

Report

	final.contigs
# contigs (>= 1000 bp)	504
# contigs (>= 5000 bp)	314
# contigs (>= 10000 bp)	169
# contigs (>= 25000 bp)	26
# contigs (>= 50000 bp)	0
Total length (>= 1000 bp)	4646757
Total length (>= 5000 bp)	4102233
Total length (>= 10000 bp)	3055684
Total length (>= 25000 bp)	826549
Total length (>= 50000 bp)	0
# contigs	546
Largest contig	45186
Total length	4679094
Reference length	4641652
GC (℥)	50.79
Reference GC (℥)	50.78
N50	14276
NG50	14303
N75	7753
NG75	7811
L50	108
LG50	107
L75	221
LG75	218
# misassemblies	0
# misassembled contigs	0
Misassembled contigs length	0
# local misassemblies	0
# unaligned contigs	0 + 0 part
Unaligned length	0
Genome fraction (℥)	99.866
Duplication ratio	1.009
# N's per 100 kbp	0.00
# mismatches per 100 kbp	0.00
# indels per 100 kbp	0.00
Largest alignment	45186
NA50	14276
NGA50	14303
NA75	7753
NGA75	7811
LA50	108
LGA50	107
LA75	221
LGA75	218

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

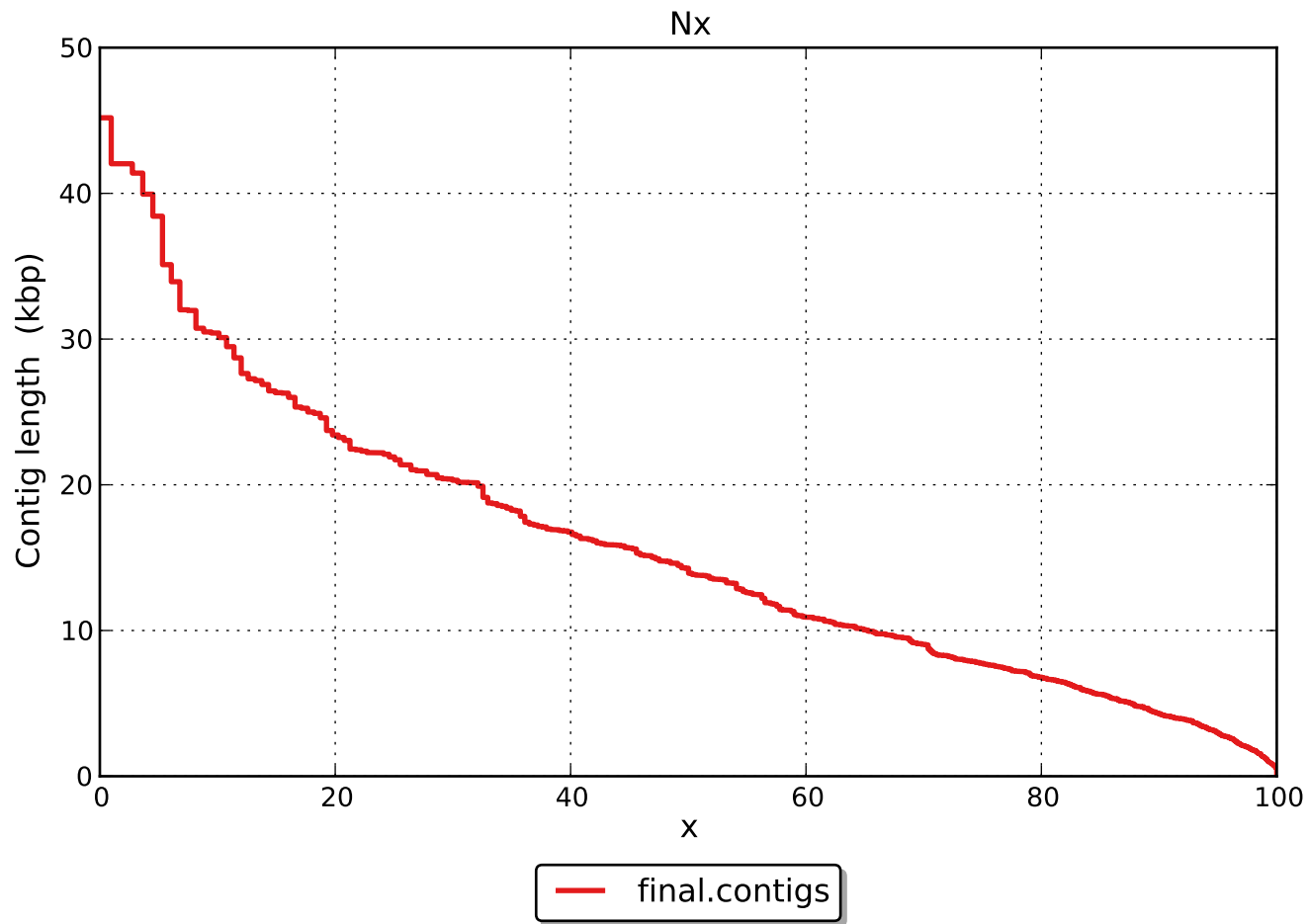
	final.contigs
# misassemblies	0
# relocations	0
# translocations	0
# inversions	0
# possibly misassembled contigs	0
# misassembled contigs	0
Misassembled contigs length	0
# local misassemblies	0
# mismatches	0
# indels	0
# short indels	0
# long indels	0
Indels length	0

