

Report

	final.contigs
# contigs (>= 1000 bp)	718
# contigs (>= 5000 bp)	375
# contigs (>= 10000 bp)	148
# contigs (>= 25000 bp)	11
# contigs (>= 50000 bp)	0
Total length (>= 1000 bp)	4873822
Total length (>= 5000 bp)	3945475
Total length (>= 10000 bp)	2329902
Total length (>= 25000 bp)	343826
Total length (>= 50000 bp)	0
# contigs	778
Largest contig	43604
Total length	4917350
Reference length	4857432
GC (%)	52.23
Reference GC (%)	52.23
N50	9436
NG50	9585
N75	5772
NG75	5871
L50	162
LG50	159
L75	328
LG75	320
# misassemblies	0
# misassembled contigs	0
Misassembled contigs length	0
# local misassemblies	0
# unaligned contigs	0 + 0 part
Unaligned length	0
Genome fraction (%)	99.715
Duplication ratio	1.015
# N's per 100 kbp	0.00
# mismatches per 100 kbp	16.04
# indels per 100 kbp	0.00
Largest alignment	43604
NA50	9436
NGA50	9585
NA75	5772
NGA75	5871
LA50	162
LGA50	159
LA75	328
LGA75	320

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

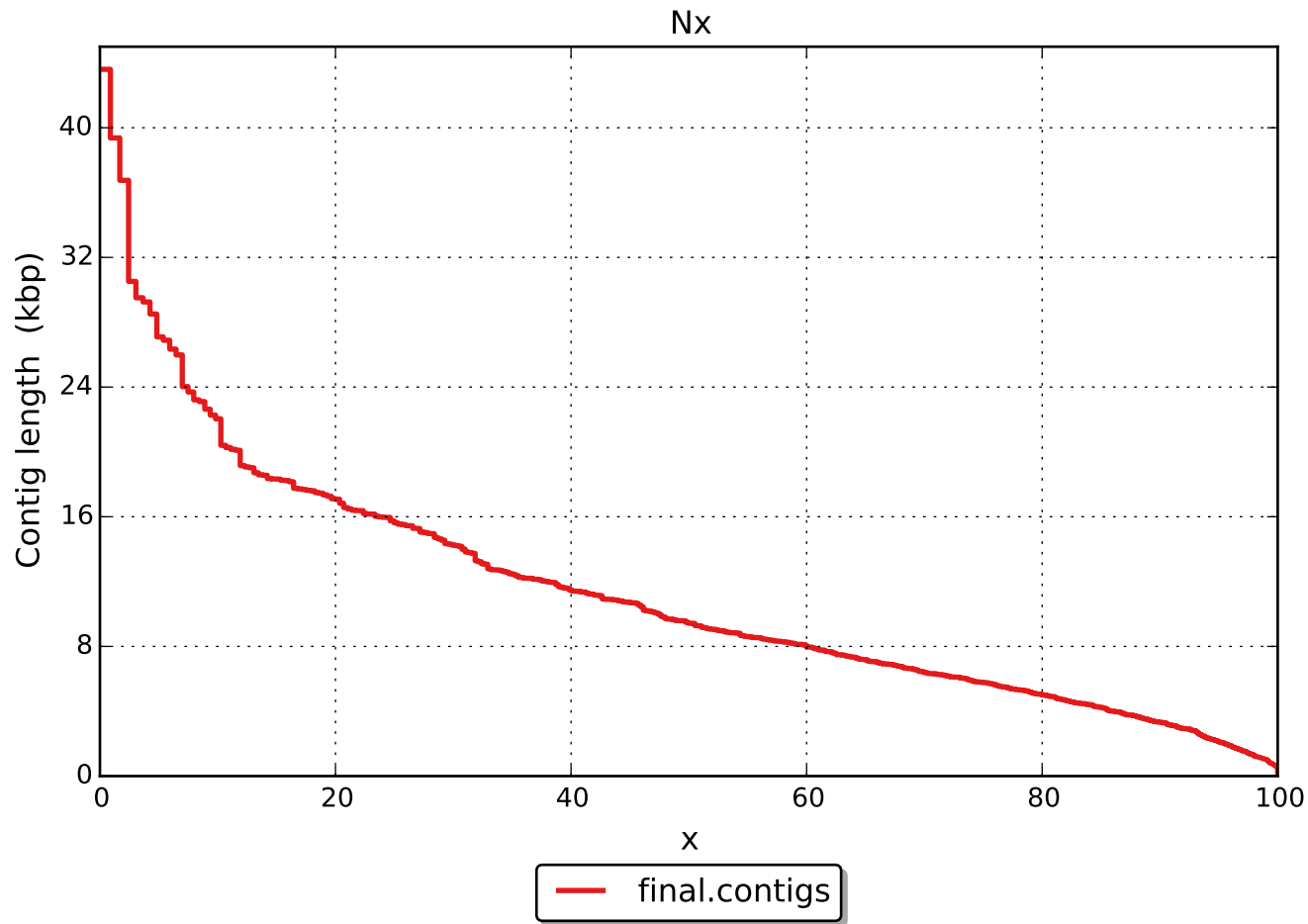
	final.contigs
# misassemblies	0
# relocations	0
# translocations	0
# inversions	0
# possibly misassembled contigs	0
# misassembled contigs	0
Misassembled contigs length	0
# local misassemblies	0
# mismatches	777
# indels	0
# short indels	0
# long indels	0
Indels length	0

