

Report

	final.contigs
# contigs (>= 1000 bp)	1419
# contigs (>= 5000 bp)	255
# contigs (>= 10000 bp)	27
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 1000 bp)	4705967
Total length (>= 5000 bp)	1856419
Total length (>= 10000 bp)	344528
Total length (>= 25000 bp)	31965
Total length (>= 50000 bp)	0
# contigs	1731
Largest contig	31965
Total length	4937583
Reference length	4857432
GC (℥)	52.22
Reference GC (℥)	52.23
N50	3970
NG50	4024
N75	2327
NG75	2414
L50	393
LG50	383
L75	794
LG75	768
# misassemblies	1
# misassembled contigs	1
Misassembled contigs length	9233
# local misassemblies	0
# unaligned contigs	0 + 0 part
Unaligned length	0
Genome fraction (℥)	98.913
Duplication ratio	1.028
# N's per 100 kbp	0.00
# mismatches per 100 kbp	8.74
# indels per 100 kbp	0.04
Largest alignment	31965
NA50	3964
NGA50	4024
NA75	2327
NGA75	2411
LA50	394
LGA50	383
LA75	794
LGA75	769

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

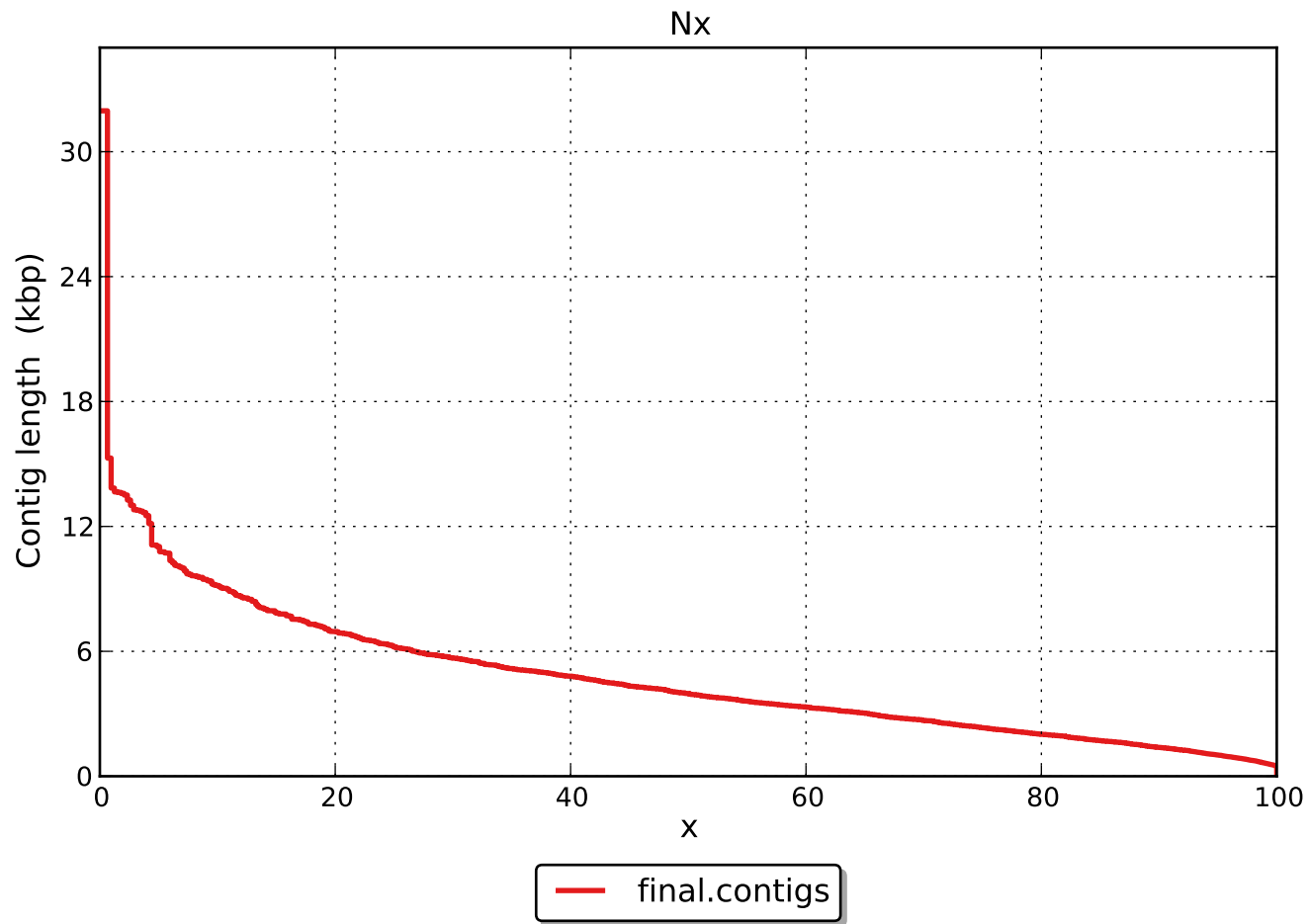
	final.contigs
# misassemblies	1
# relocations	1
# translocations	0
# inversions	0
# possibly misassembled contigs	0
# misassembled contigs	1
Misassembled contigs length	9233
# local misassemblies	0
# mismatches	420
# indels	2
# short indels	2
# long indels	0
Indels length	2