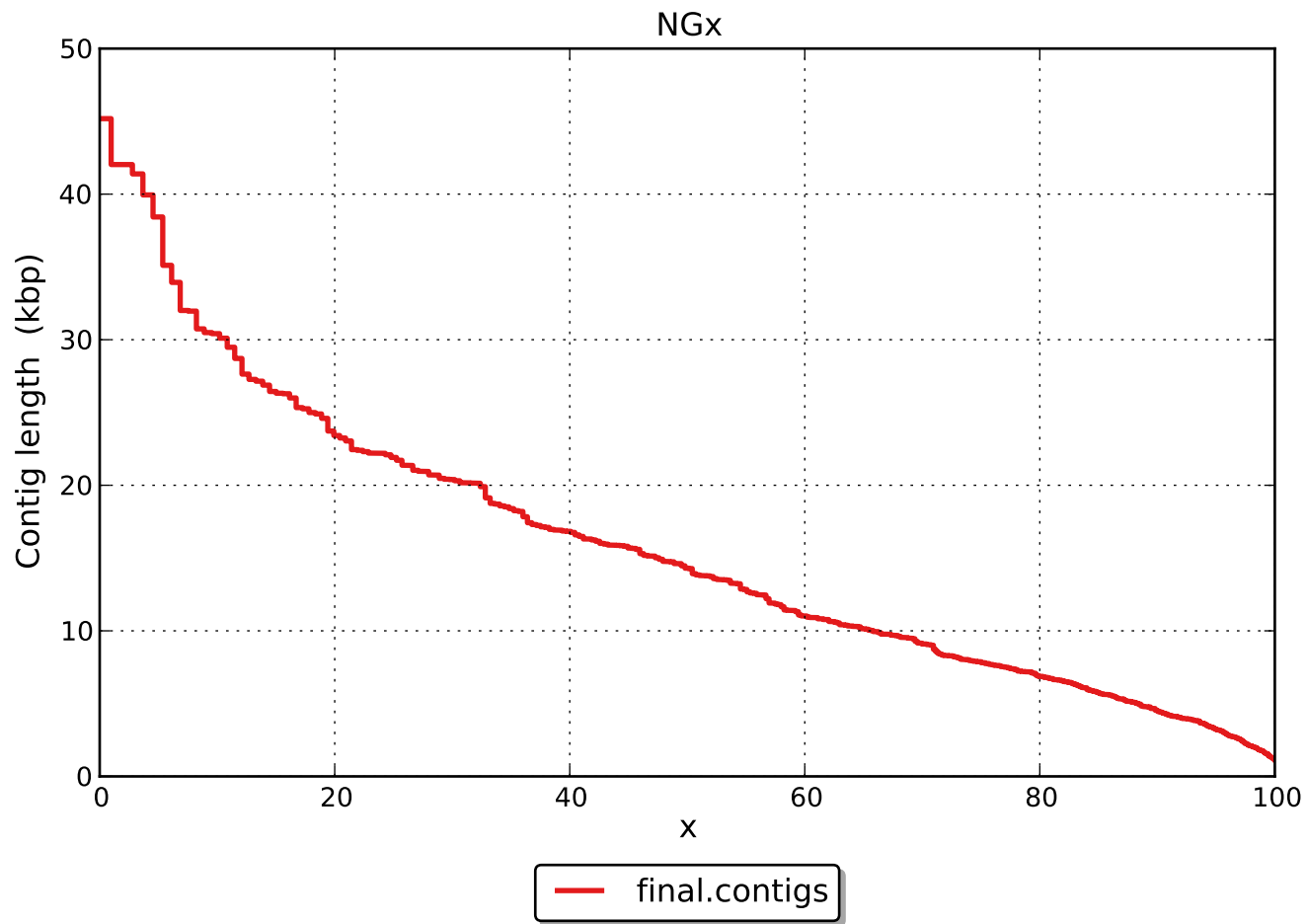
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

