

Report

	final.contigs
# contigs (>= 1000 bp)	1503
# contigs (>= 5000 bp)	1
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 1000 bp)	2312290
Total length (>= 5000 bp)	5476
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	3828
Largest contig	5476
Total length	3974624
Reference length	4641652
GC (℥)	50.74
Reference GC (℥)	50.79
N50	1117
NG50	996
N75	784
NG75	642
L50	1195
LG50	1512
L75	2260
LG75	2963
# misassemblies	1
# misassembled contigs	1
Misassembled contigs length	3047
# local misassemblies	0
# unaligned contigs	0 + 0 part
Unaligned length	0
Genome fraction (℥)	81.896
Duplication ratio	1.046
# N's per 100 kbp	0.00
# mismatches per 100 kbp	39.28
# indels per 100 kbp	0.00
Largest alignment	5476
NA50	1117
NGA50	996
NA75	784
NGA75	642
LA50	1195
LGA50	1512
LA75	2260
LGA75	2963

All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Misassemblies report

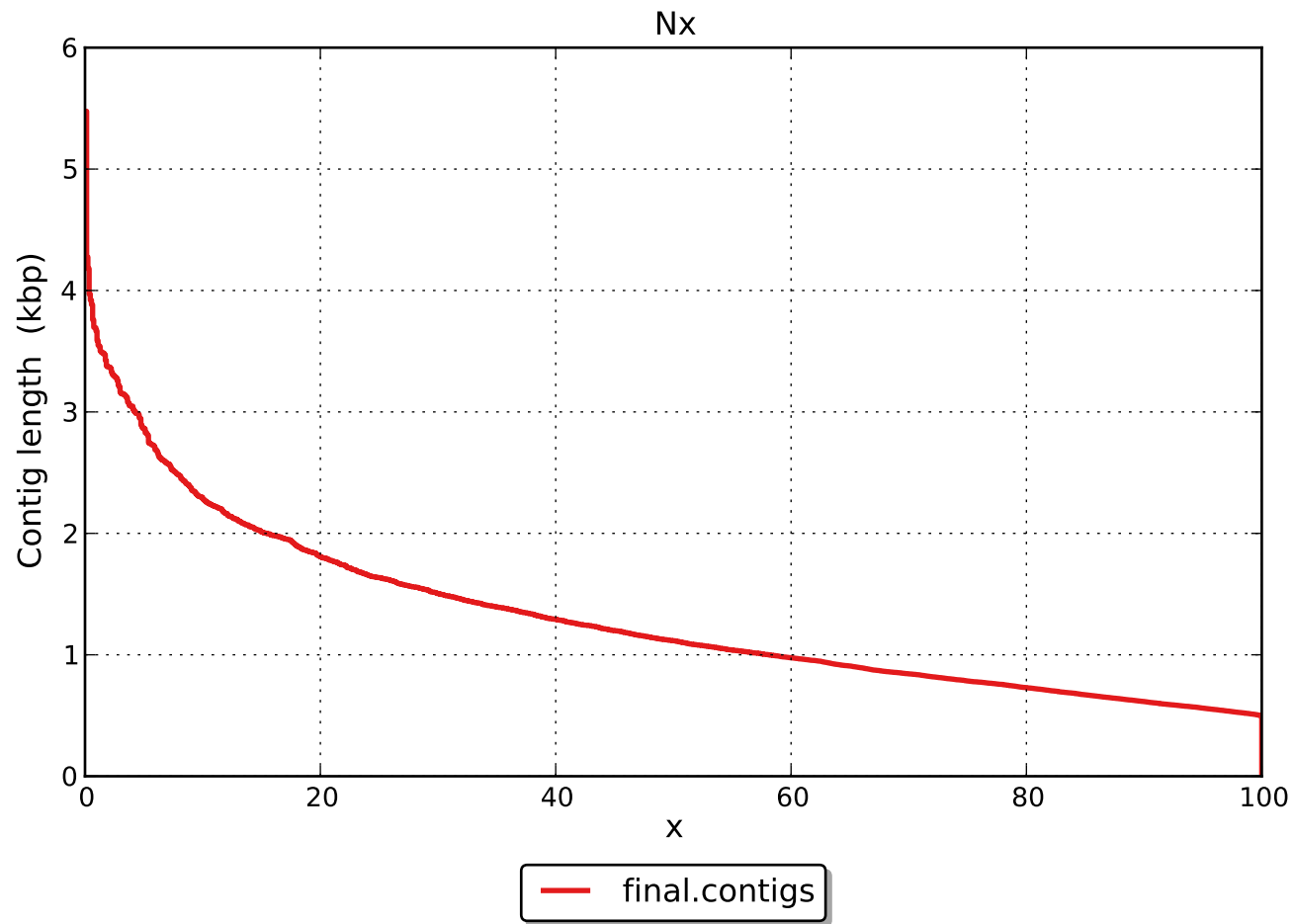
	final.contigs
# misassemblies	1
# relocations	1
# translocations	0
# inversions	0
# possibly misassembled contigs	0
# misassembled contigs	1
Misassembled contigs length	3047
# local misassemblies	0
# mismatches	1493
# indels	0
# short indels	0
# long indels	0
Indels length	0

