

Report

	final.contigs
# contigs (>= 1000 bp)	1699
# contigs (>= 5000 bp)	86
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 1000 bp)	3915920
Total length (>= 5000 bp)	533575
Total length (>= 10000 bp)	21899
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	2557
Largest contig	11062
Total length	4549027
Reference length	4641652
GC (℥)	50.73
Reference GC (℥)	50.79
N50	2267
NG50	2235
N75	1389
NG75	1333
L50	635
LG50	656
L75	1275
LG75	1326
# misassemblies	0
# misassembled contigs	0
Misassembled contigs length	0
# local misassemblies	0
# unaligned contigs	0 + 0 part
Unaligned length	0
Genome fraction (℥)	94.823
Duplication ratio	1.034
# N's per 100 kbp	0.00
# mismatches per 100 kbp	5.13
# indels per 100 kbp	0.00
Largest alignment	11062
NA50	2267
NGA50	2235
NA75	1389
NGA75	1333
LA50	635
LGA50	656
LA75	1275
LGA75	1326

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

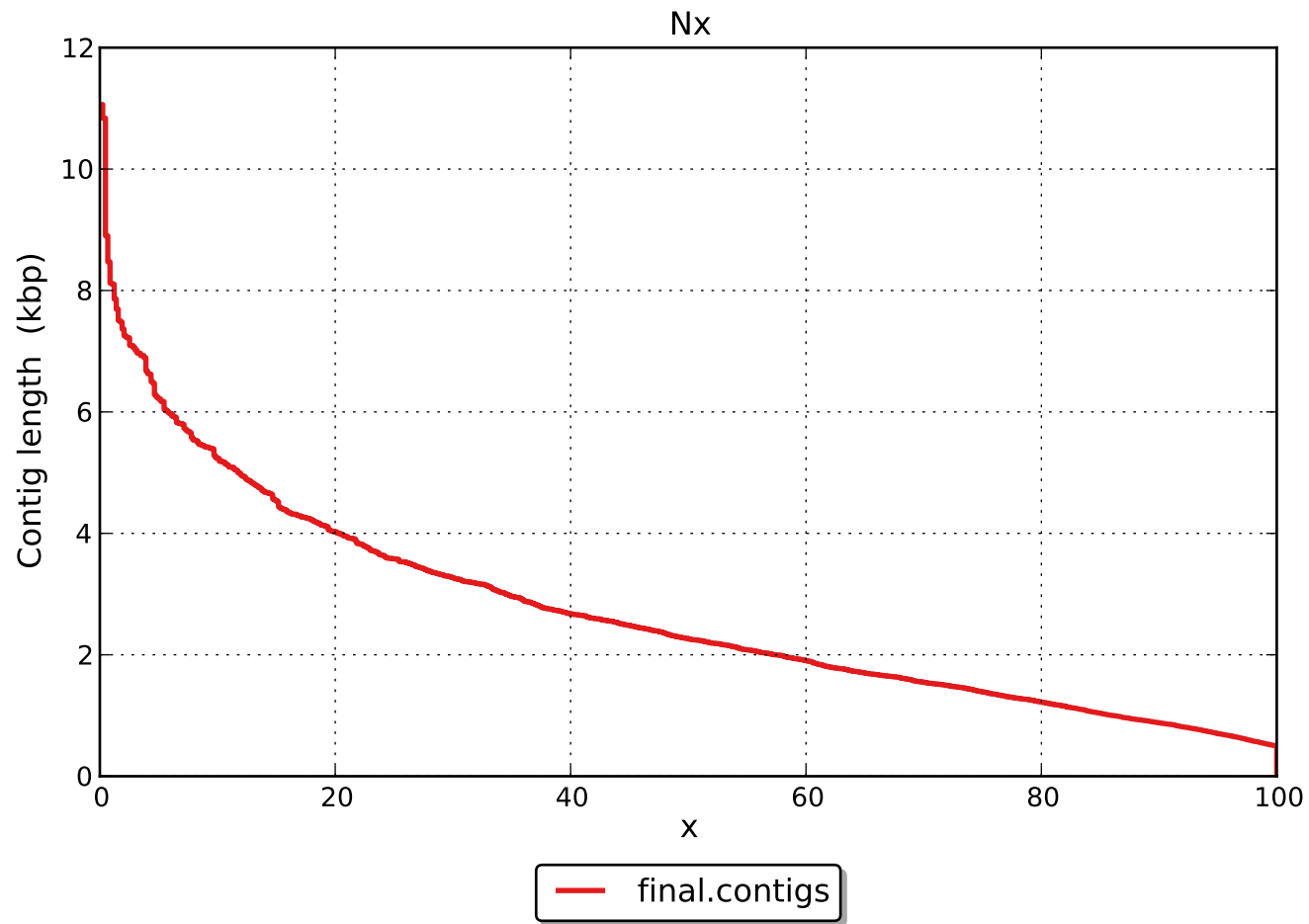
	final.contigs
# misassemblies	0
# relocations	0
# translocations	0
# inversions	0
# possibly misassembled contigs	0
# misassembled contigs	0
Misassembled contigs length	0
# local misassemblies	0
# mismatches	226
# indels	0
# short indels	0
# long indels	0
Indels length	0