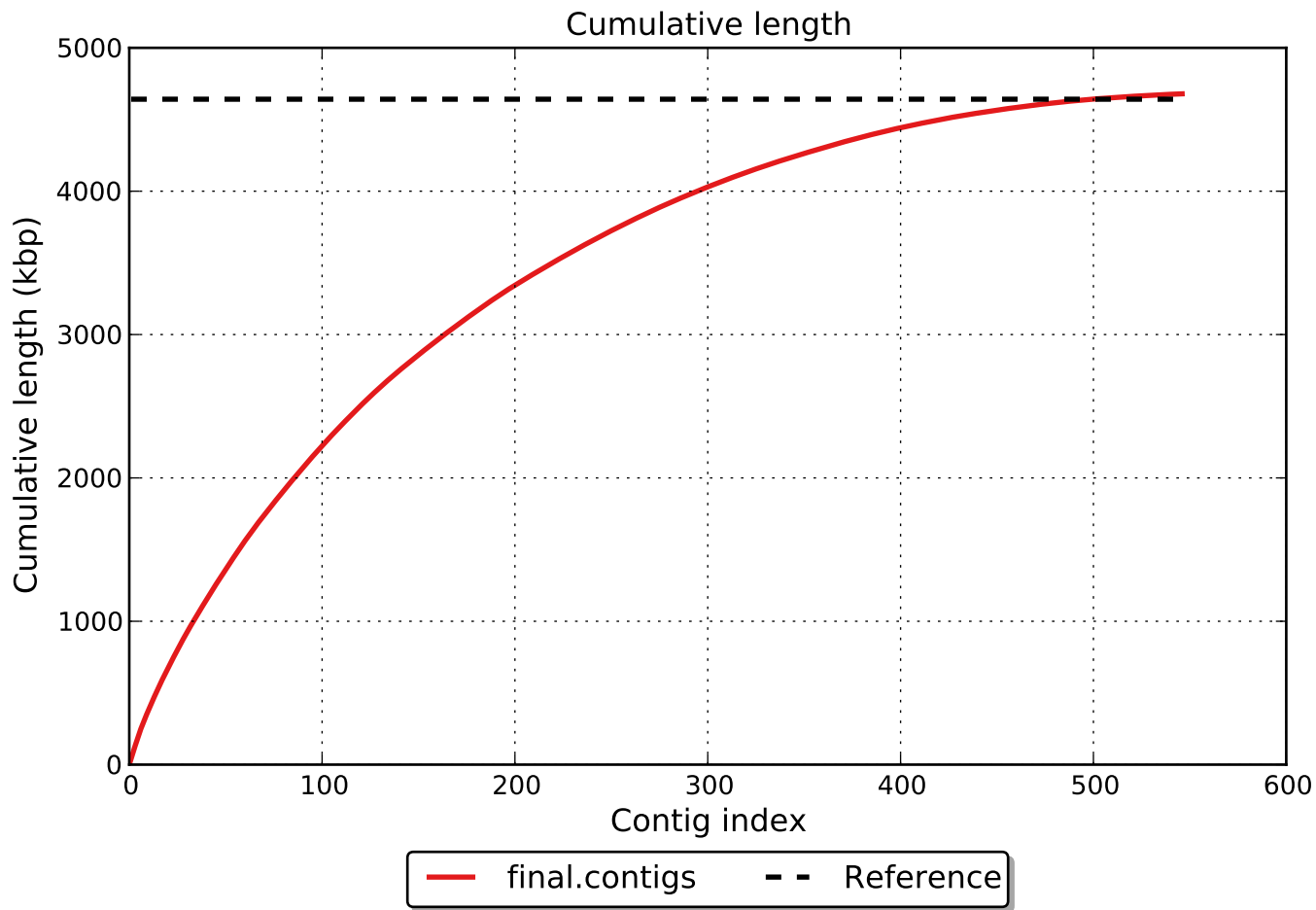
Unaligned report

	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