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).

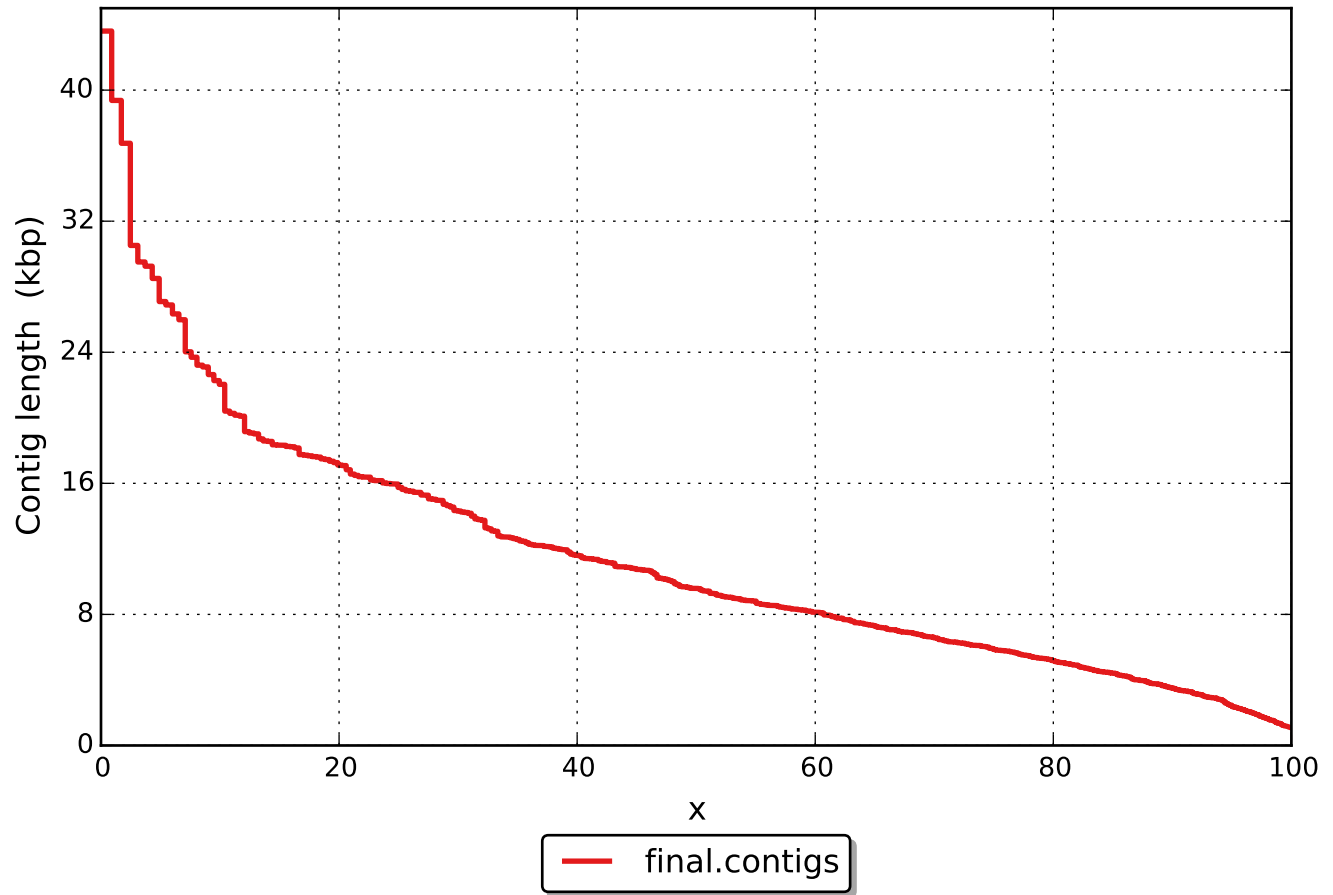
Unaligned report

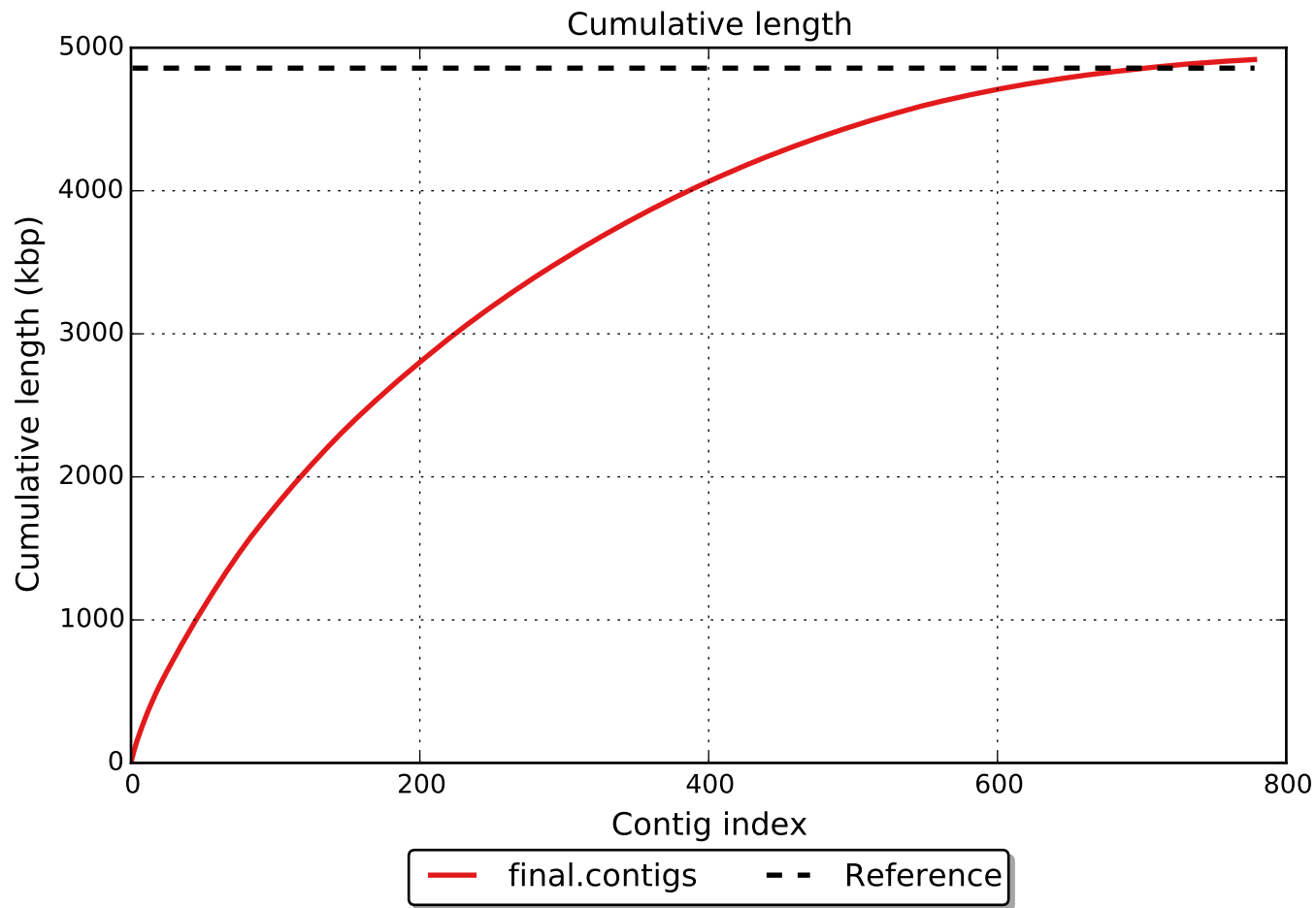
	final.contigs
# fully unaligned contigs	0
Fully unaligned length	0
# partially unaligned contigs	0
# with misassembly	0
# both parts are significant	0
Partially unaligned length	0
# N's	0

