

Report

	final.contigs
# contigs (>= 1000 bp)	1829
# contigs (>= 5000 bp)	71
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 1000 bp)	4124239
Total length (>= 5000 bp)	425185
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	2714
Largest contig	9591
Total length	4782387
Reference length	4857432
GC (℥)	52.19
Reference GC (℥)	52.22
N50	2241
NG50	2213
N75	1373
NG75	1327
L50	692
LG50	709
L75	1370
LG75	1412
# misassemblies	0
# misassembled contigs	0
Misassembled contigs length	0
# local misassemblies	0
# unaligned contigs	0 + 0 part
Unaligned length	0
Genome fraction (℥)	95.216
Duplication ratio	1.034
# N's per 100 kbp	0.00
# mismatches per 100 kbp	4.30
# indels per 100 kbp	0.00
Largest alignment	9591
NA50	2241
NGA50	2213
NA75	1373
NGA75	1327
LA50	692
LGA50	709
LA75	1370
LGA75	1412

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

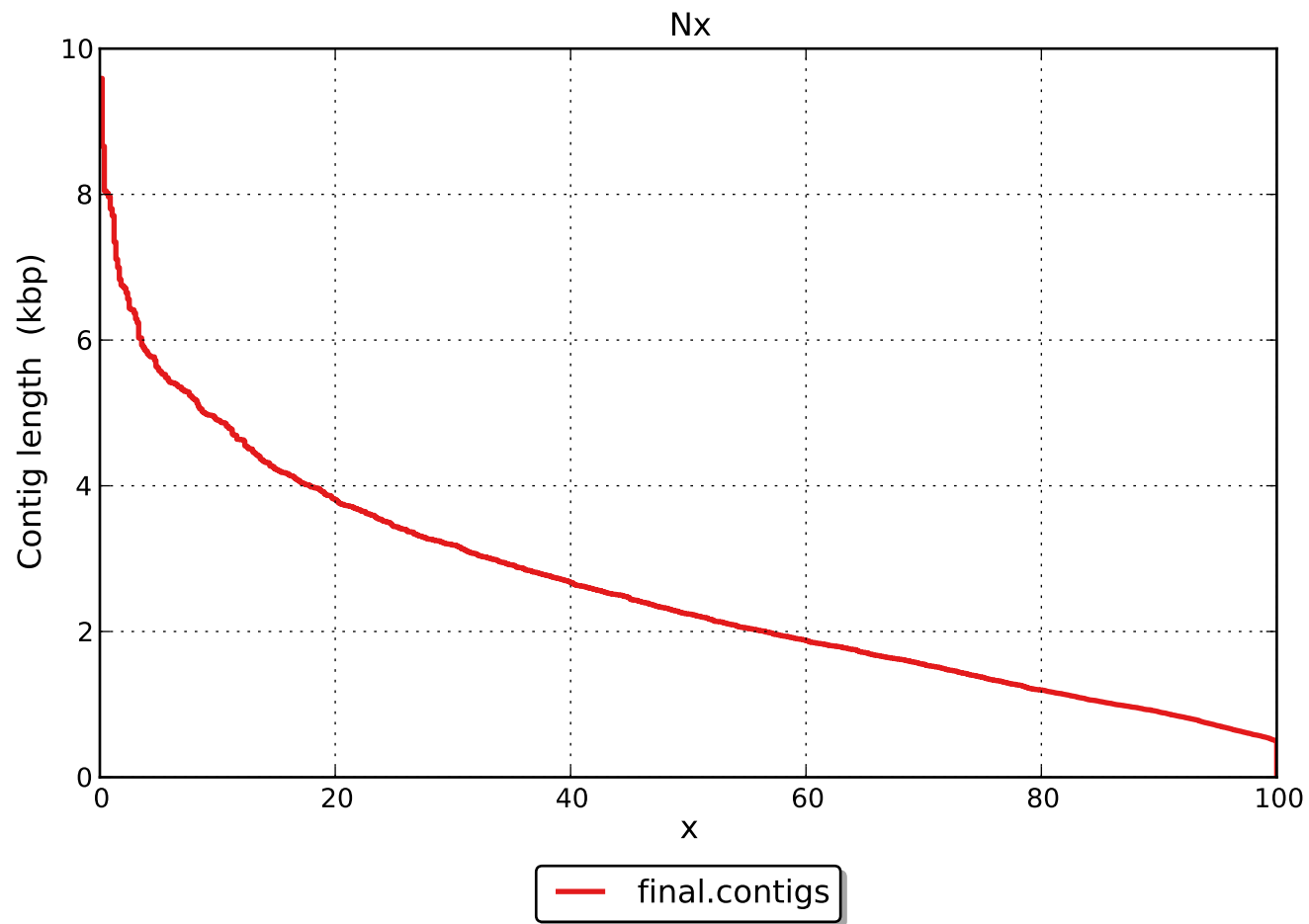
	final.contigs
# misassemblies	0
# relocations	0
# translocations	0
# inversions	0
# possibly misassembled contigs	0
# misassembled contigs	0
Misassembled contigs length	0
# local misassemblies	0
# mismatches	199
# indels	0
# short indels	0
# long indels	0
Indels length	0