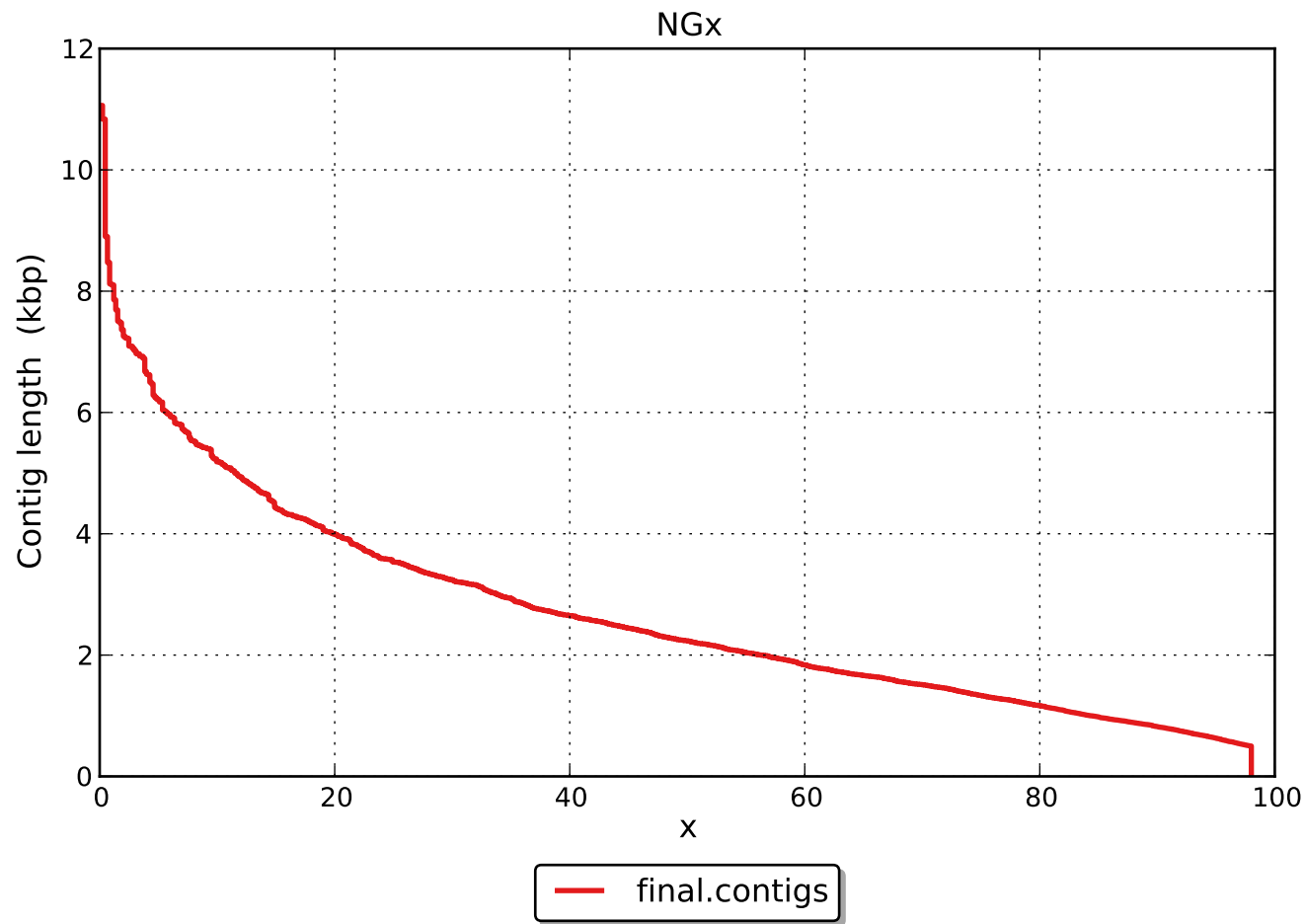
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

