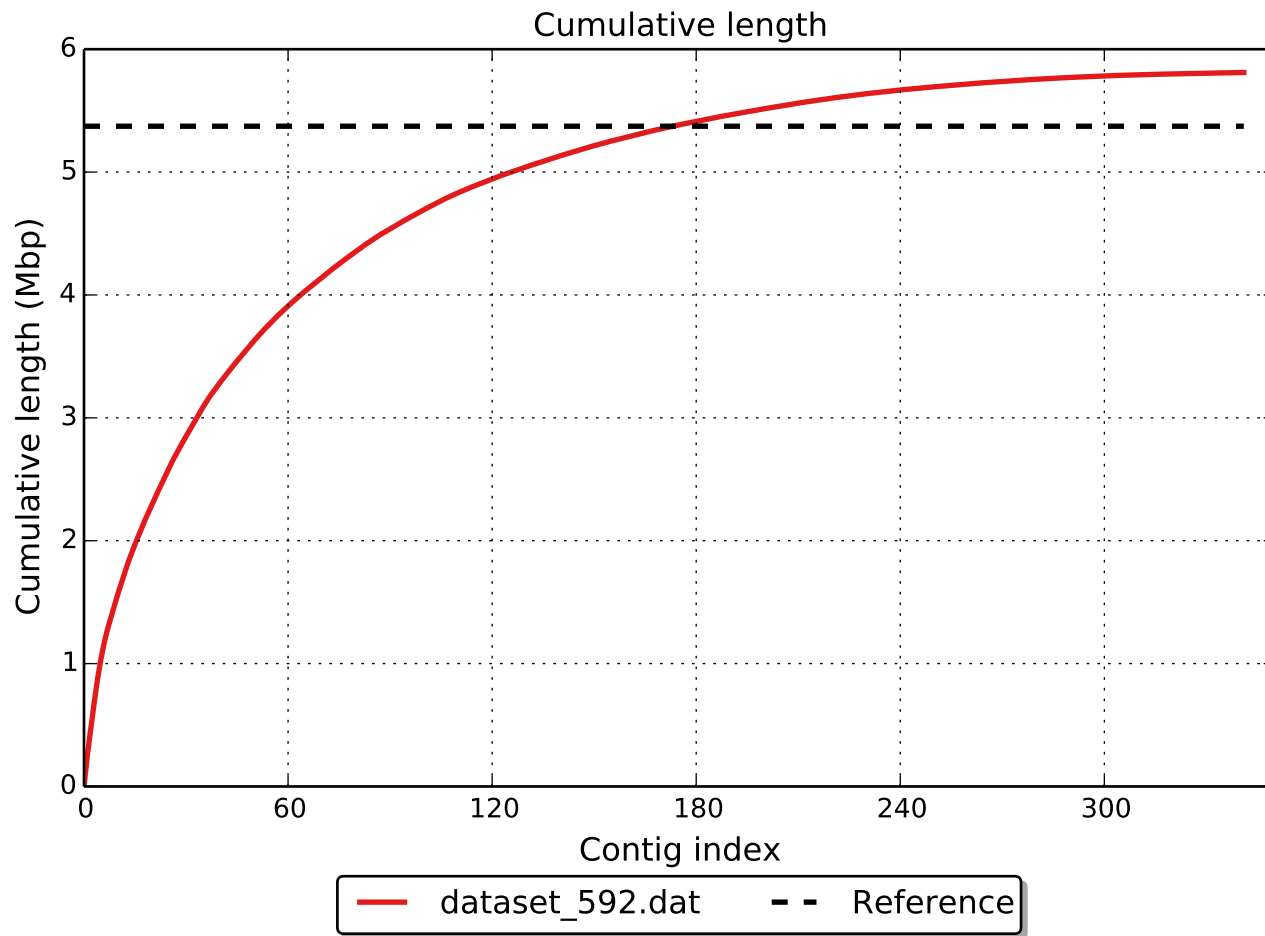
Cumulative # complete genes



