

Report

	final.contigs
# contigs (>= 1000 bp)	1073
# contigs (>= 5000 bp)	347
# contigs (>= 10000 bp)	68
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 1000 bp)	4592383
Total length (>= 5000 bp)	2725285
Total length (>= 10000 bp)	868571
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	1254
Largest contig	19288
Total length	4725872
Reference length	4641652
GC (%)	50.78
Reference GC (%)	50.78
N50	5607
NG50	5669
N75	3257
NG75	3391
L50	279
LG50	272
L75	551
LG75	532
# misassemblies	0
# misassembled contigs	0
Misassembled contigs length	0
# local misassemblies	0
# unaligned contigs	0 + 1 part
Unaligned length	29
Genome fraction (%)	99.354
Duplication ratio	1.025
# N's per 100 kbp	0.00
# mismatches per 100 kbp	27.67
# indels per 100 kbp	0.00
Largest alignment	19288
NA50	5607
NGA50	5669
NA75	3257
NGA75	3391
LA50	279
LGA50	272
LA75	551
LGA75	532

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

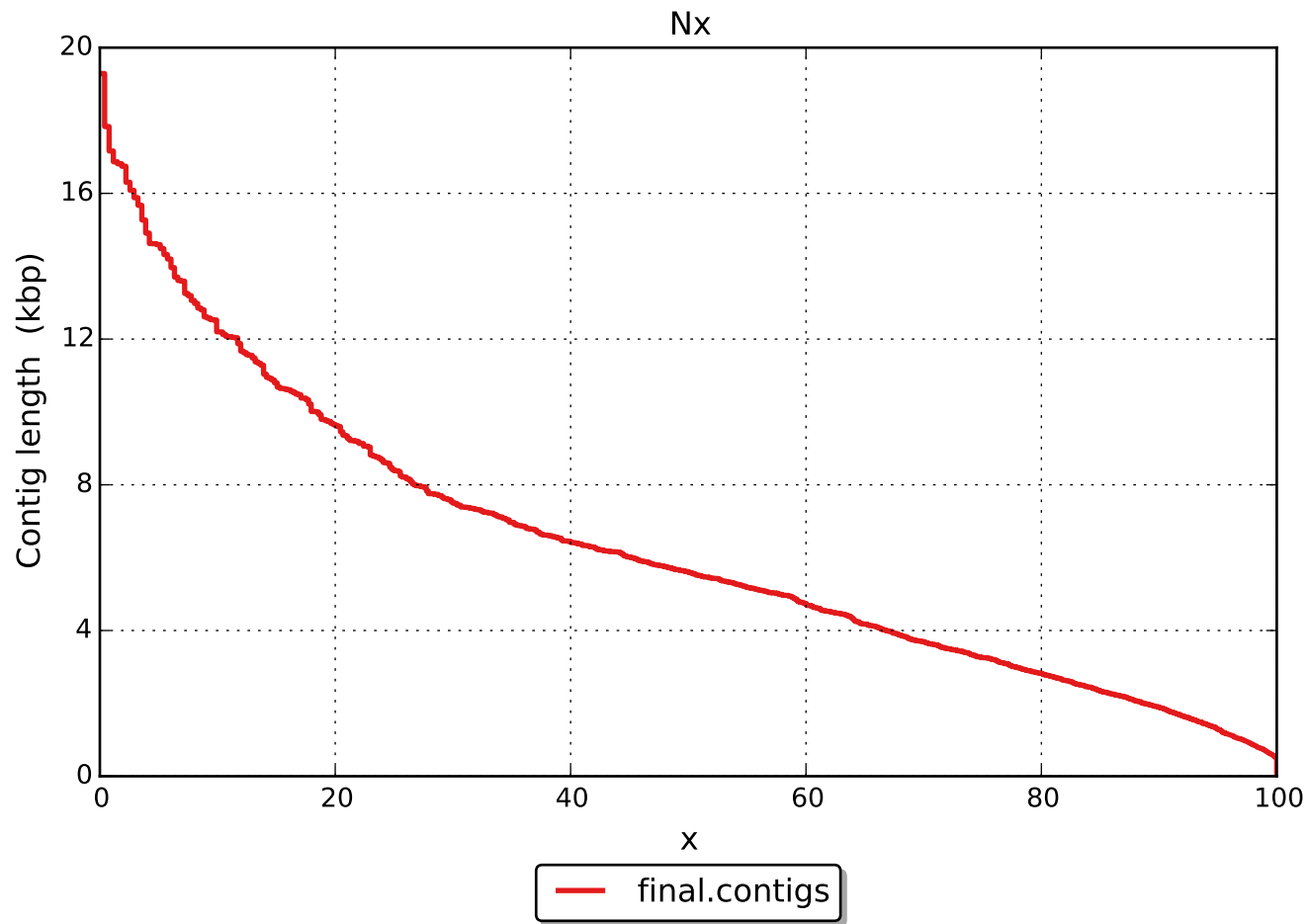
	final.contigs
# misassemblies	0
# relocations	0
# translocations	0
# inversions	0
# possibly misassembled contigs	0
# misassembled contigs	0
Misassembled contigs length	0
# local misassemblies	0
# mismatches	1276
# indels	0
# short indels	0
# long indels	0
Indels length	0