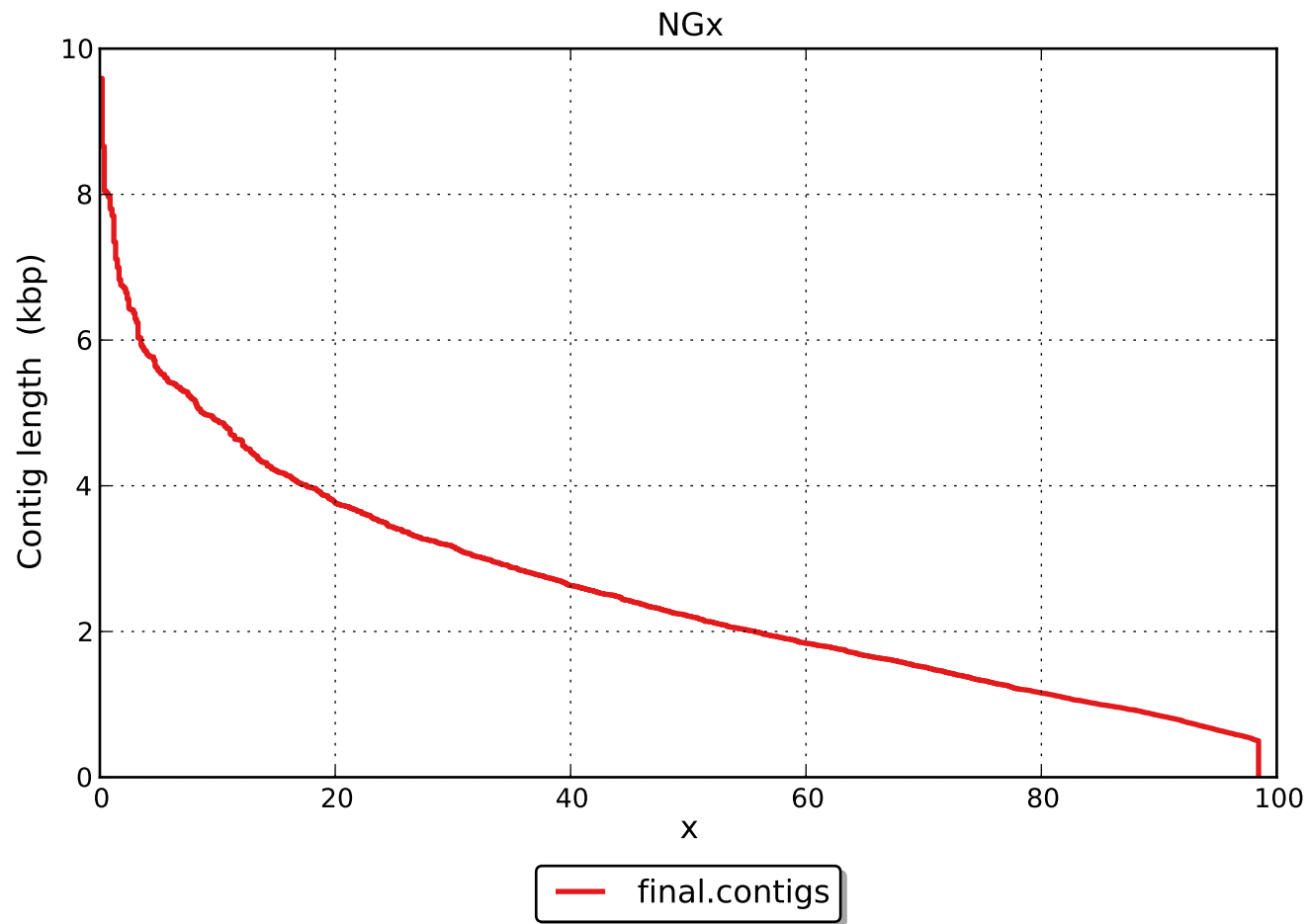
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