All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

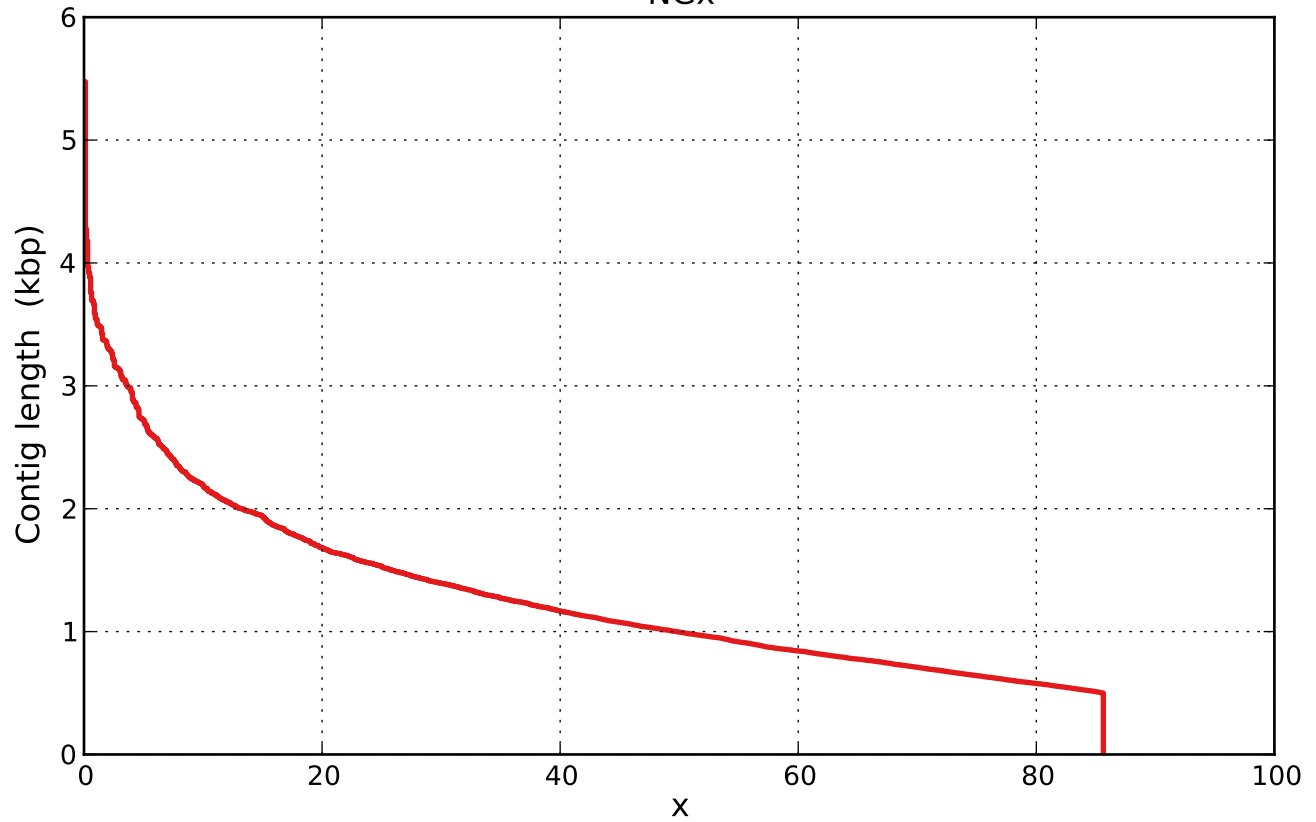
Unaligned report

	final.contigs
# fully unaligned contigs	0
Fully unaligned length	0