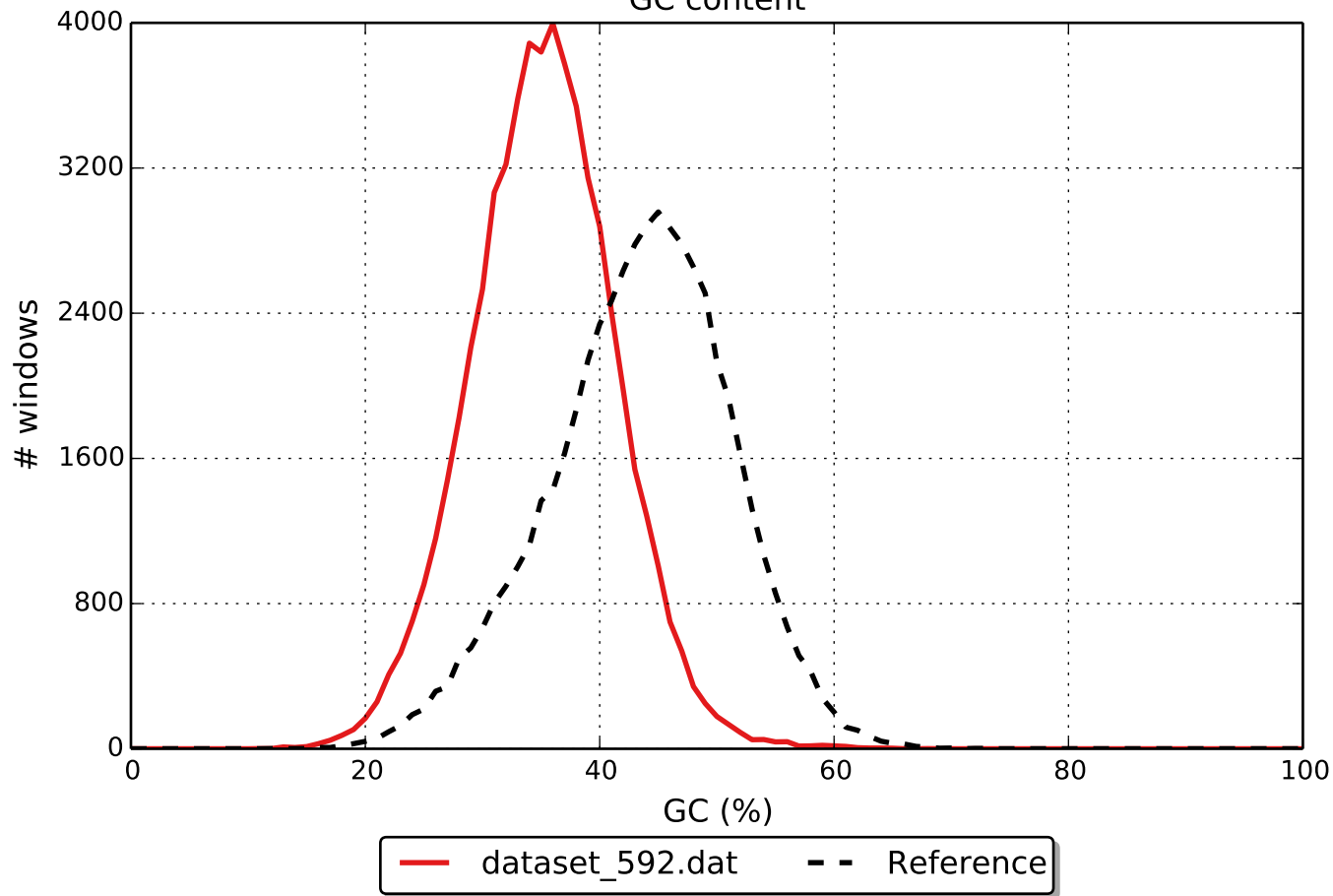




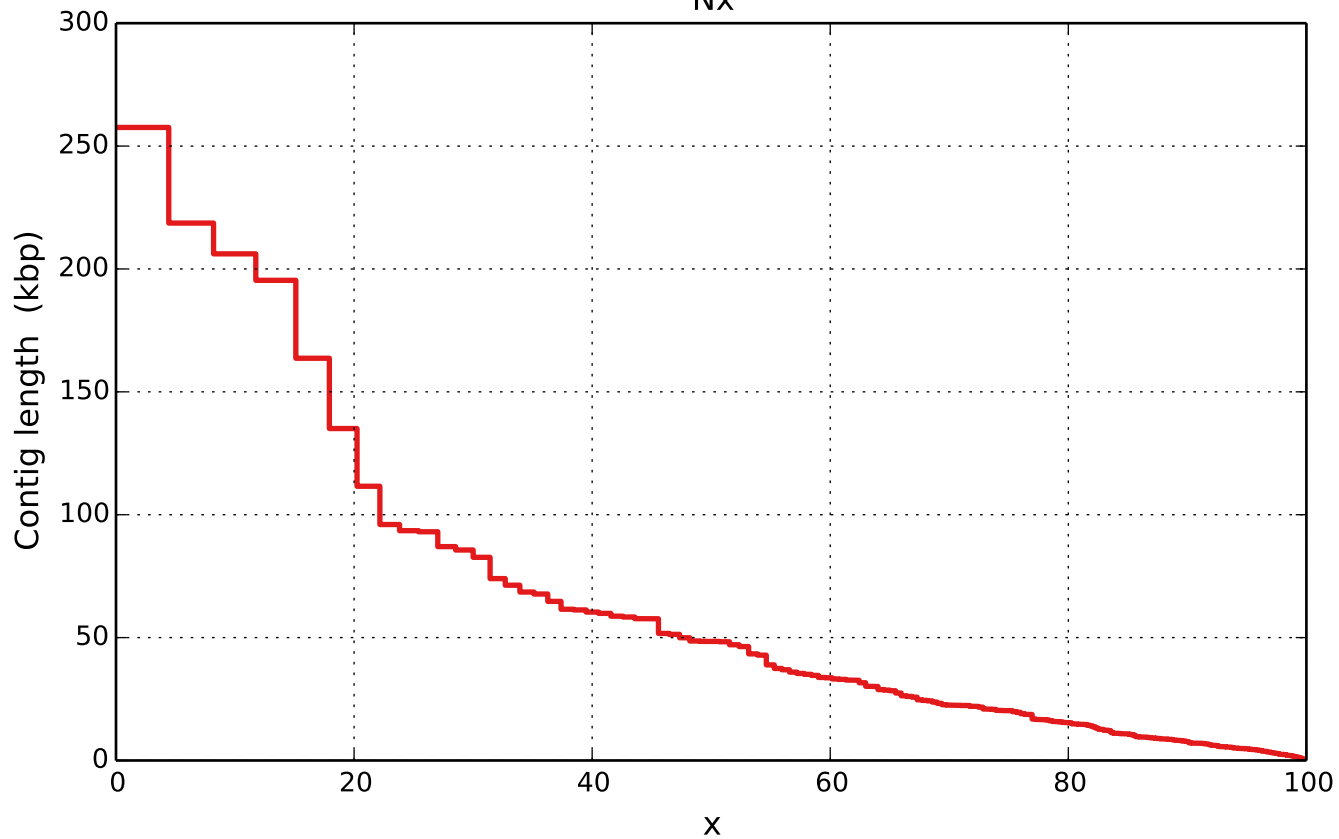