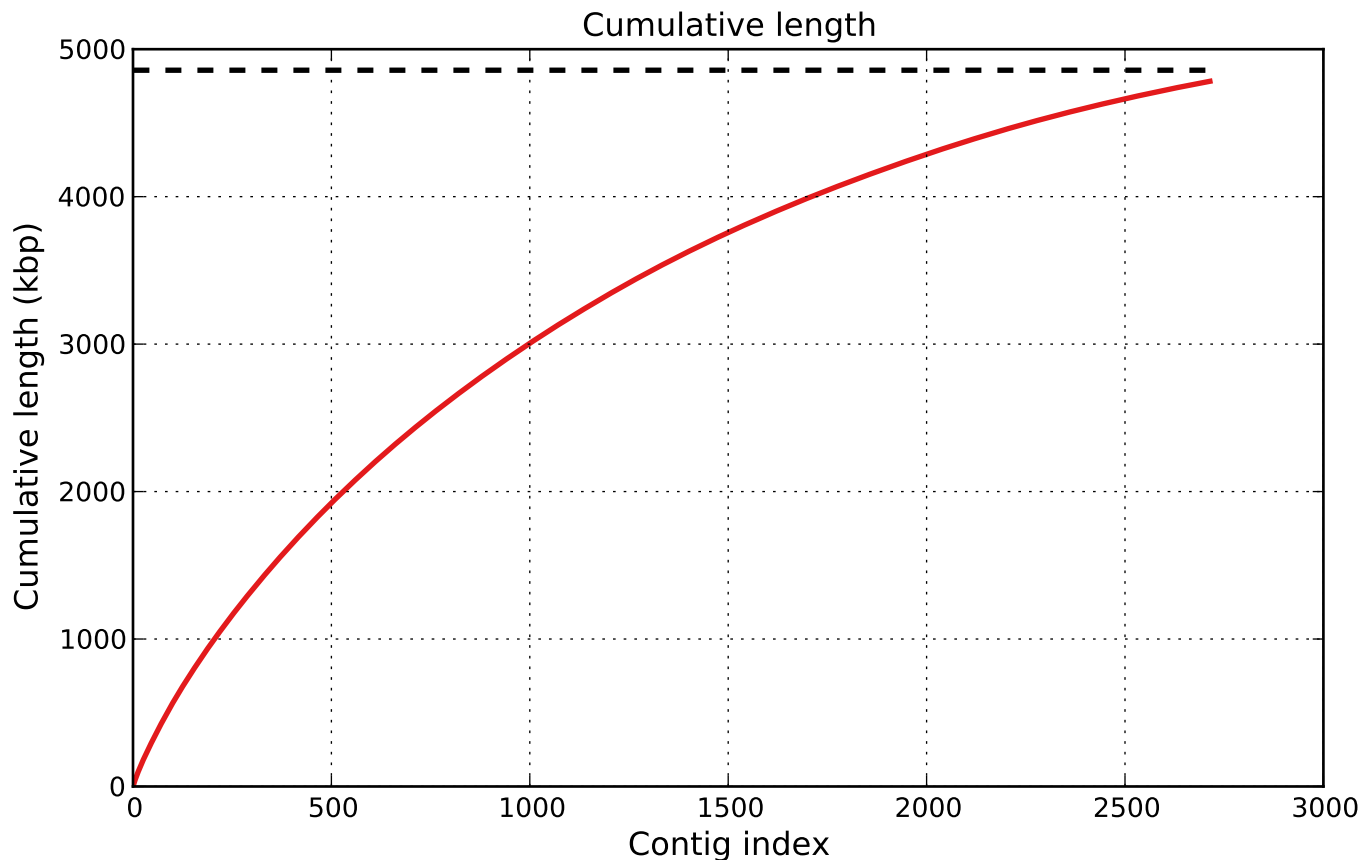
Unaligned report

	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