# partially unaligned contigs	0
# with misassembly	0
# both parts are significant	0
Partially unaligned length	0
# N's	0

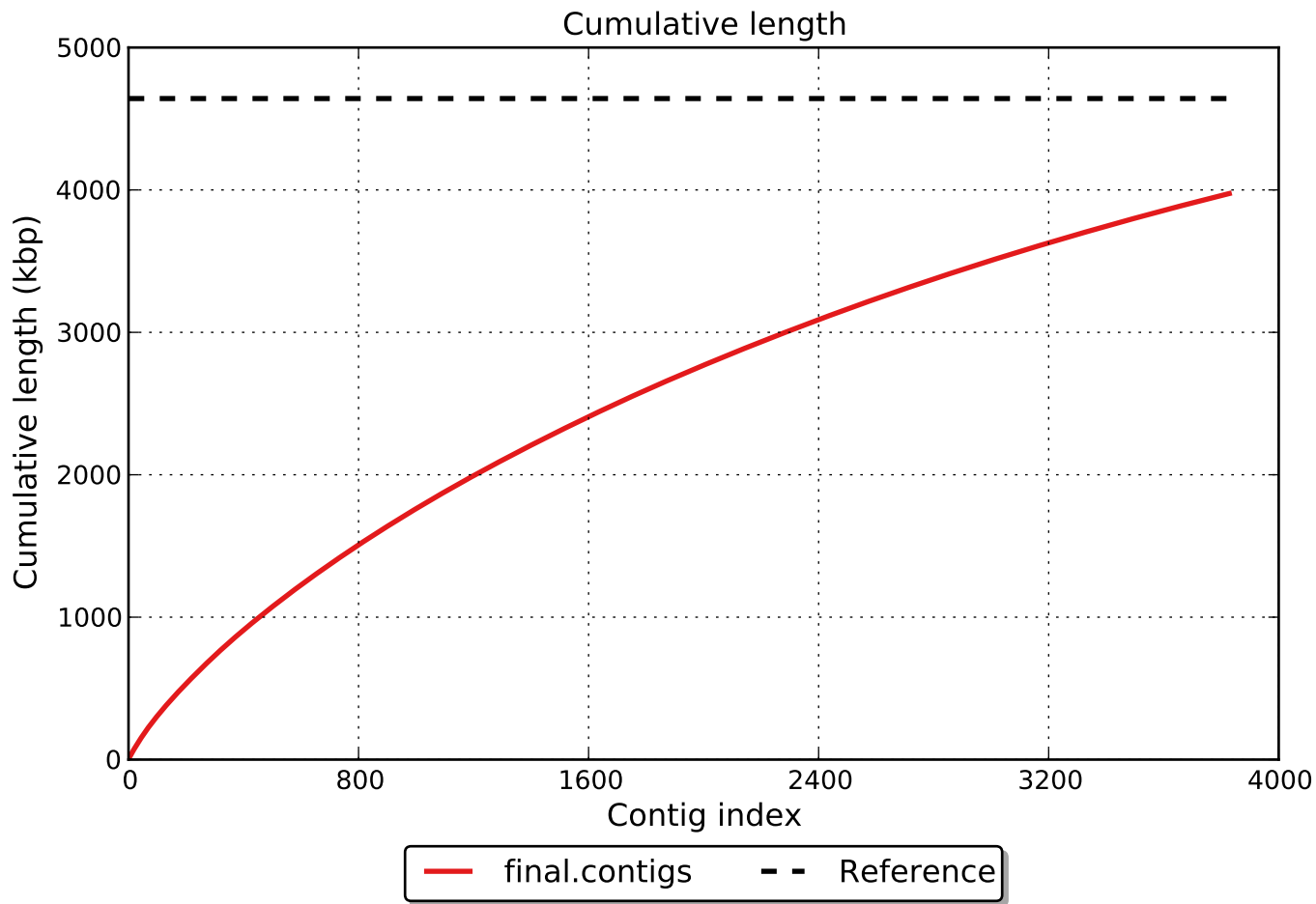
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).