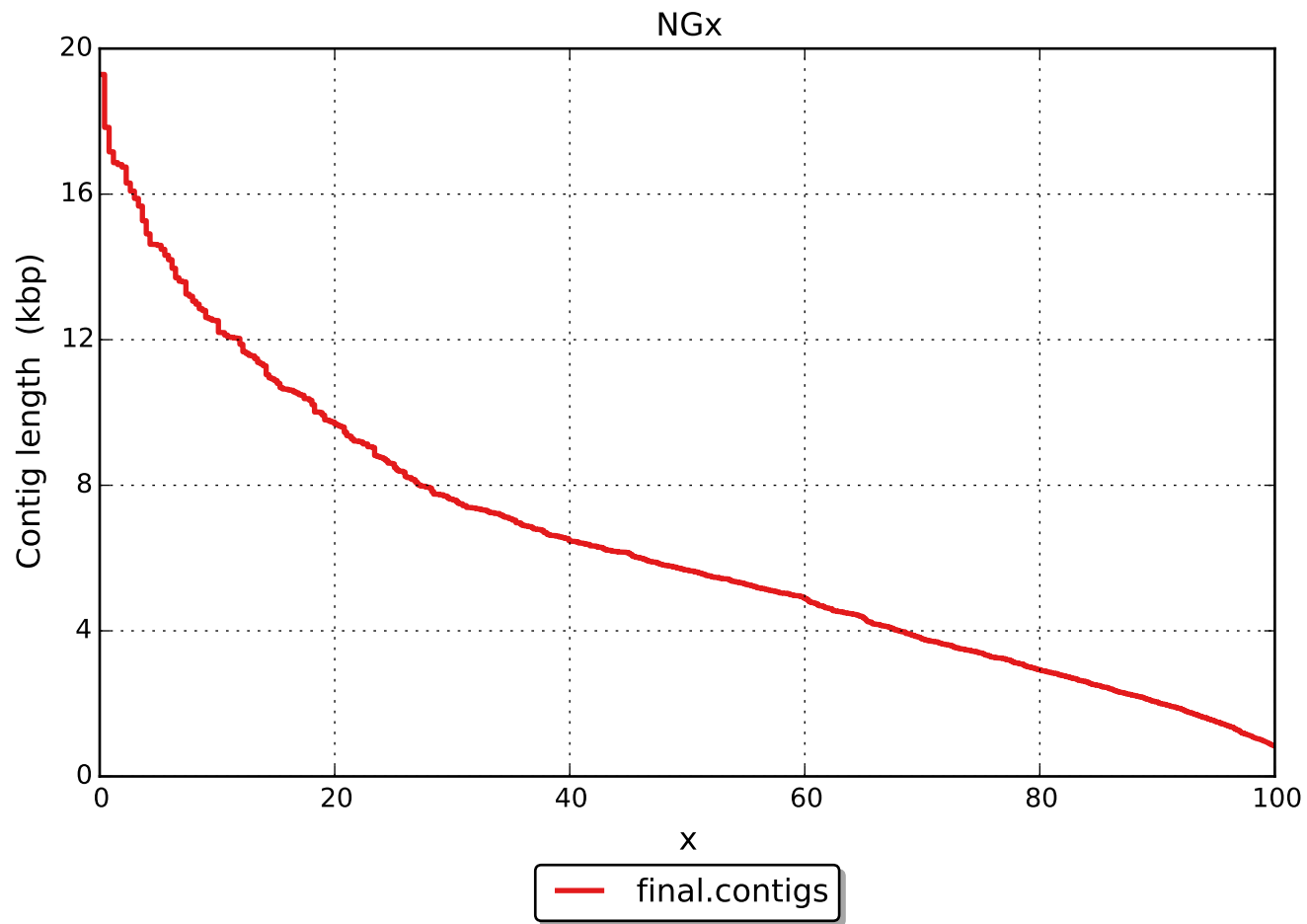
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

