Cumulative length

deBruijnGraphQUAST_s_6.first100000   deBruijnGraphQUAST1_s_6.first100000   R
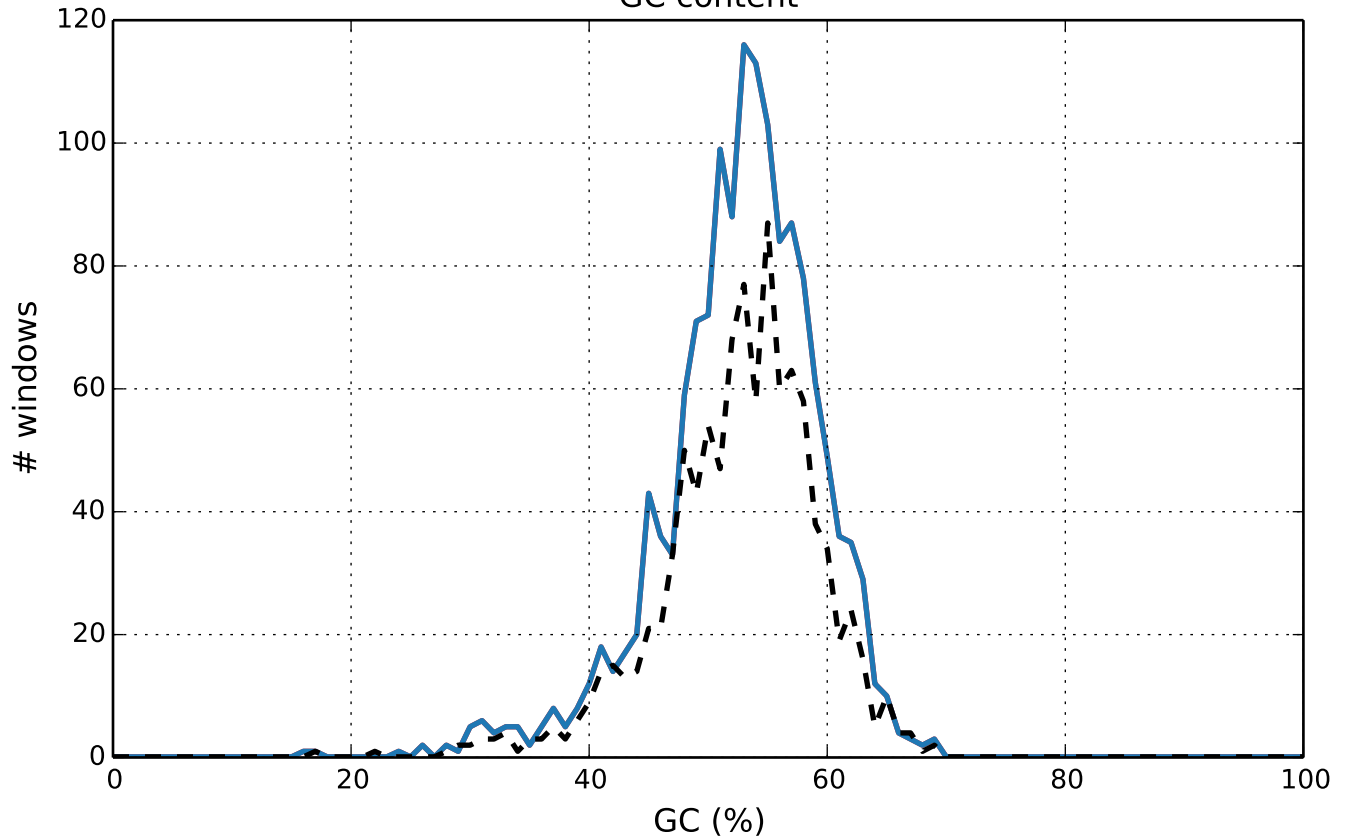
GC content

windows

GC (%)