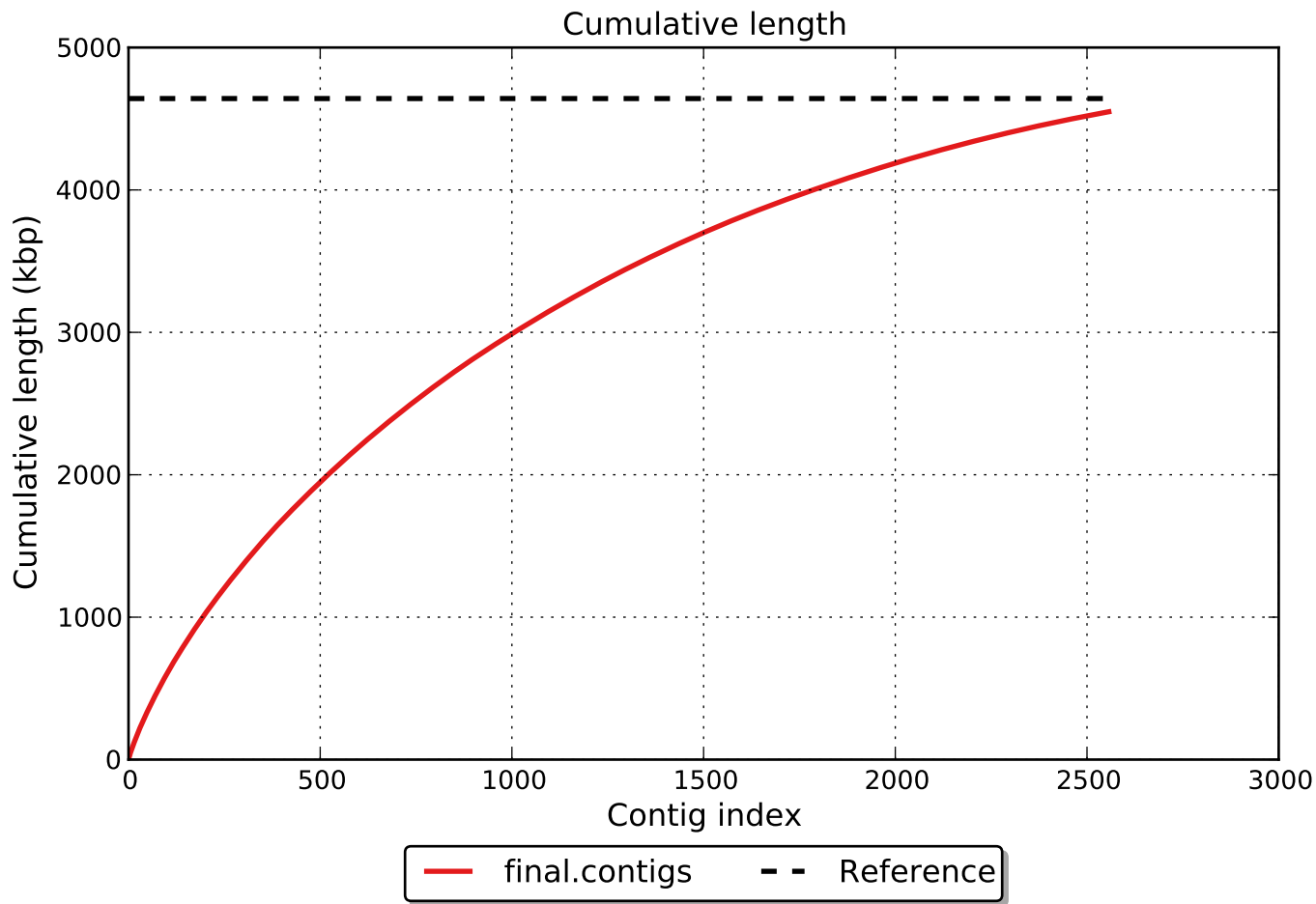
Unaligned report

	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