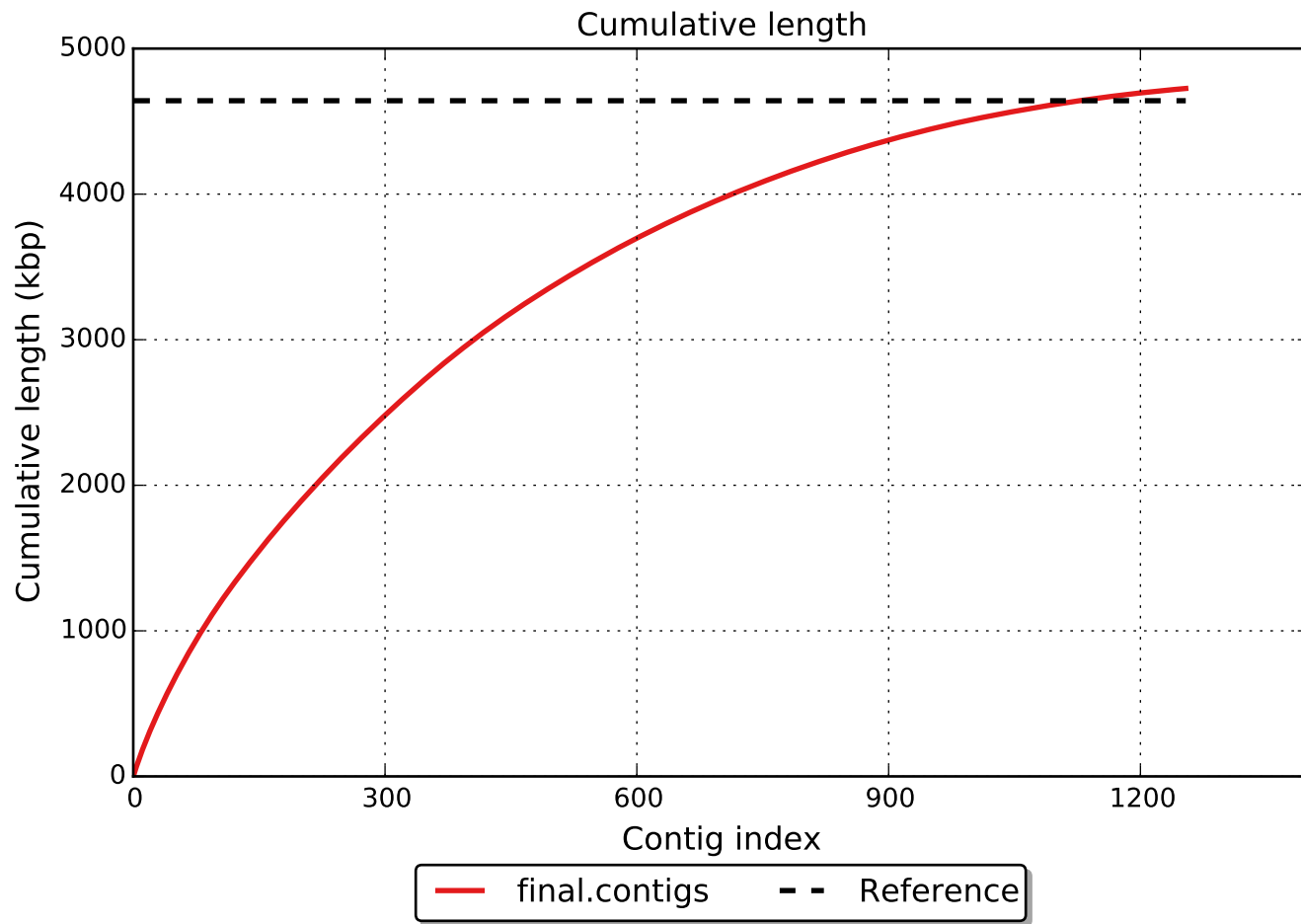
Unaligned report

	final.contigs
# fully unaligned contigs	0
Fully unaligned length	0
# partially unaligned contigs	1
# with misassembly	