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).

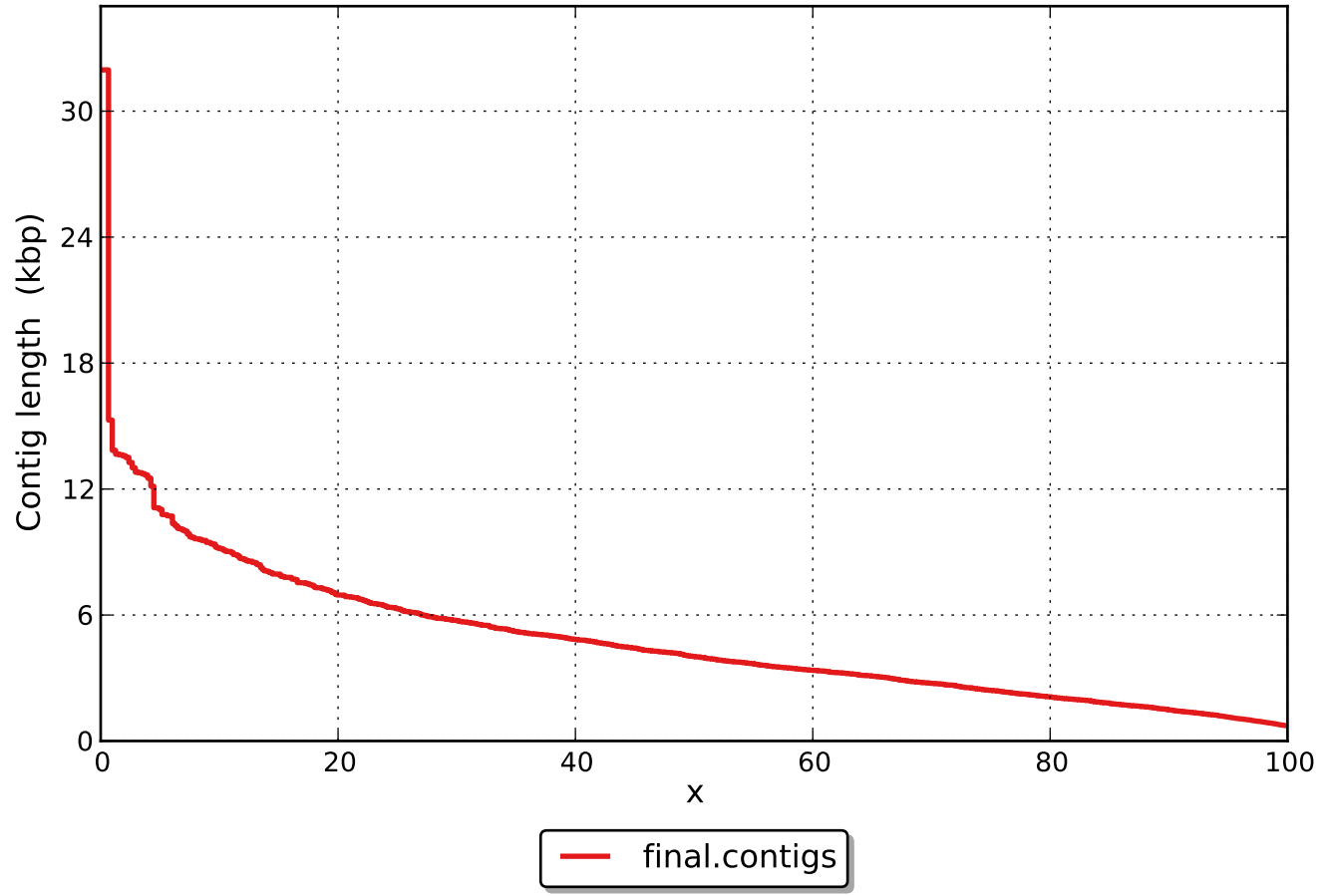
Unaligned report

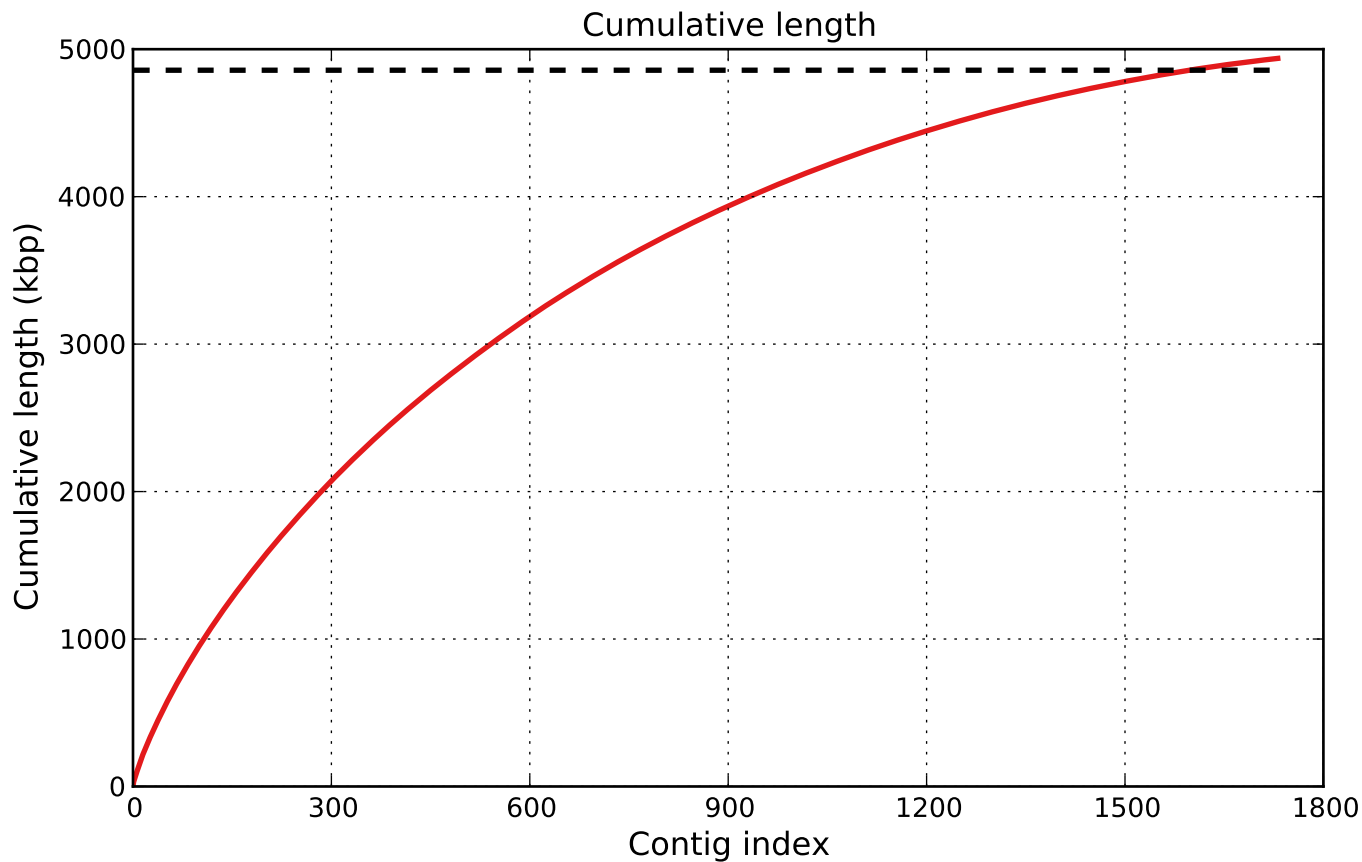
	final.contigs
# fully unaligned contigs	0
Fully unaligned length	0
# partially unaligned contigs	0
# with misassembly	0
# both parts are significant	0
Partially unaligned length	0
# N's	0