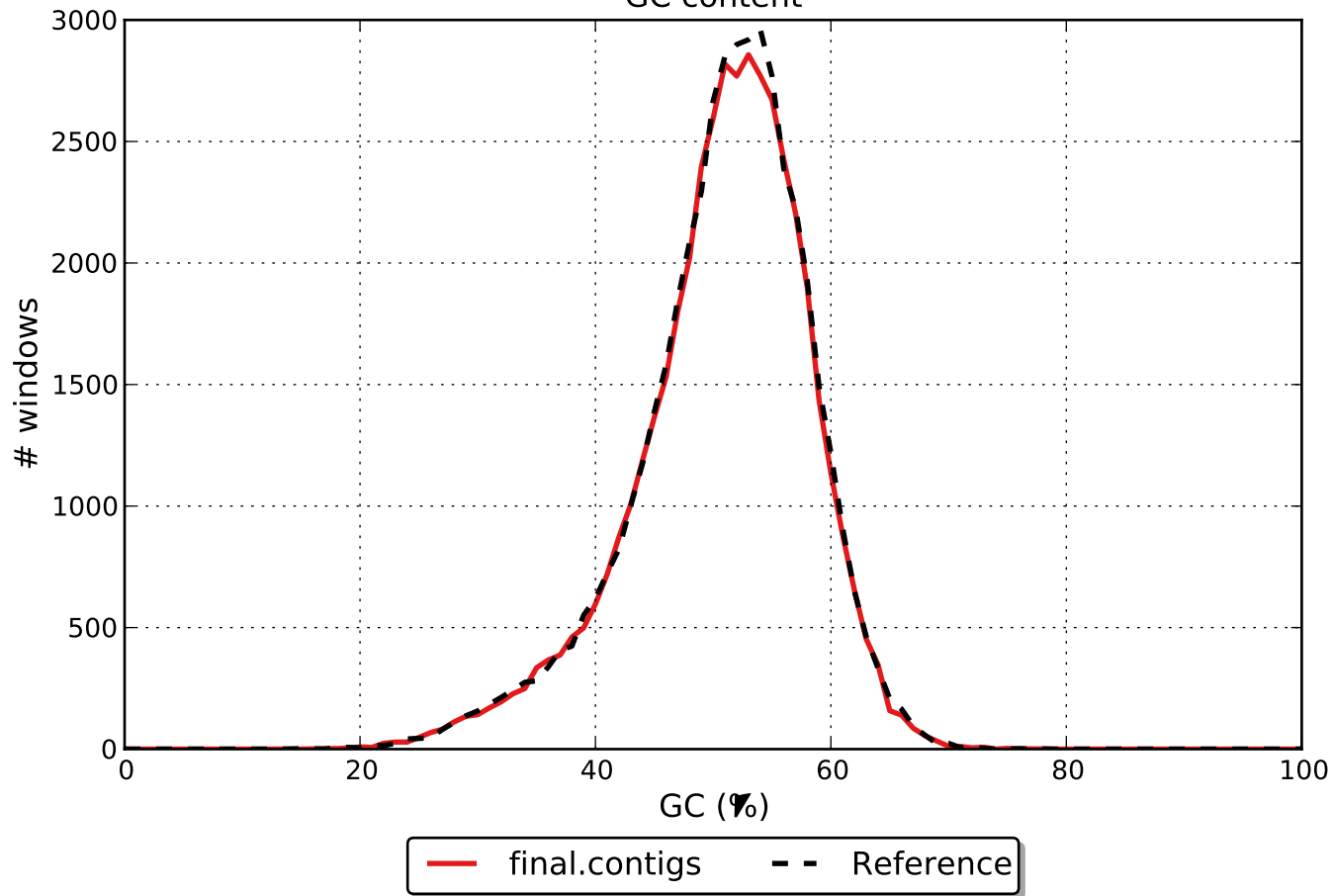
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).