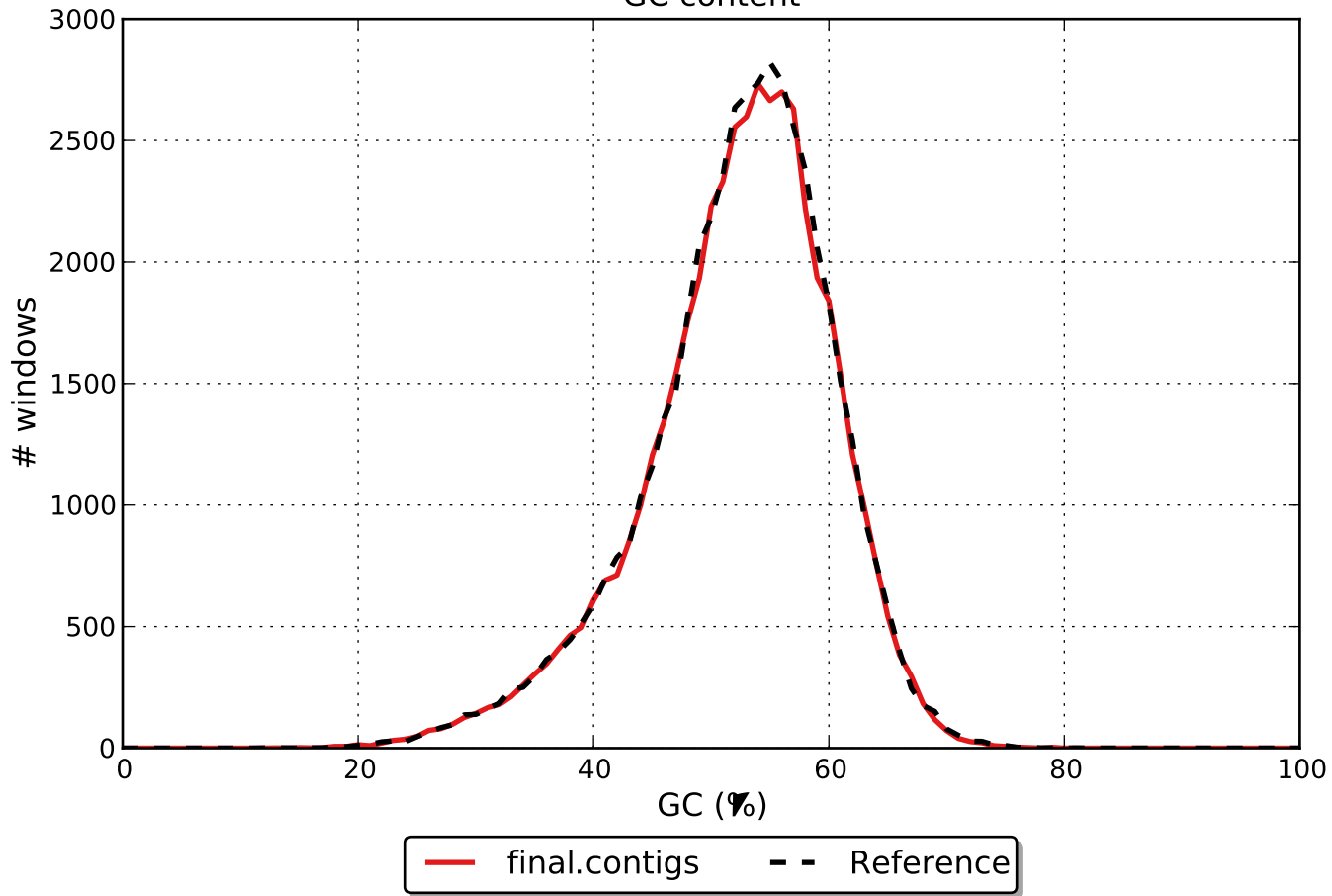
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).