B_fragilis_638R



GC content

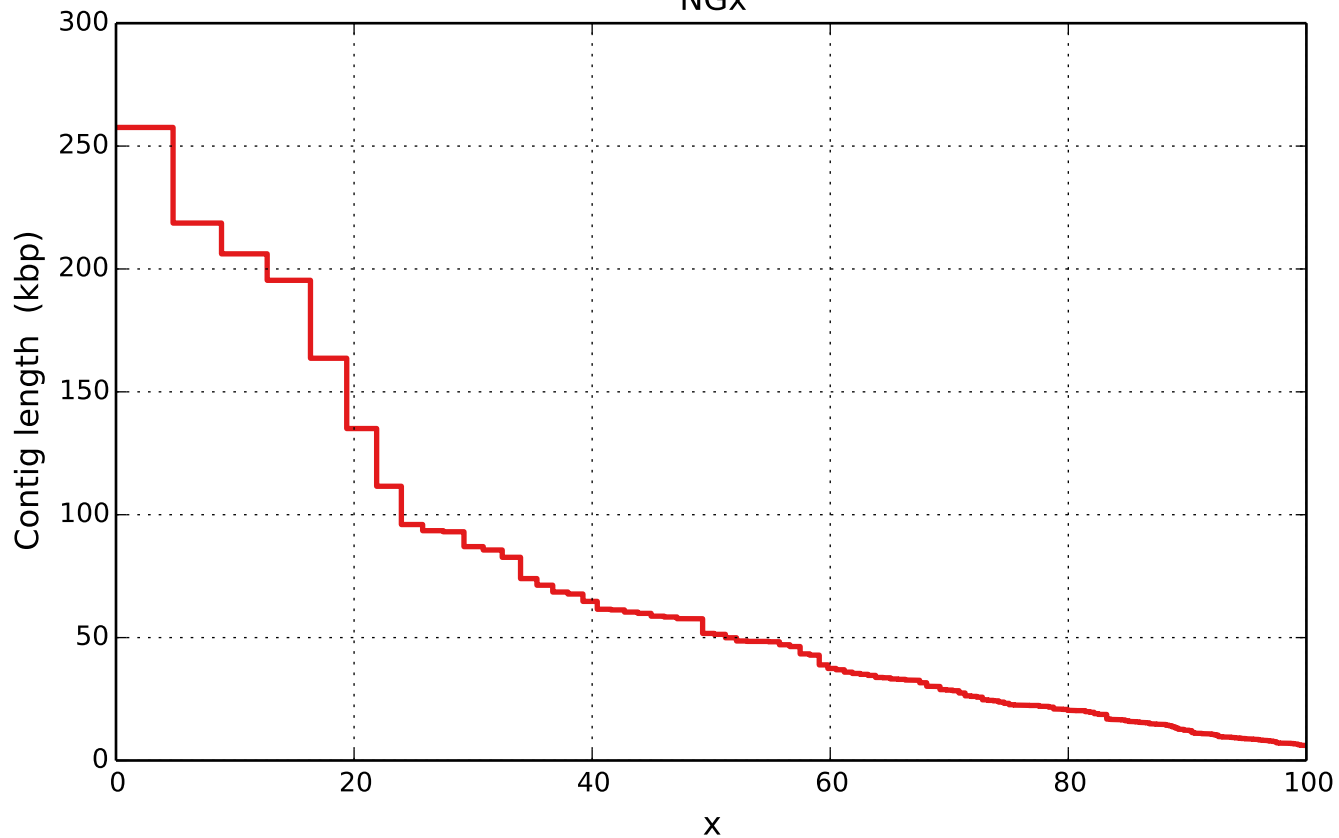