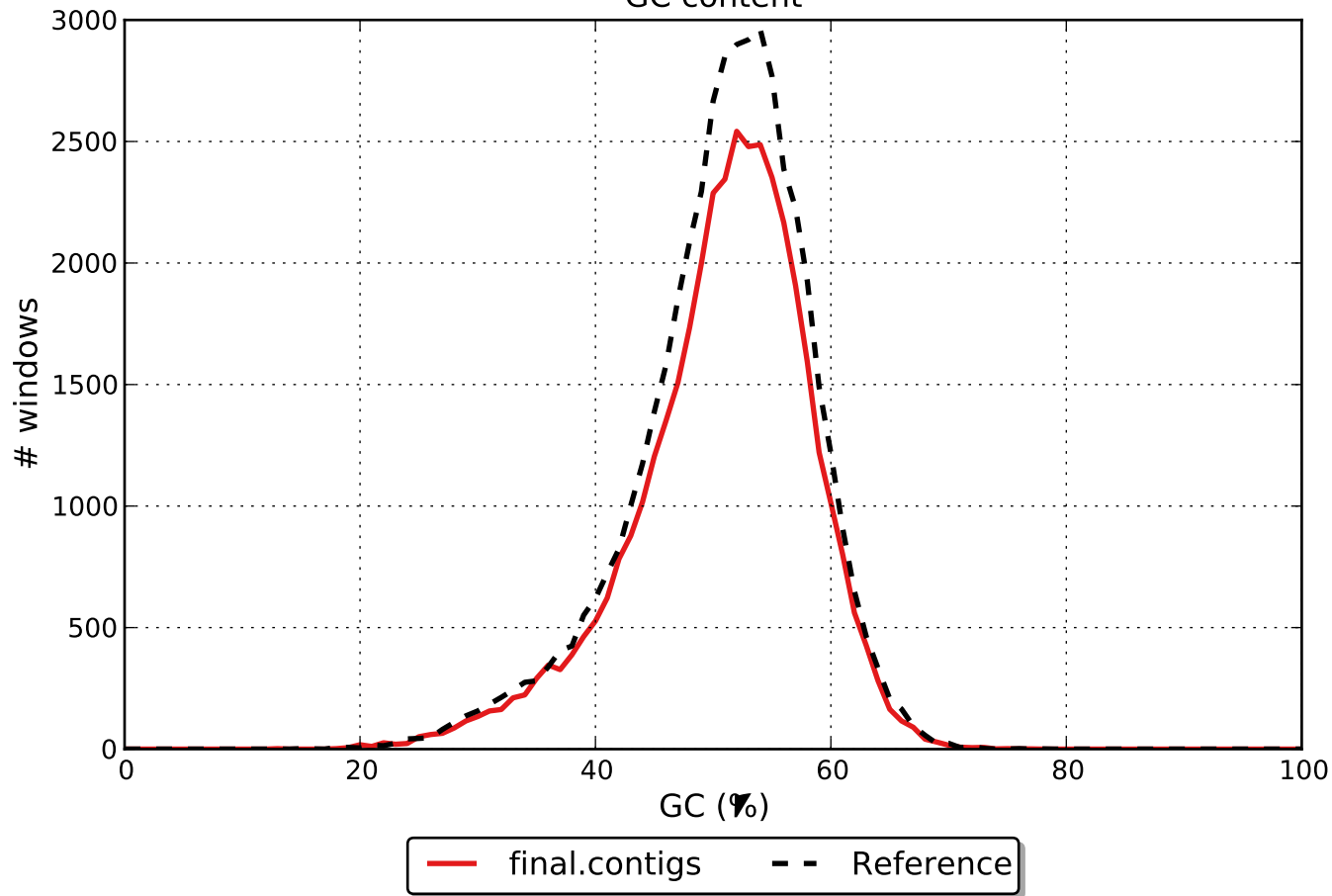
NGx



— final.contigs