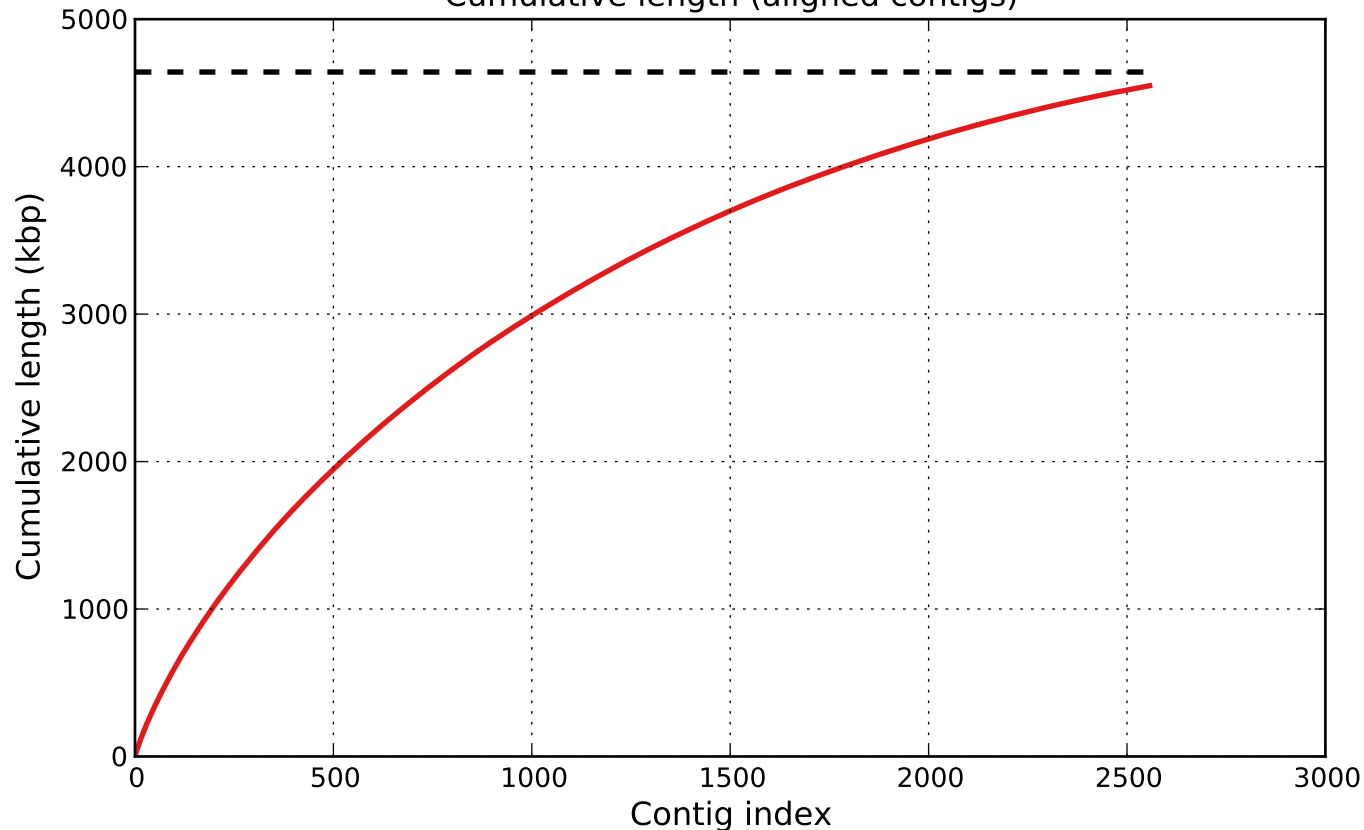
GC content



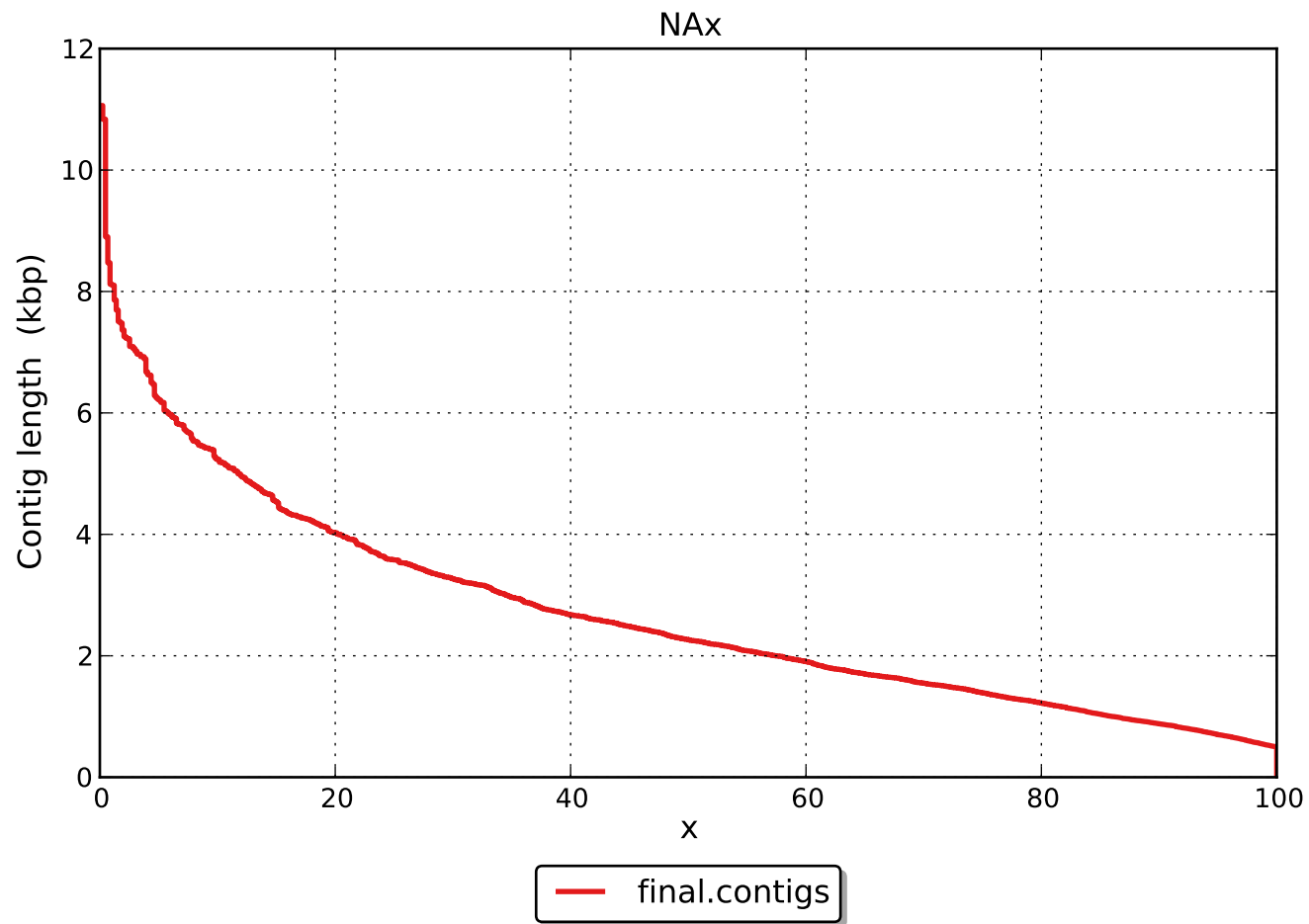