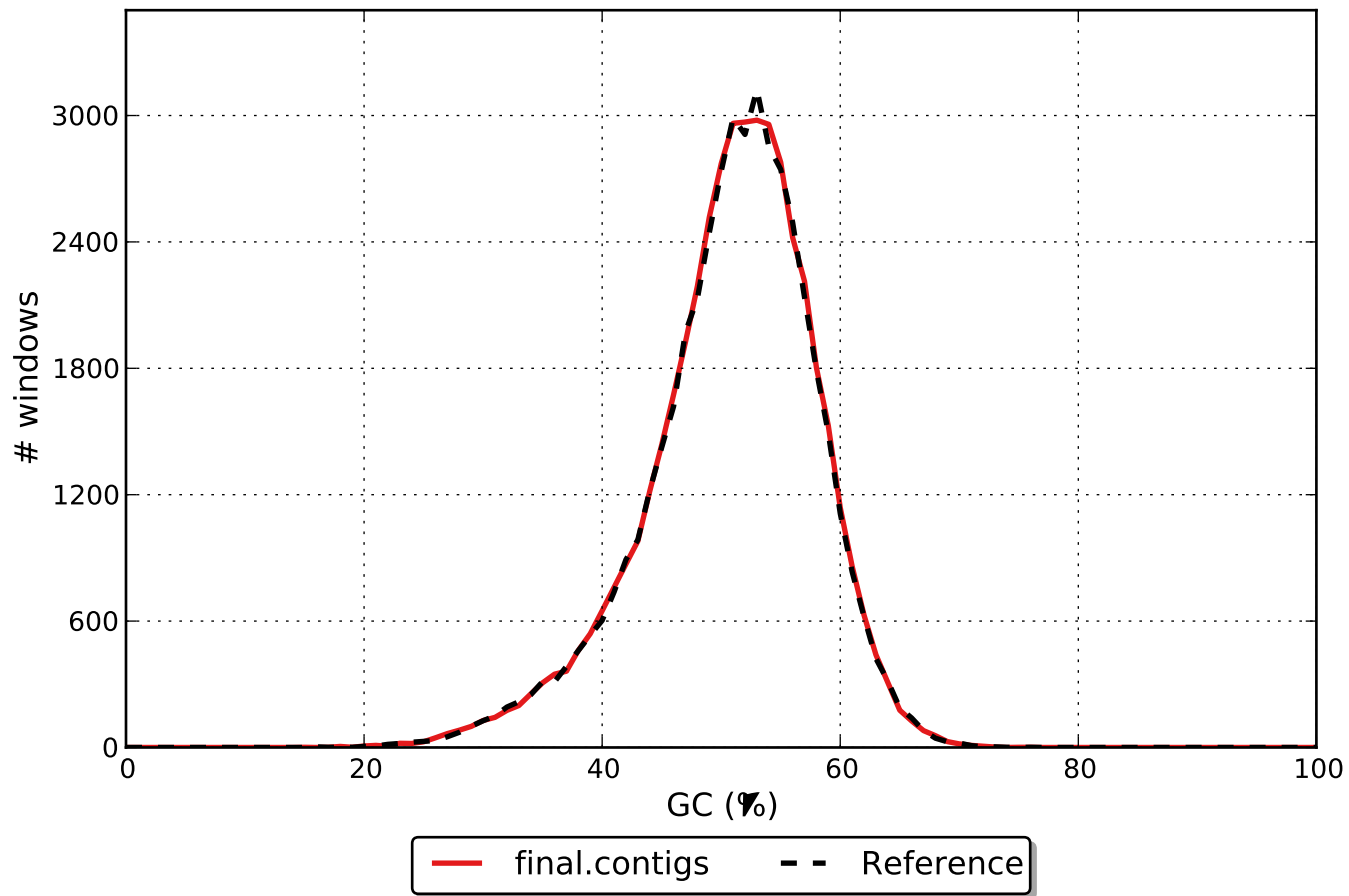
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).