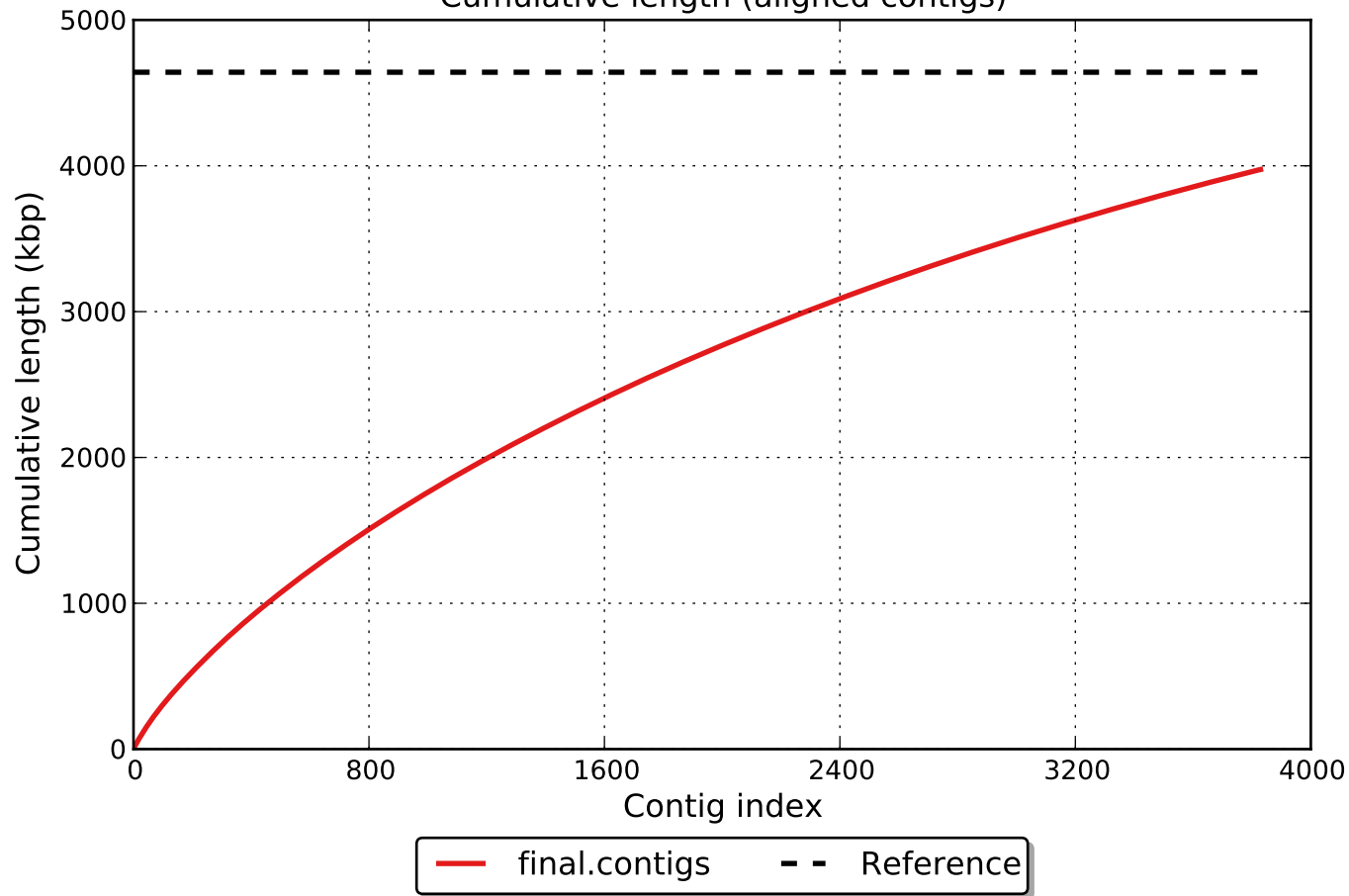


GC content