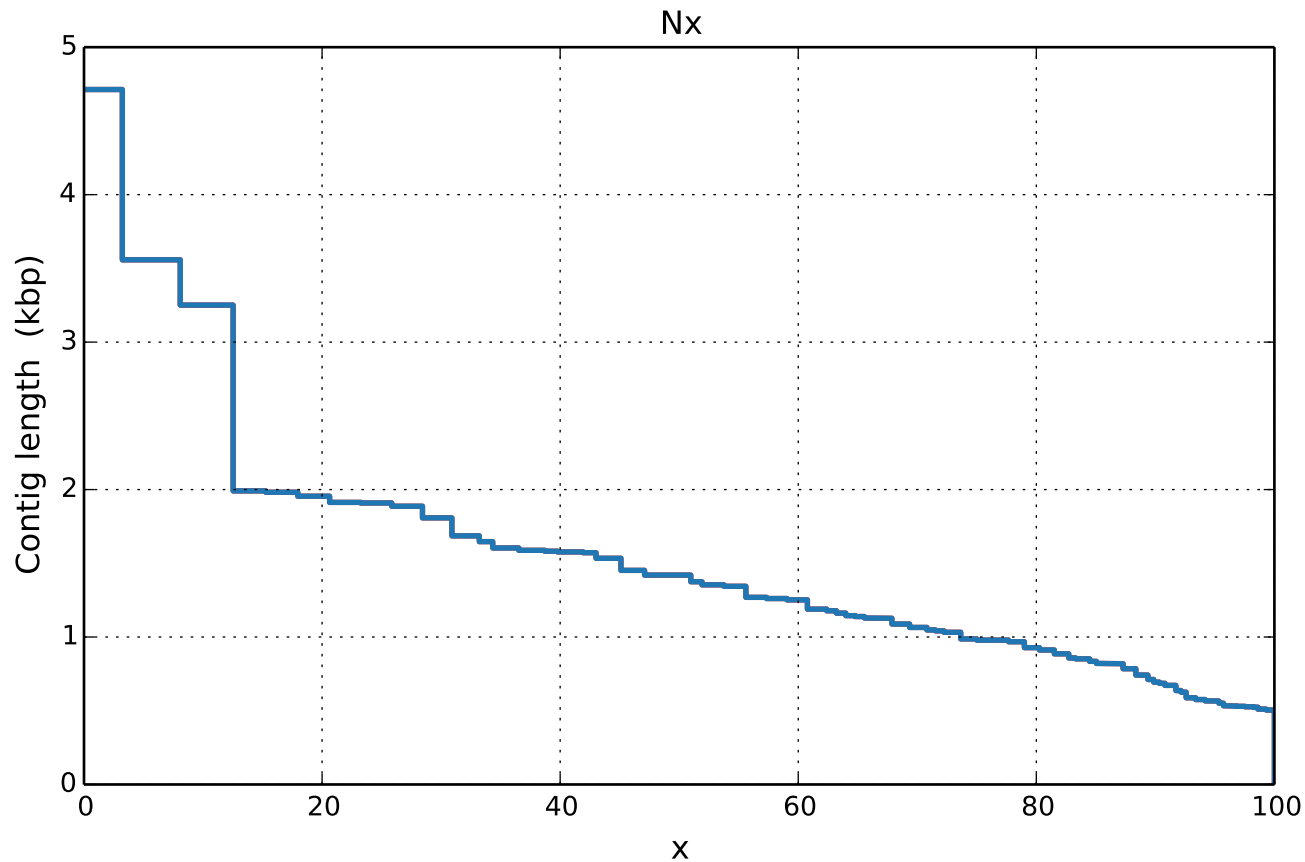deBruijnGraphQUAST_s_6.first100000 — deBruijnGraphQUAST1_s_6.first100000 - - R
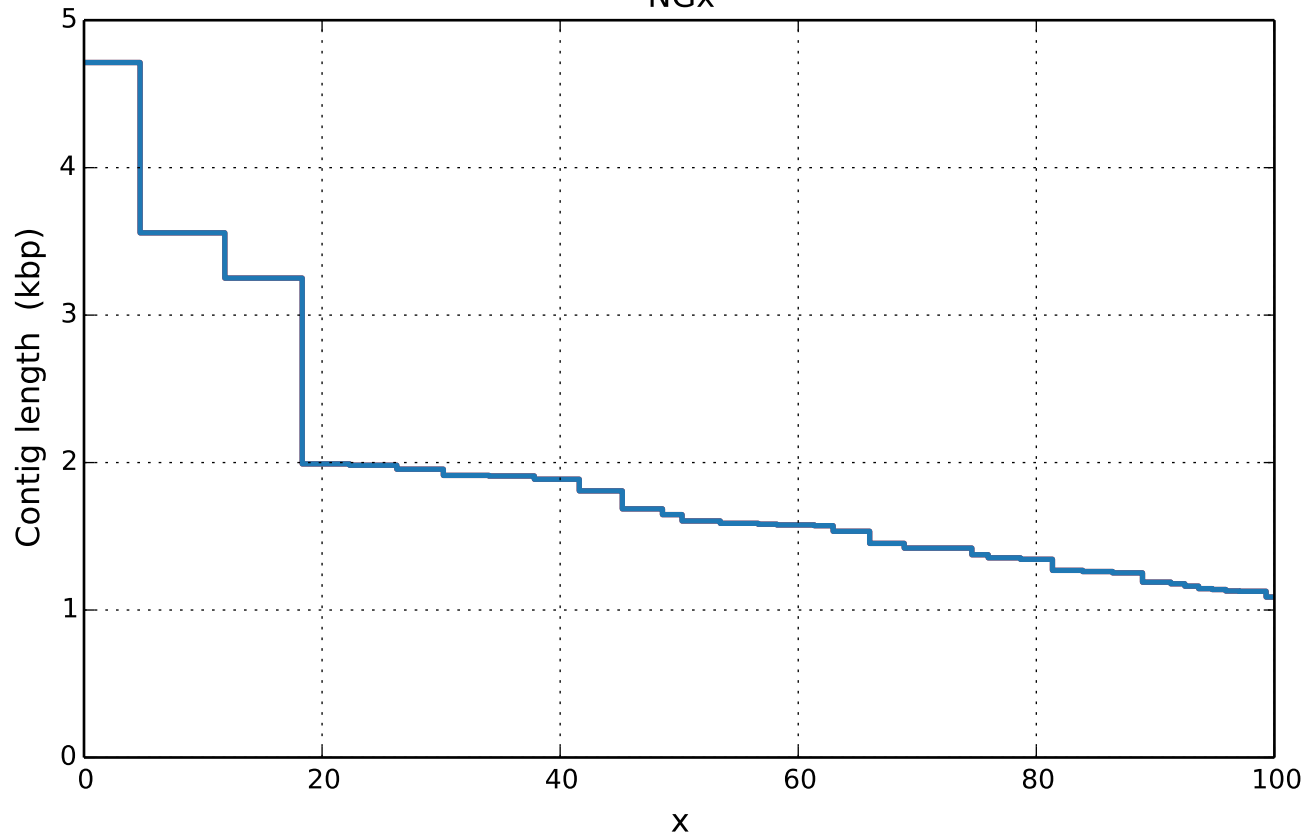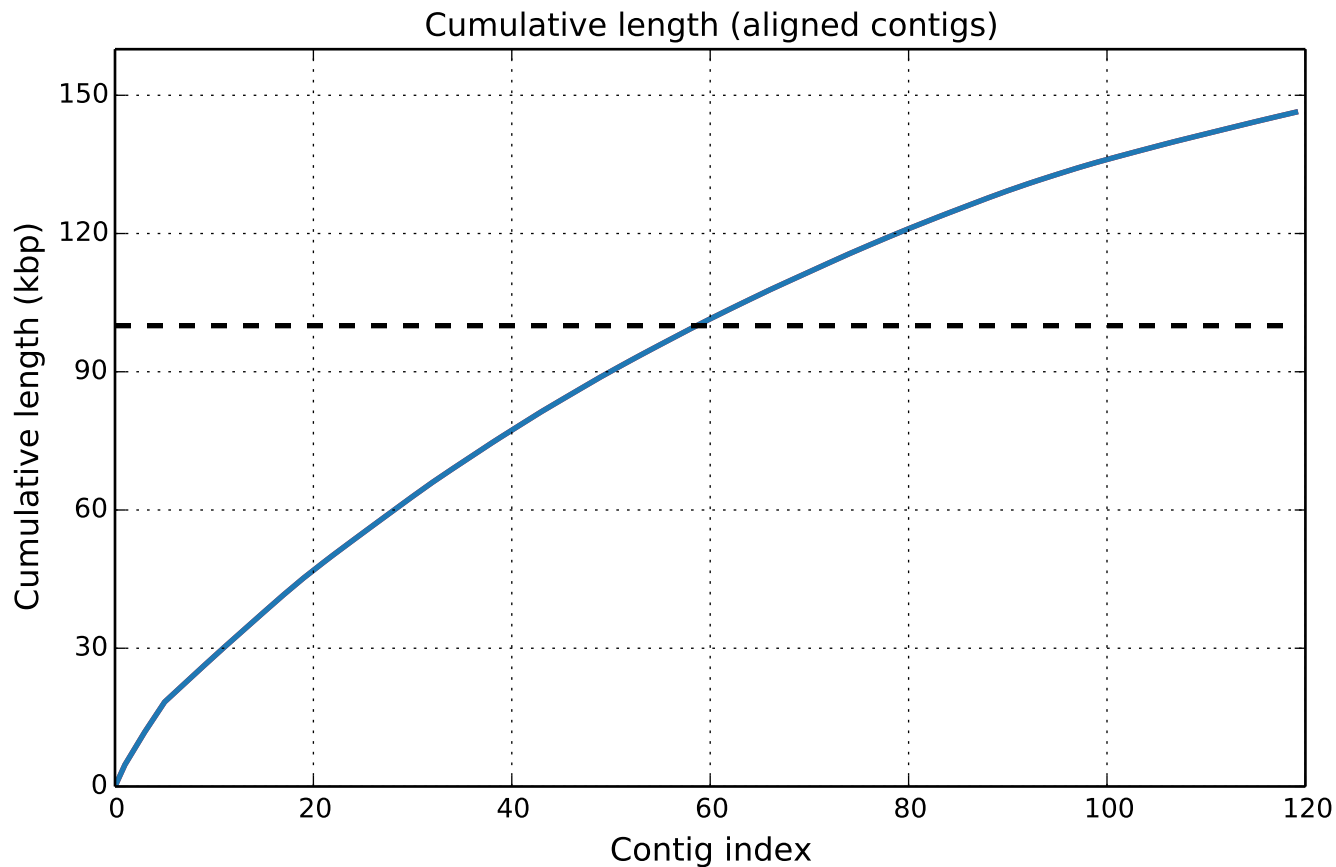
Nx