Nx



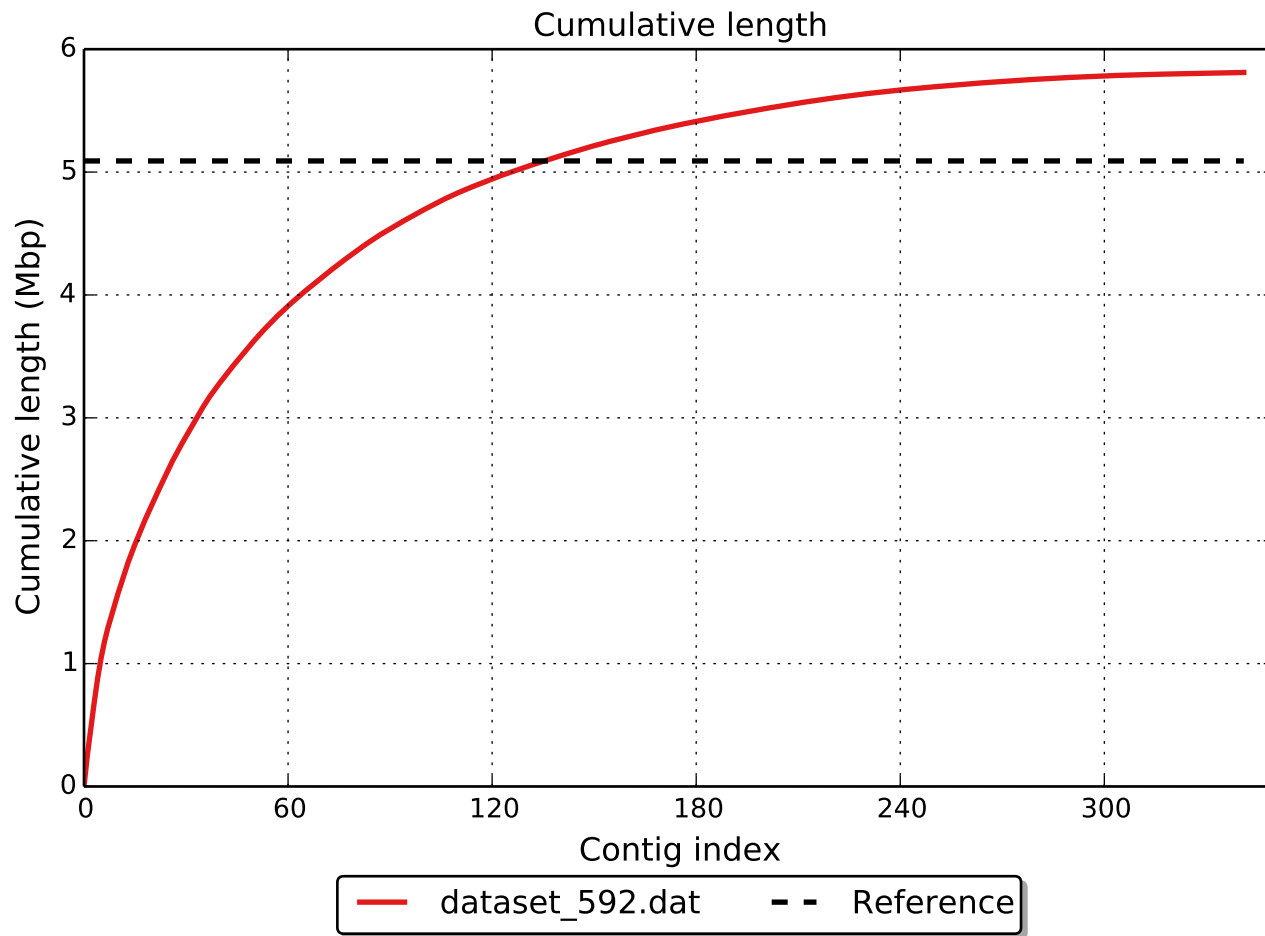
dataset_592.dat

NGx