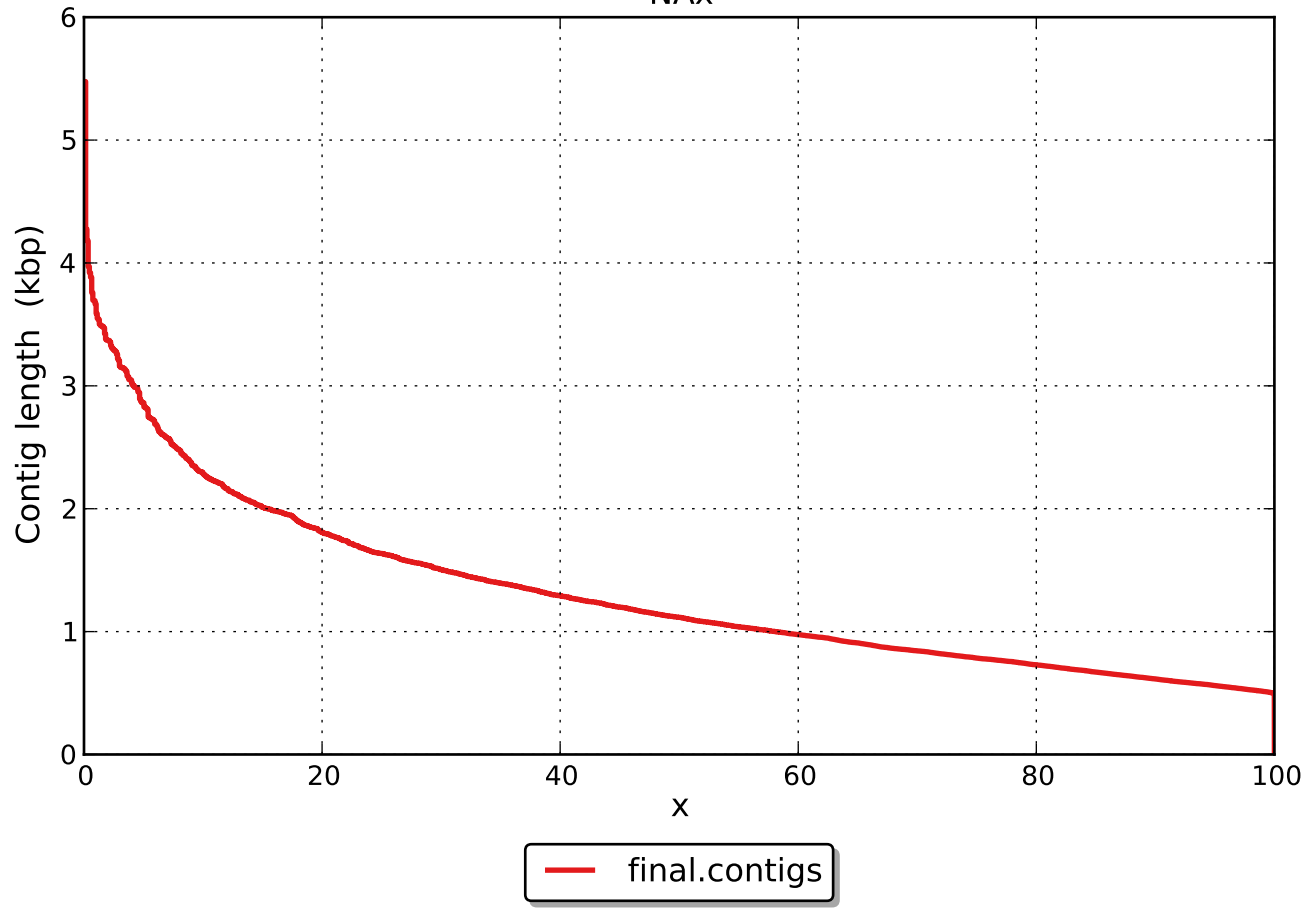




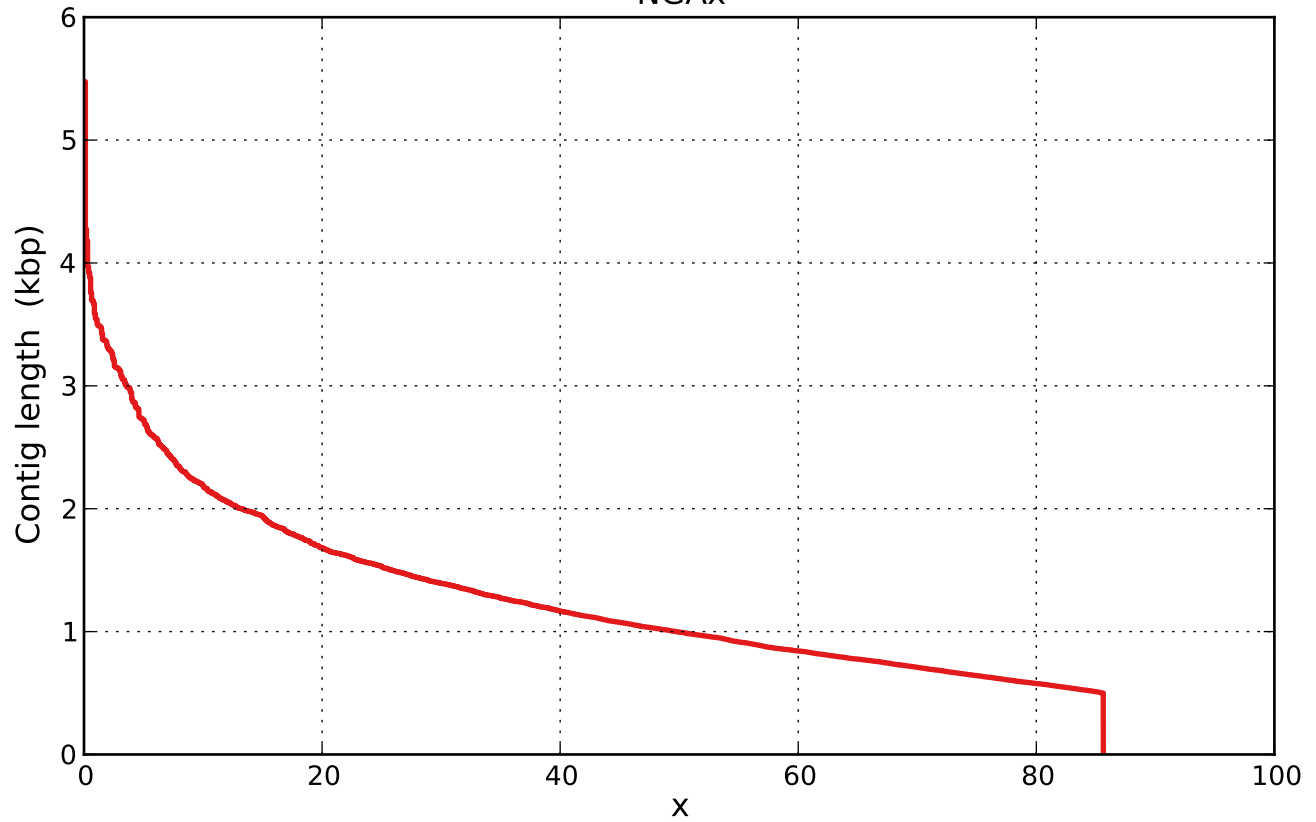
Cumulative length (aligned contigs)



NAx



NGAx



— final.contigs