Misassemblies



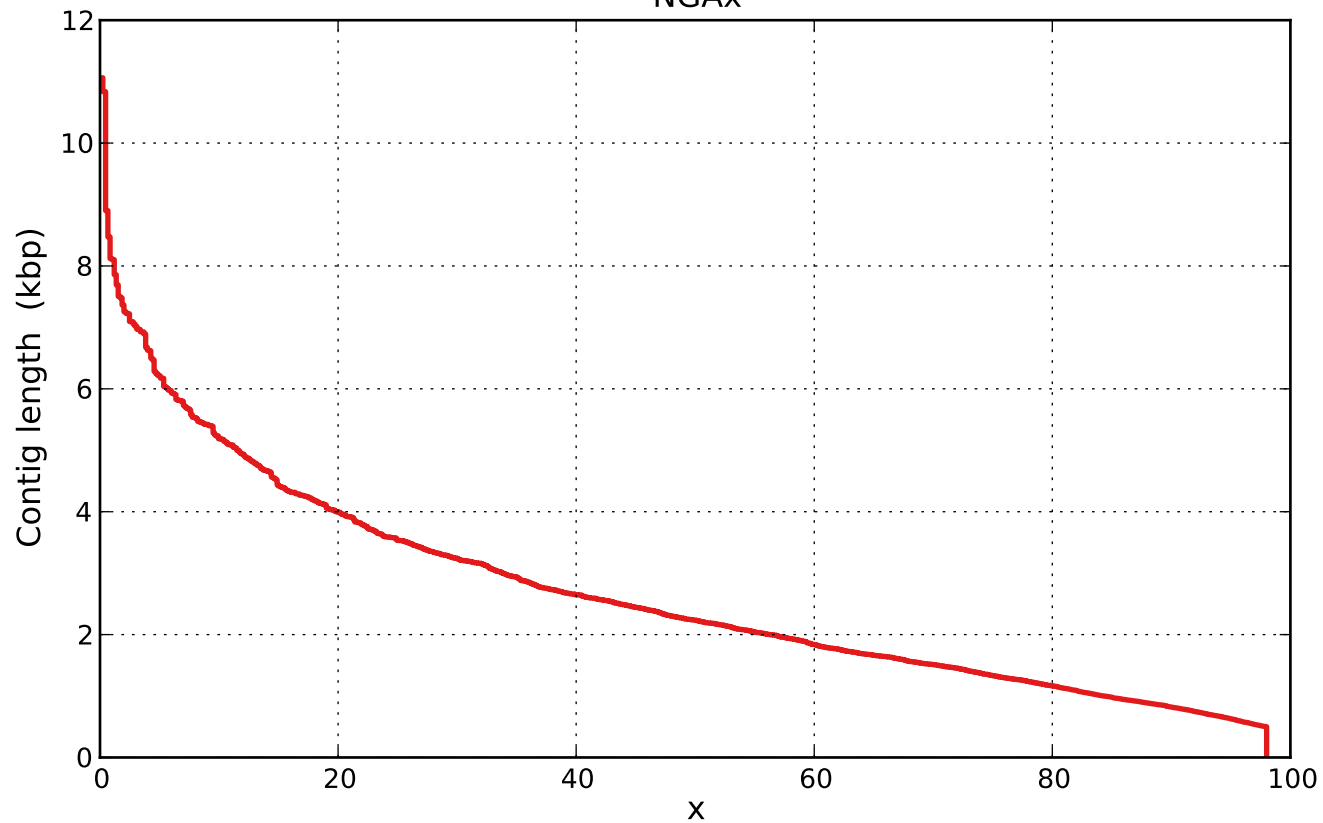
Cumulative length (aligned contigs)



— final.contigs - - Reference



NGAx



— final.contigs