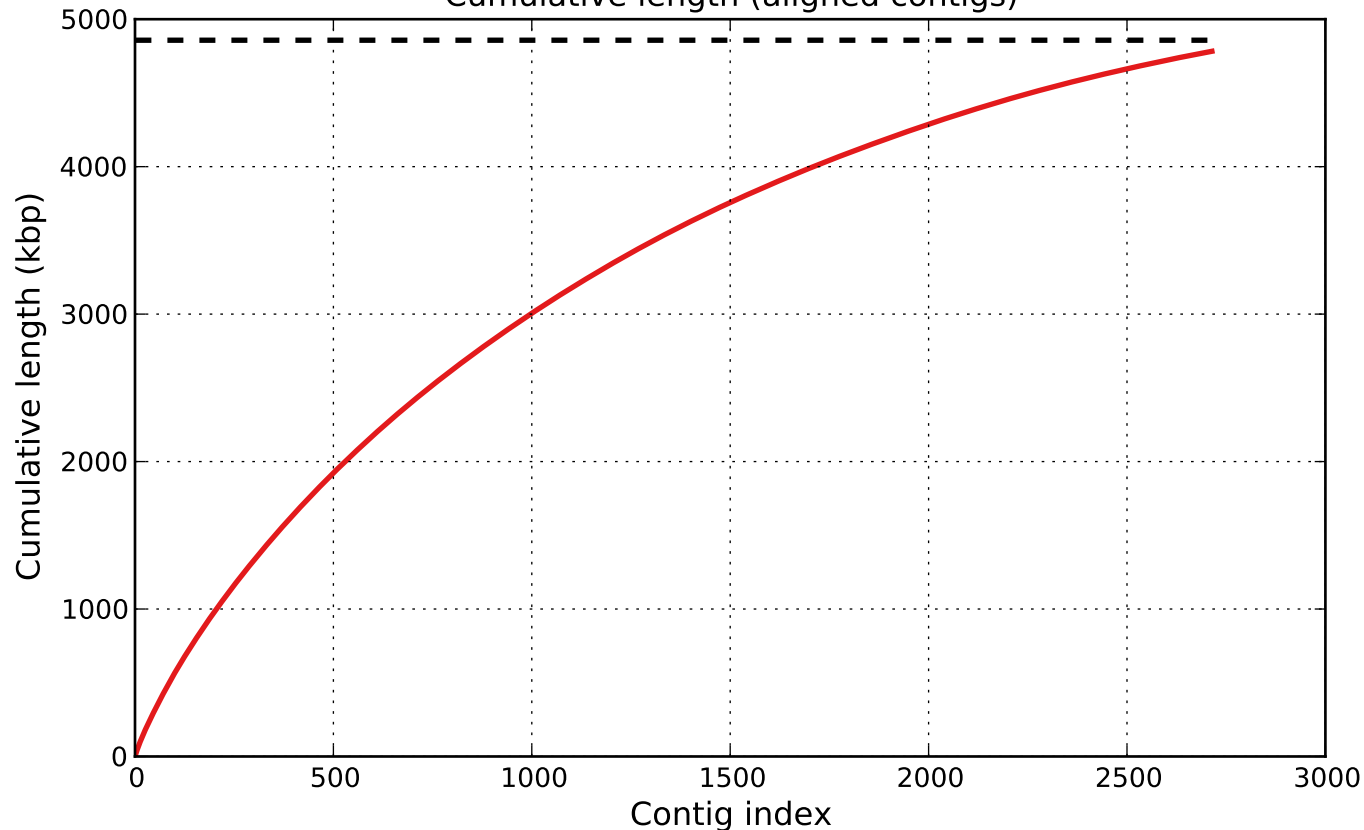
GC content



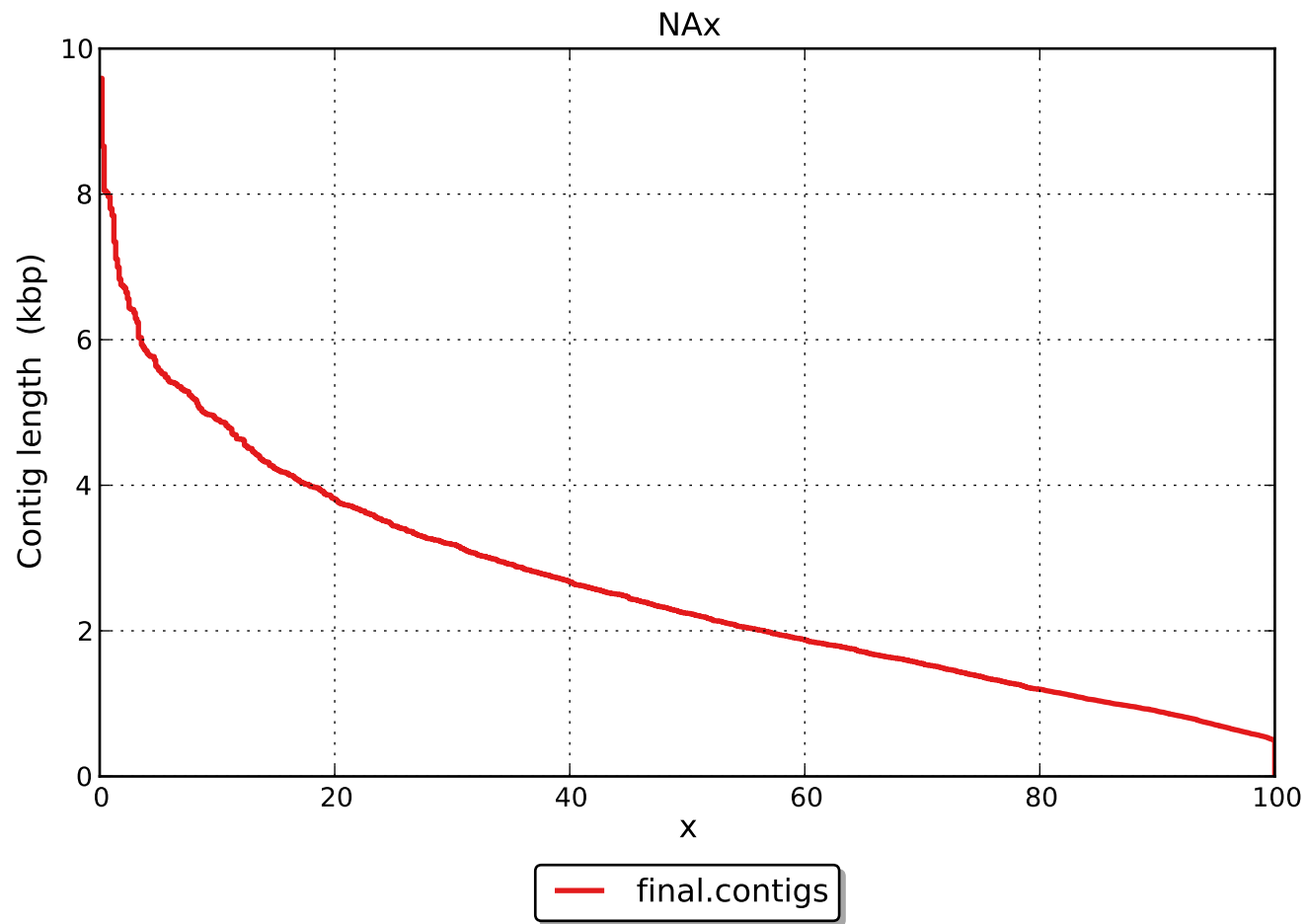
Misassemblies



Cumulative length (aligned contigs)



— final.contigs - - Reference



NGAx