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).

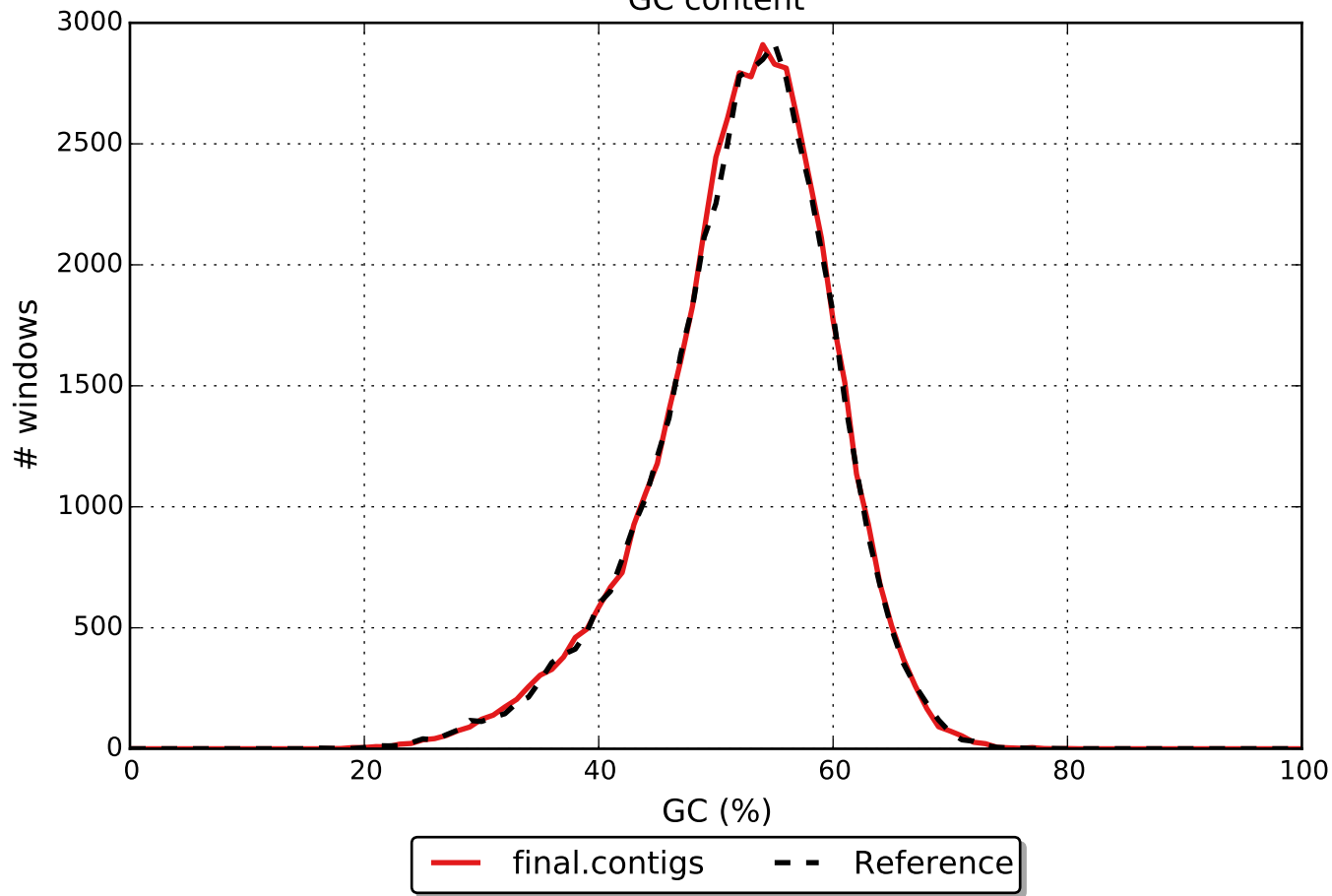