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).

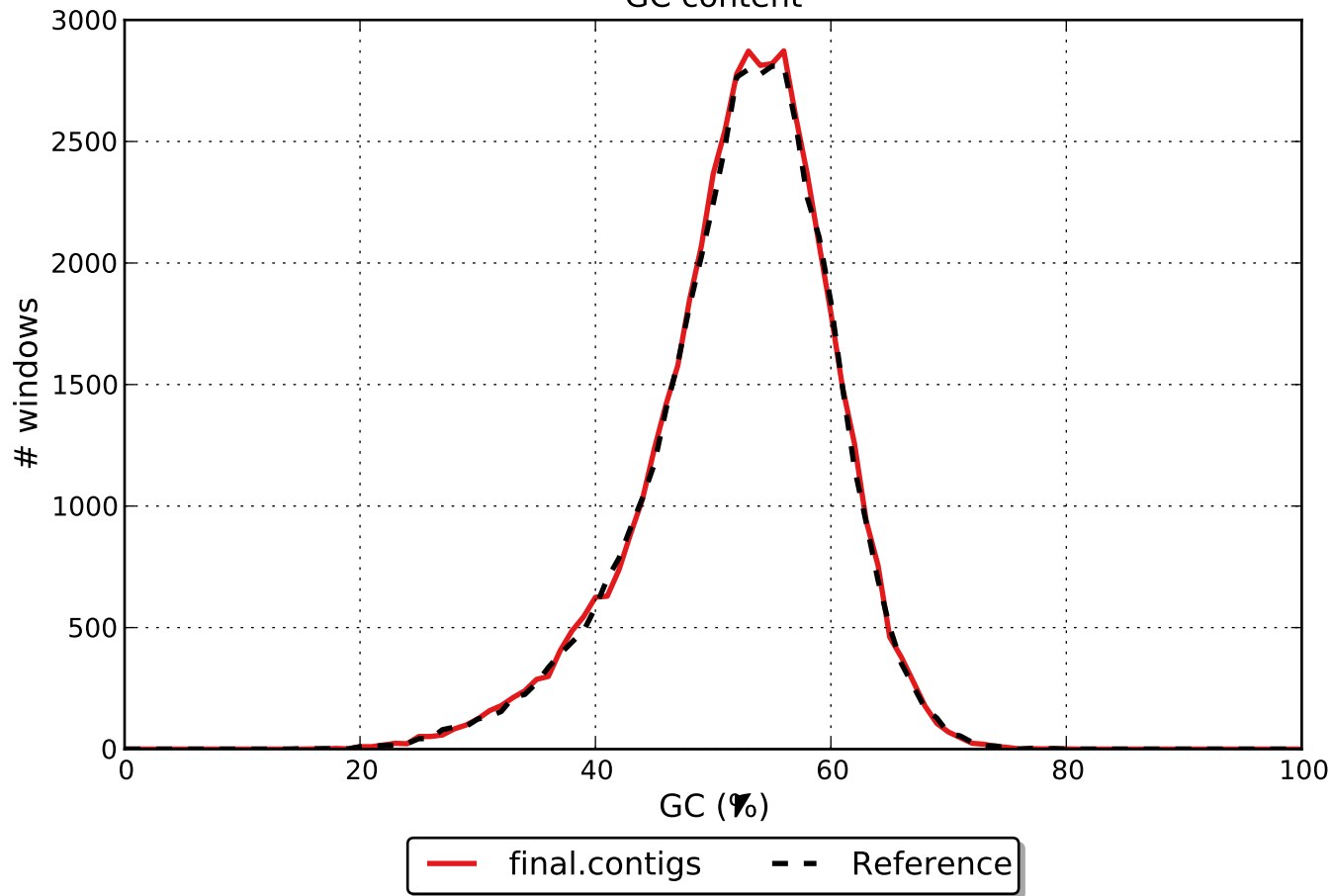