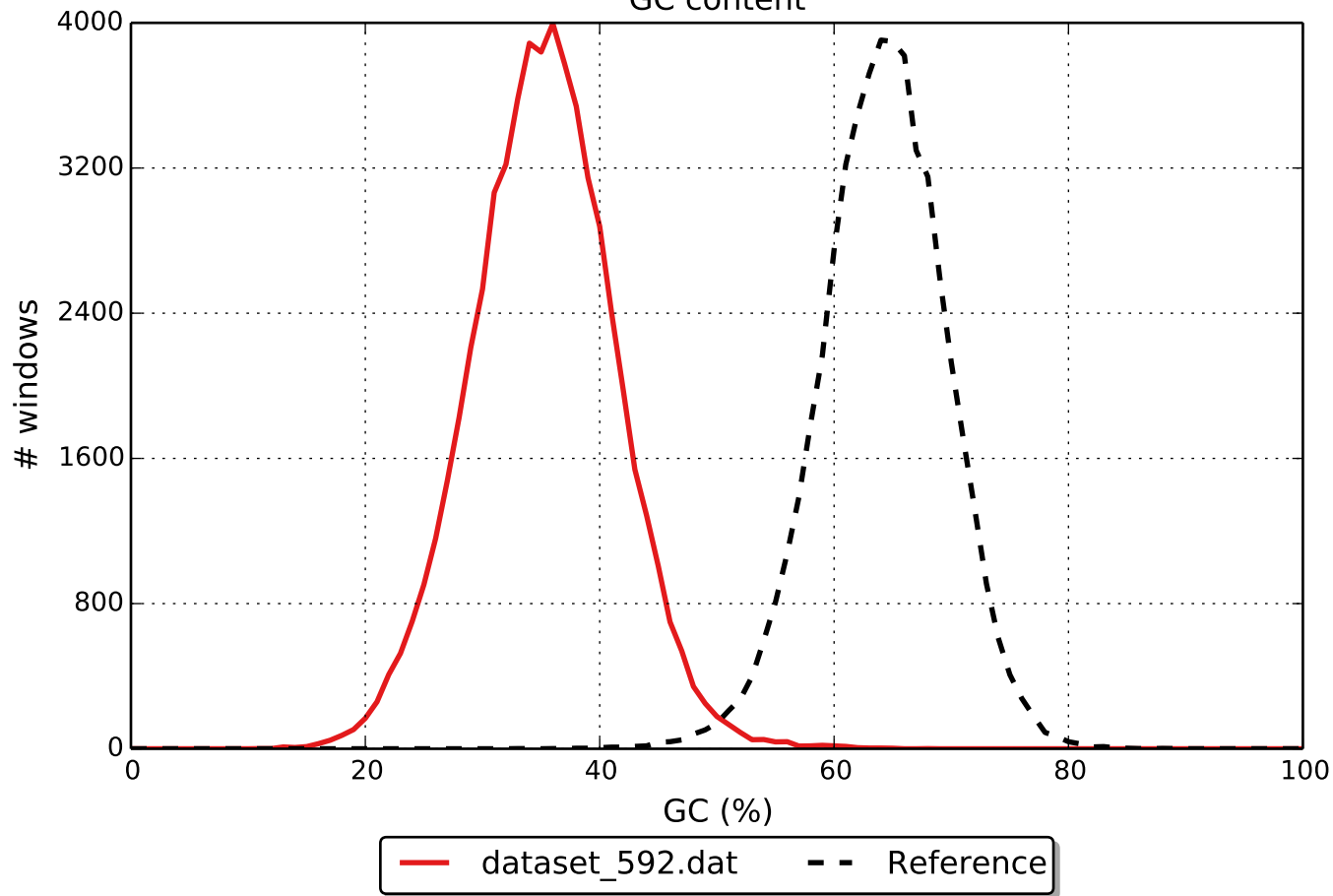


— dataset_592.dat