0
# both parts are significant	0
Partially unaligned length	29
# N's	0

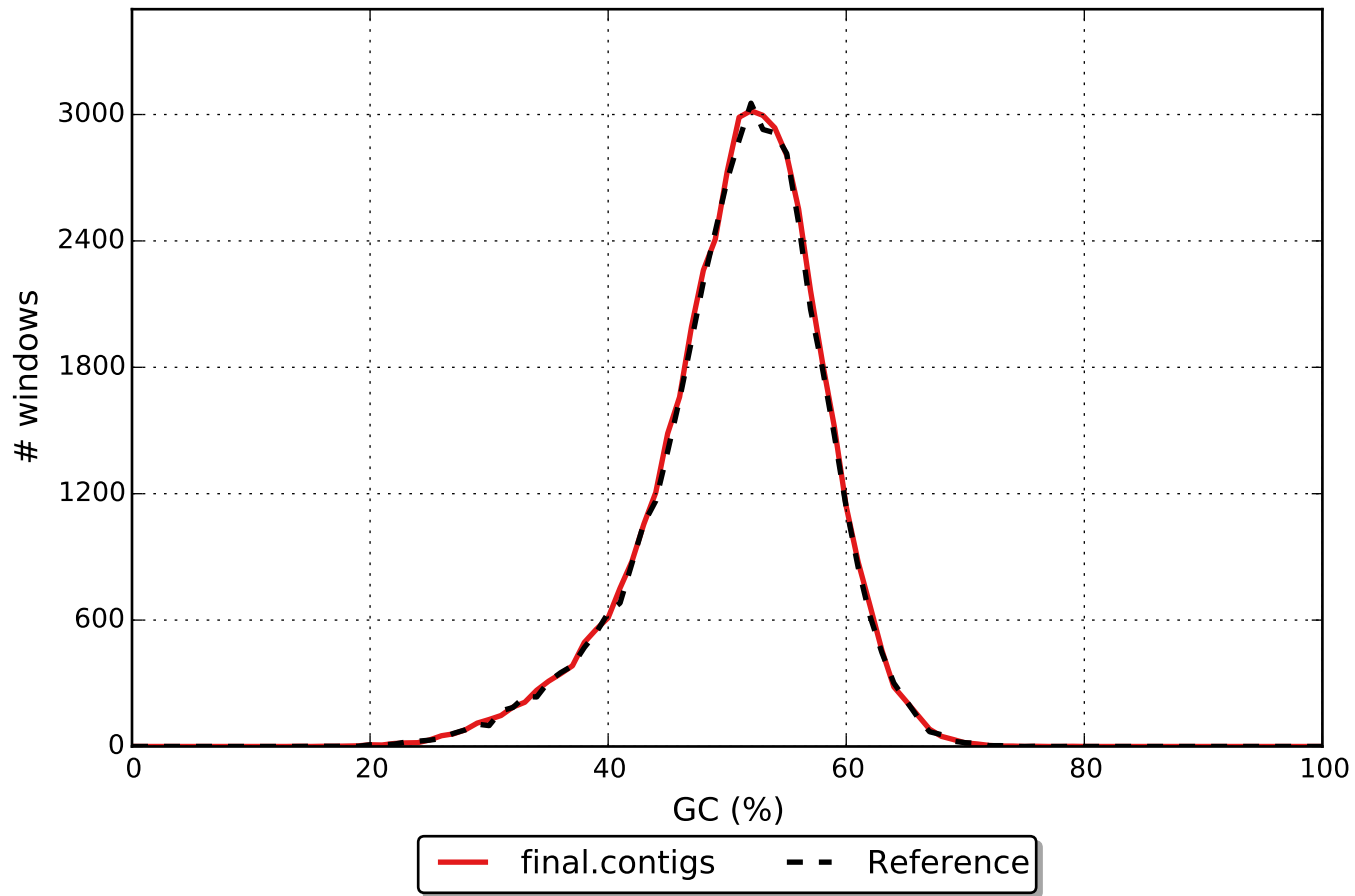
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).