NGx



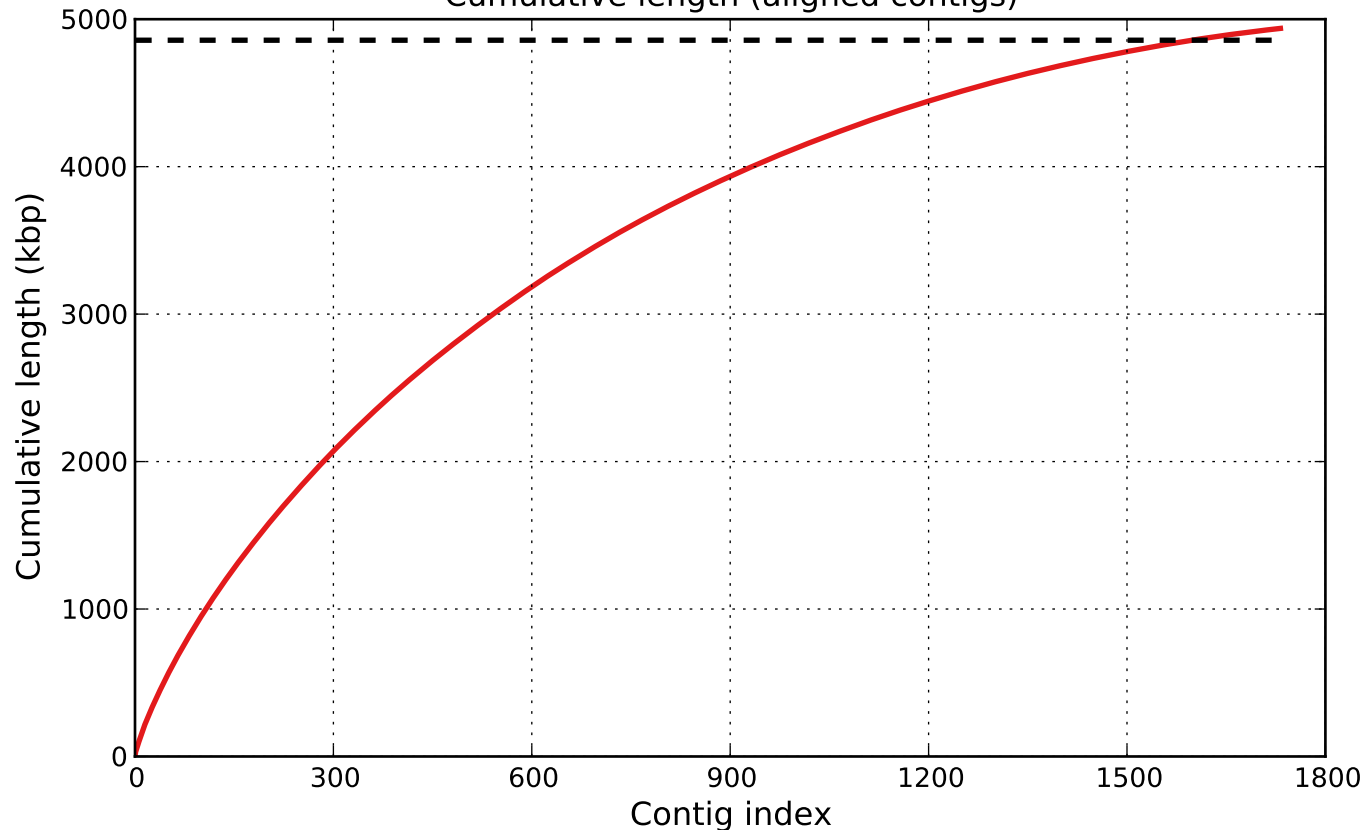


GC content