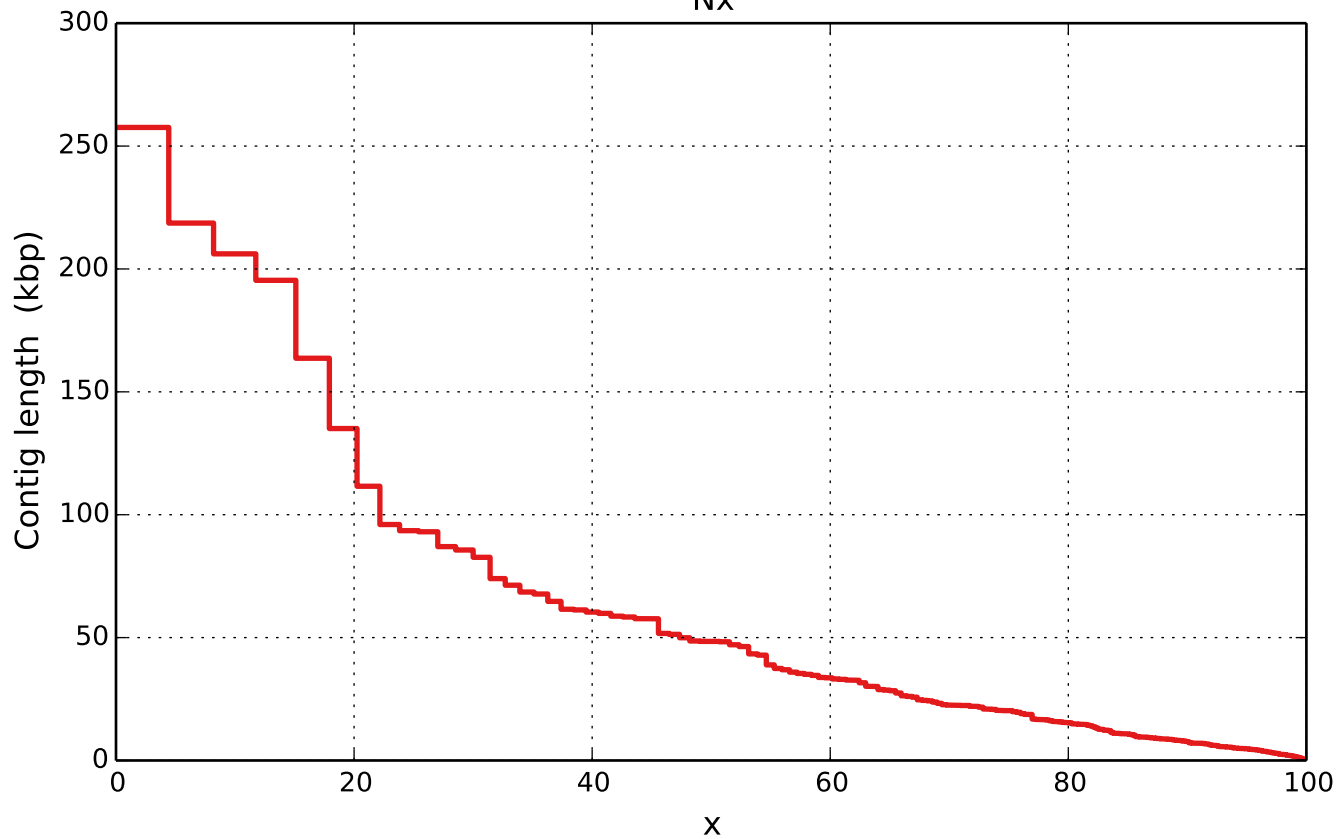
M_abscessus_ATCC_19977



GC content