Contig length (kbp)

x

deBruijnGraphQUAST_s_6.first100000   deBruijnGraphQUAST1_s_6.first100000

NGx

Contig length (kbp)

x

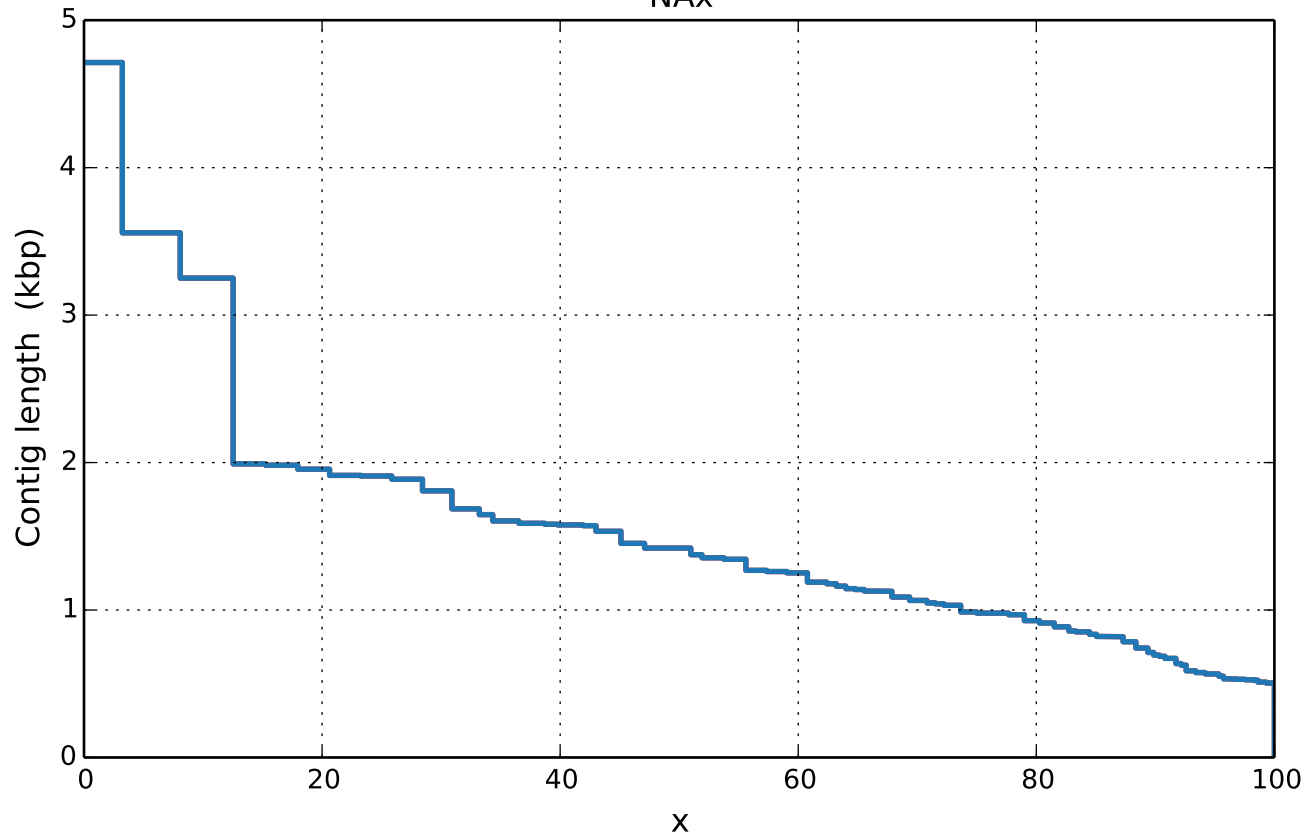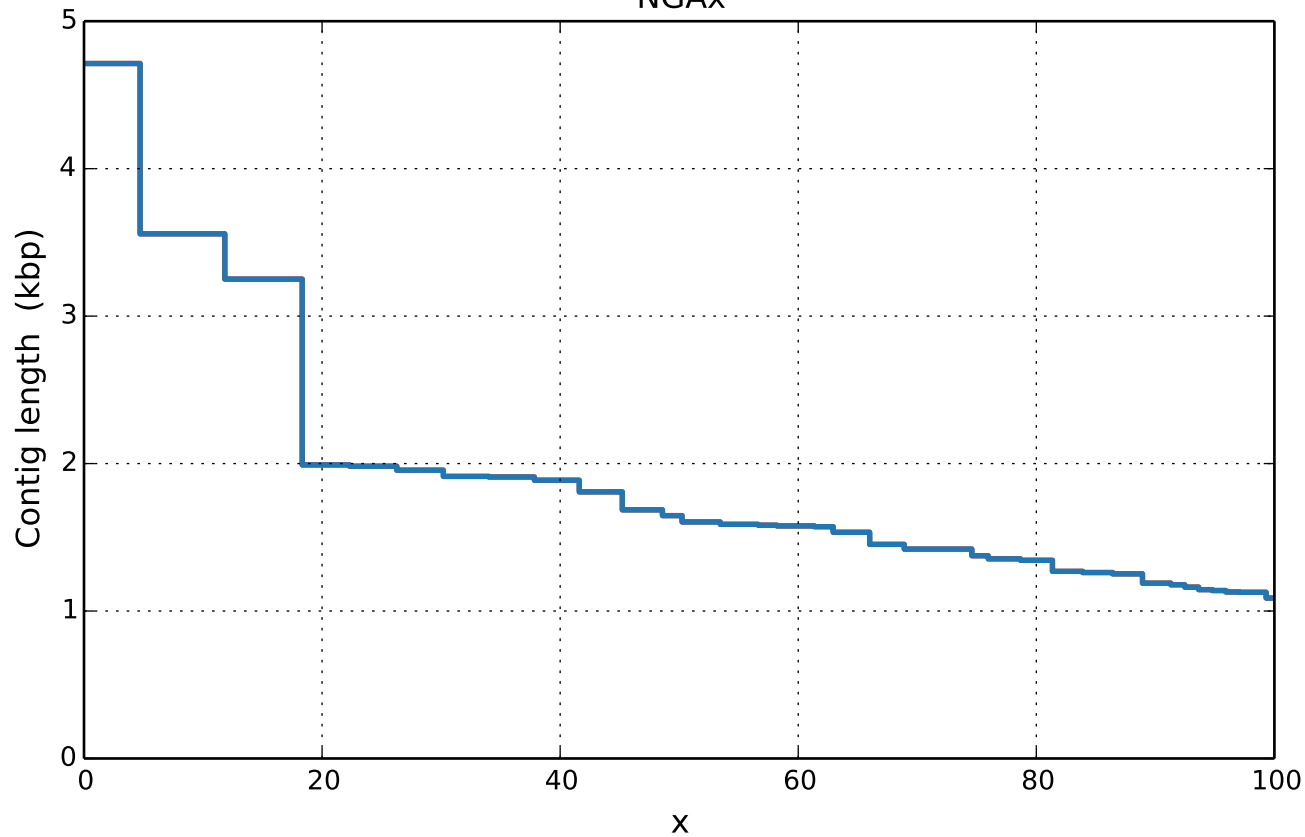deBruijnGraphQUAST_s_6.first100000   deBruijnGraphQUAST1_s_6.first100000

Cumulative length (aligned contigs)

Contig index

Cumulative length (kbp)

deBruijnGraphQUAST_s_6.first100000 — deBruijnGraphQUAST1_s_6.first100000 – – R

NAx

Contig length (kbp) vs x

deBruijnGraphQUAST_s_6.first100000

deBruijnGraphQUAST1_s_6.first100000

NGAx

Genome fraction, %

Legend: deBruijnGraphQUAST_s_6.first100000 (red), deBruijnGraphQUAST1_s_6.first100000 (blue)