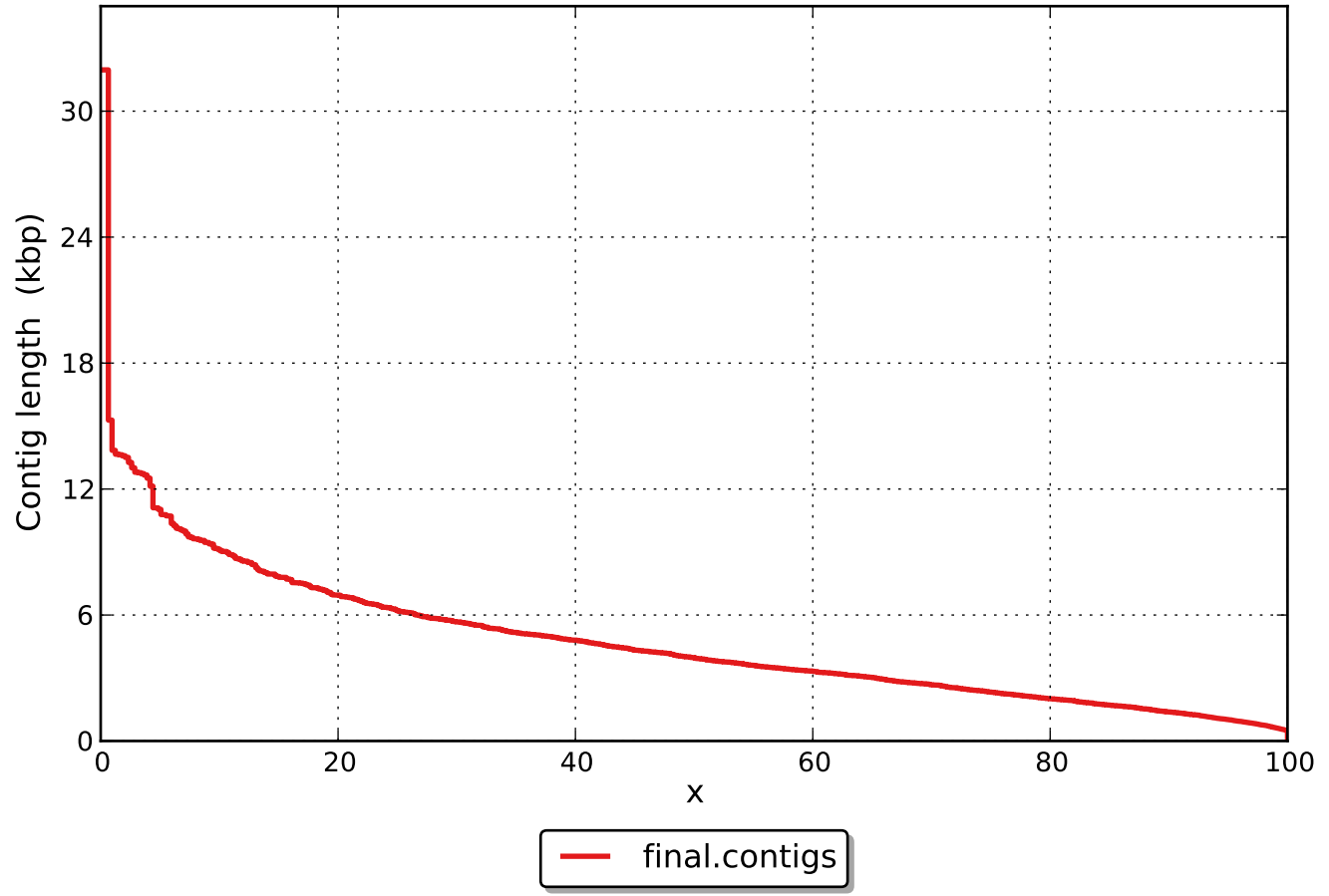




Cumulative length (aligned contigs)



NAx



NGAx