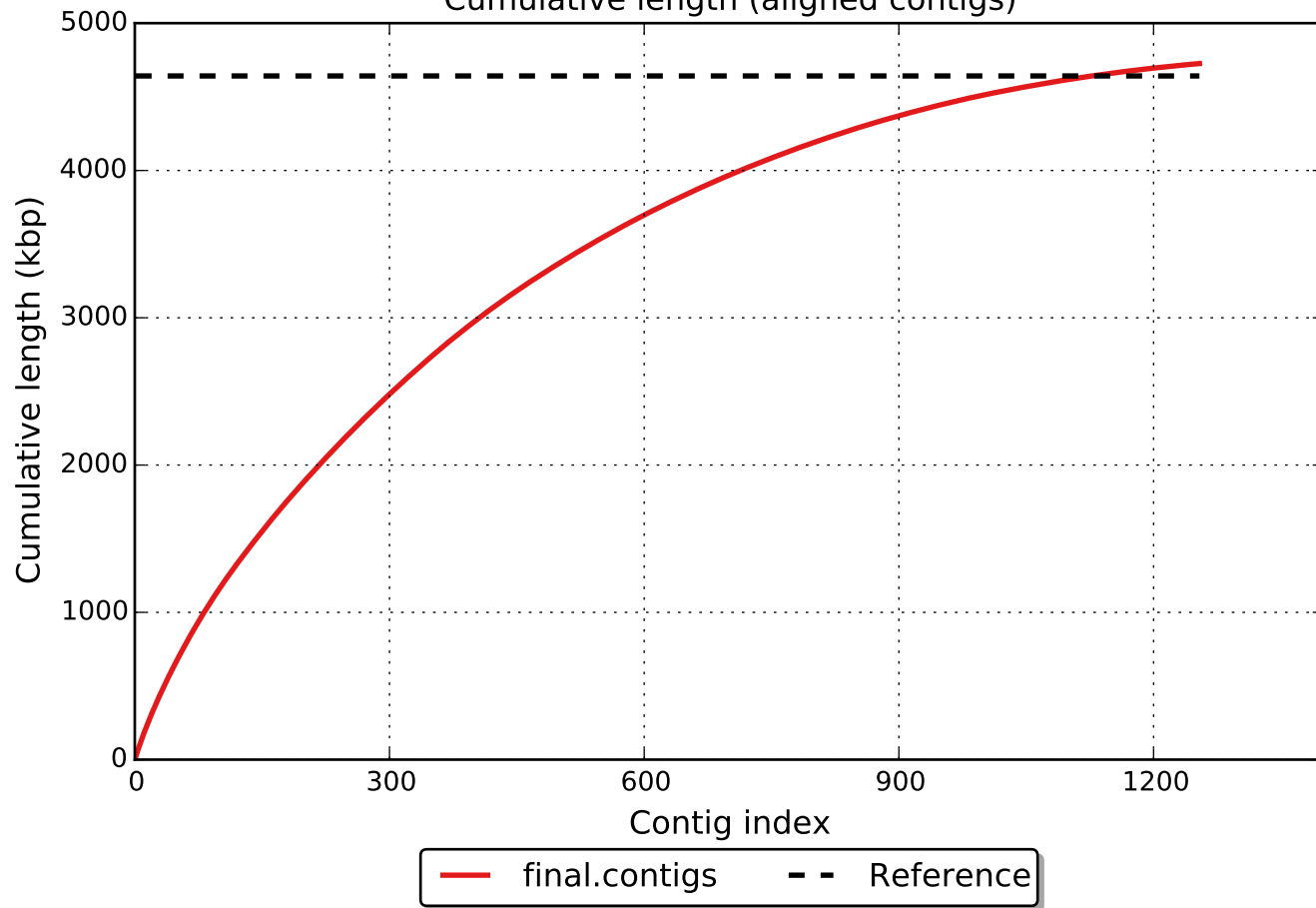
GC content

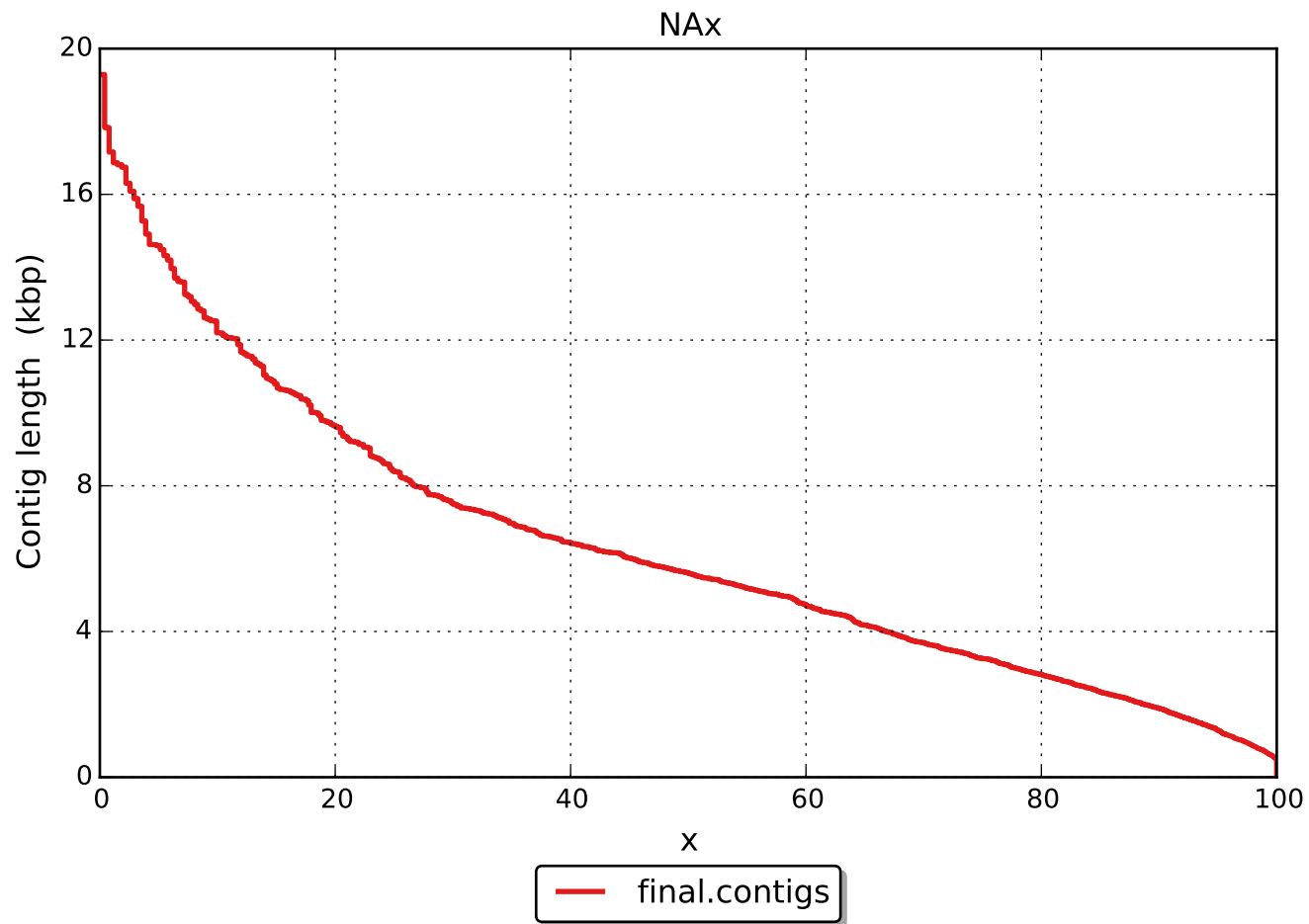


Misassemblies



Cumulative length (aligned contigs)





NGAx