NGx





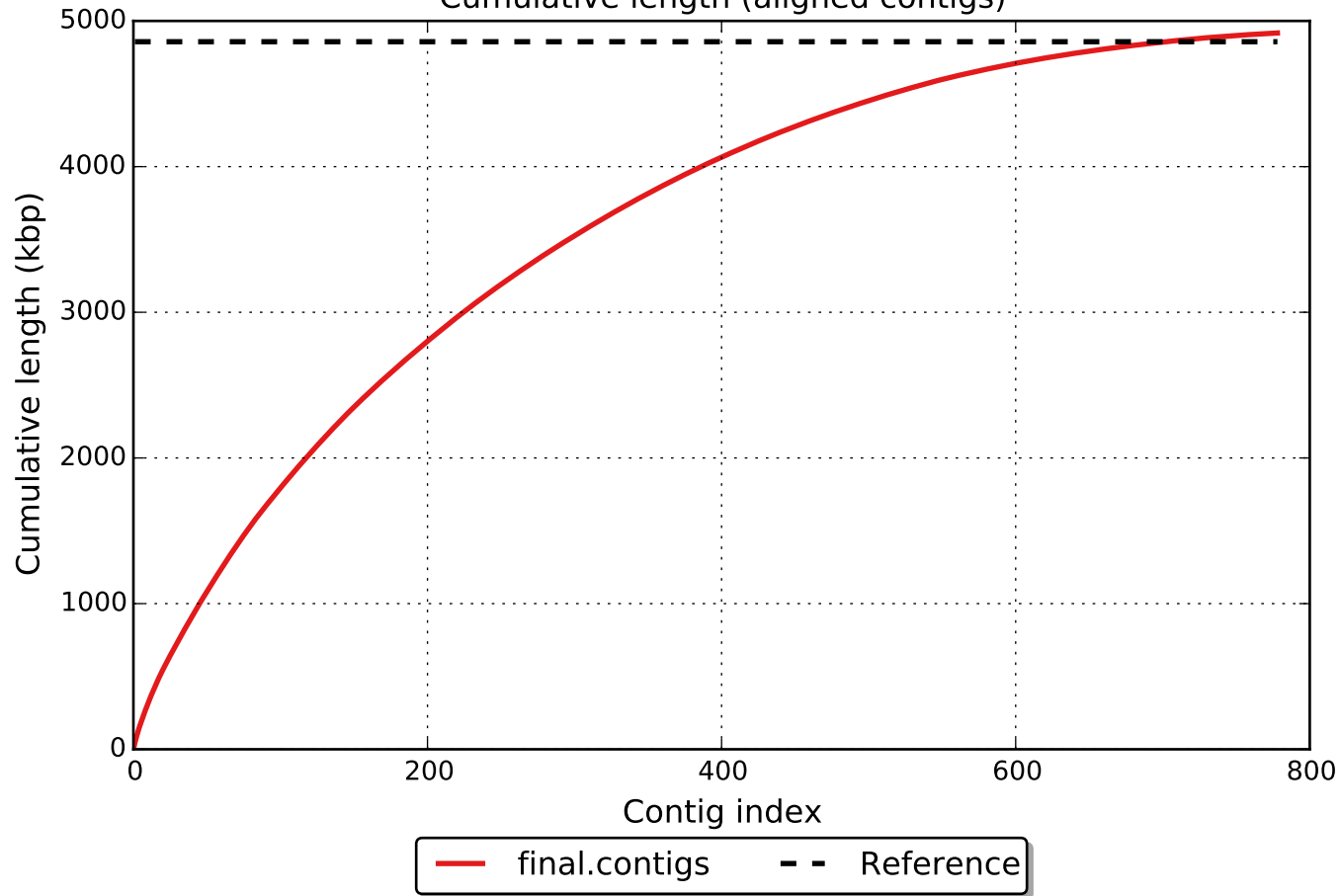
GC content