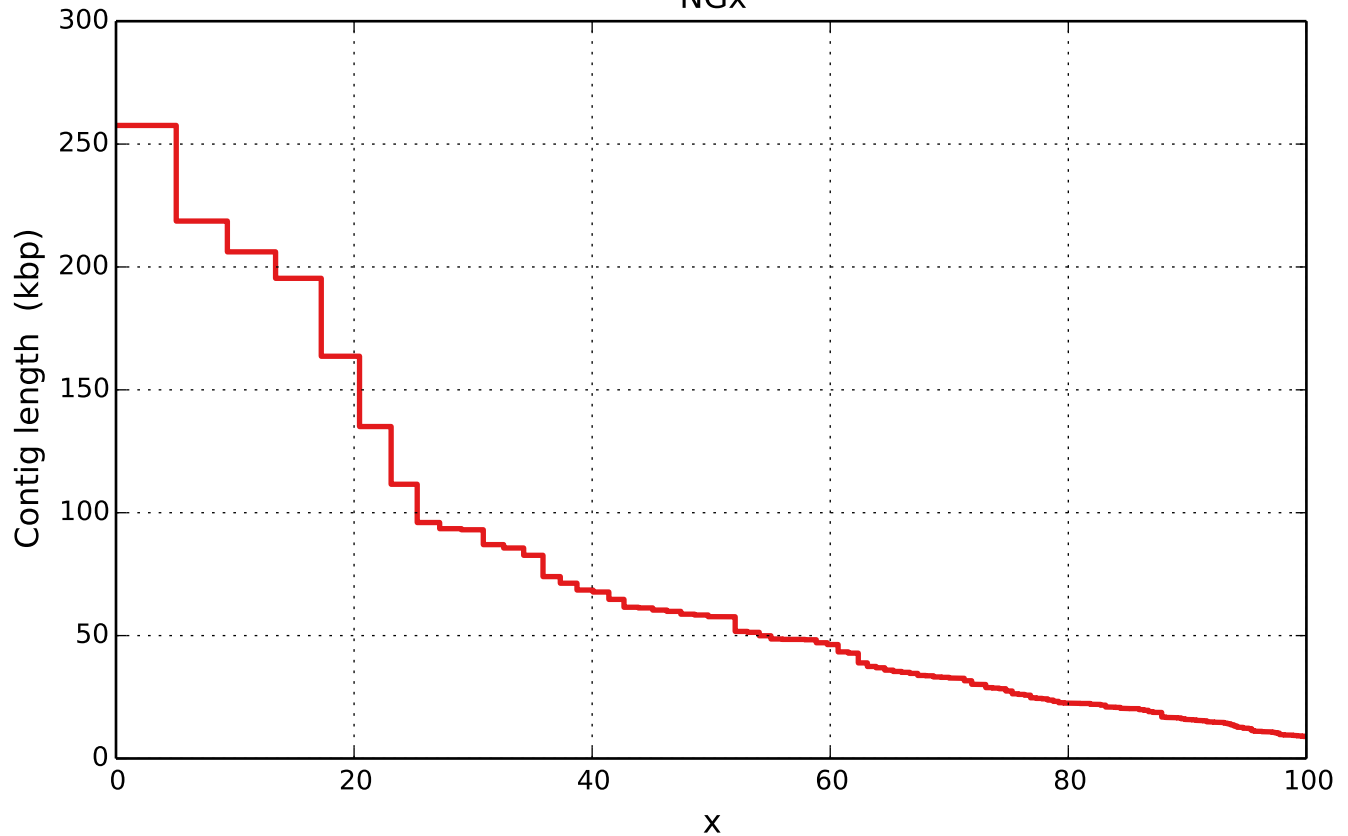


Nx



— dataset_592.dat

NGx



— dataset_592.dat