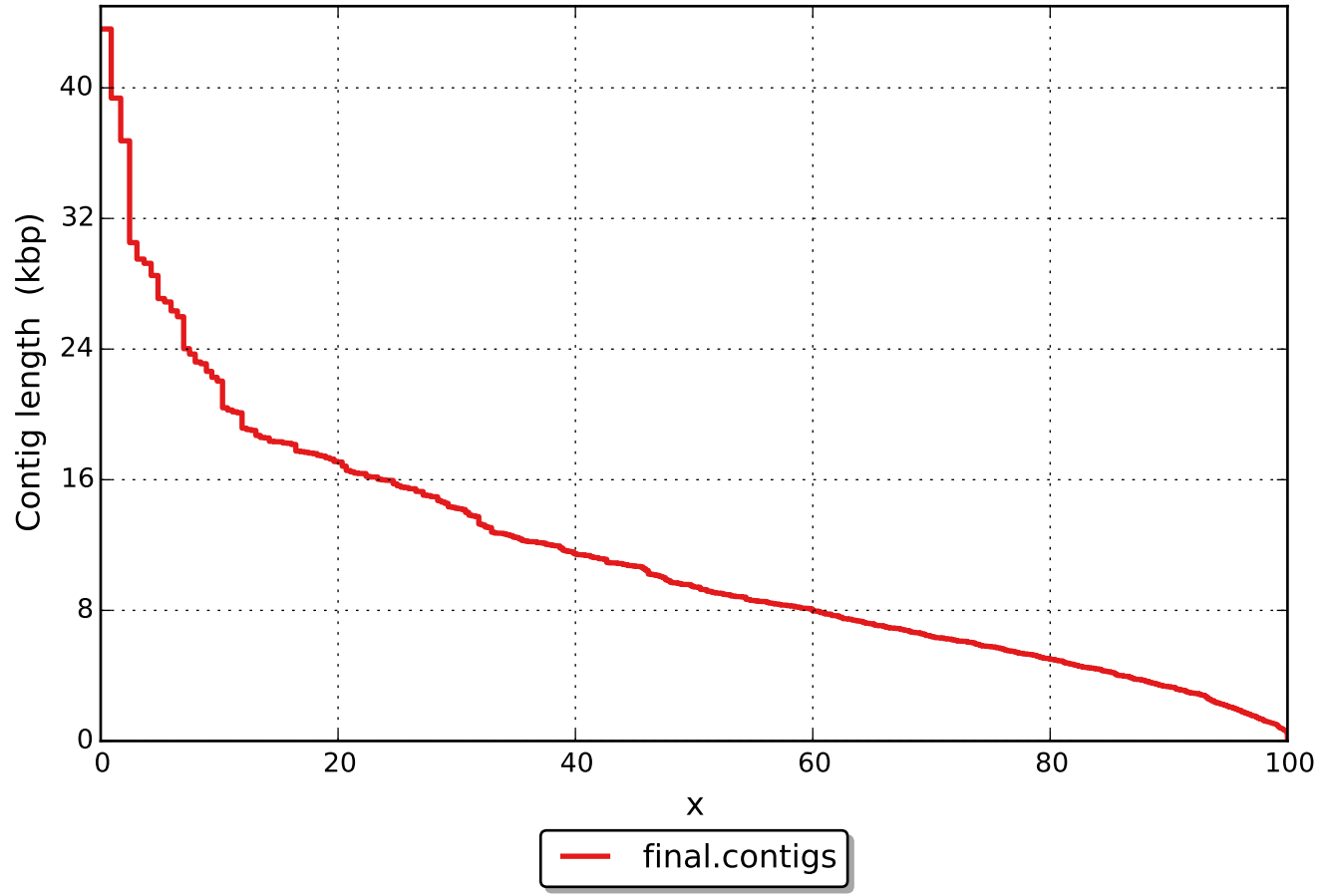
Misassemblies



Cumulative length (aligned contigs)



NAx



NGAx