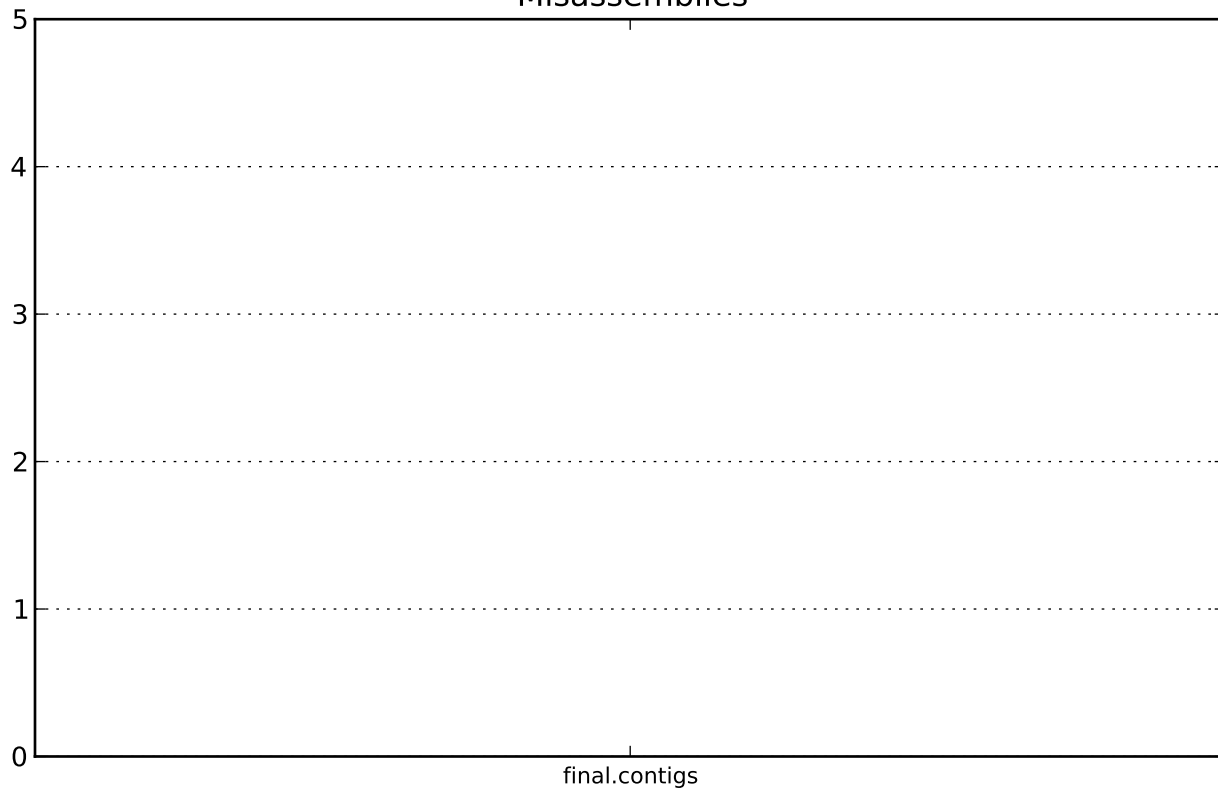




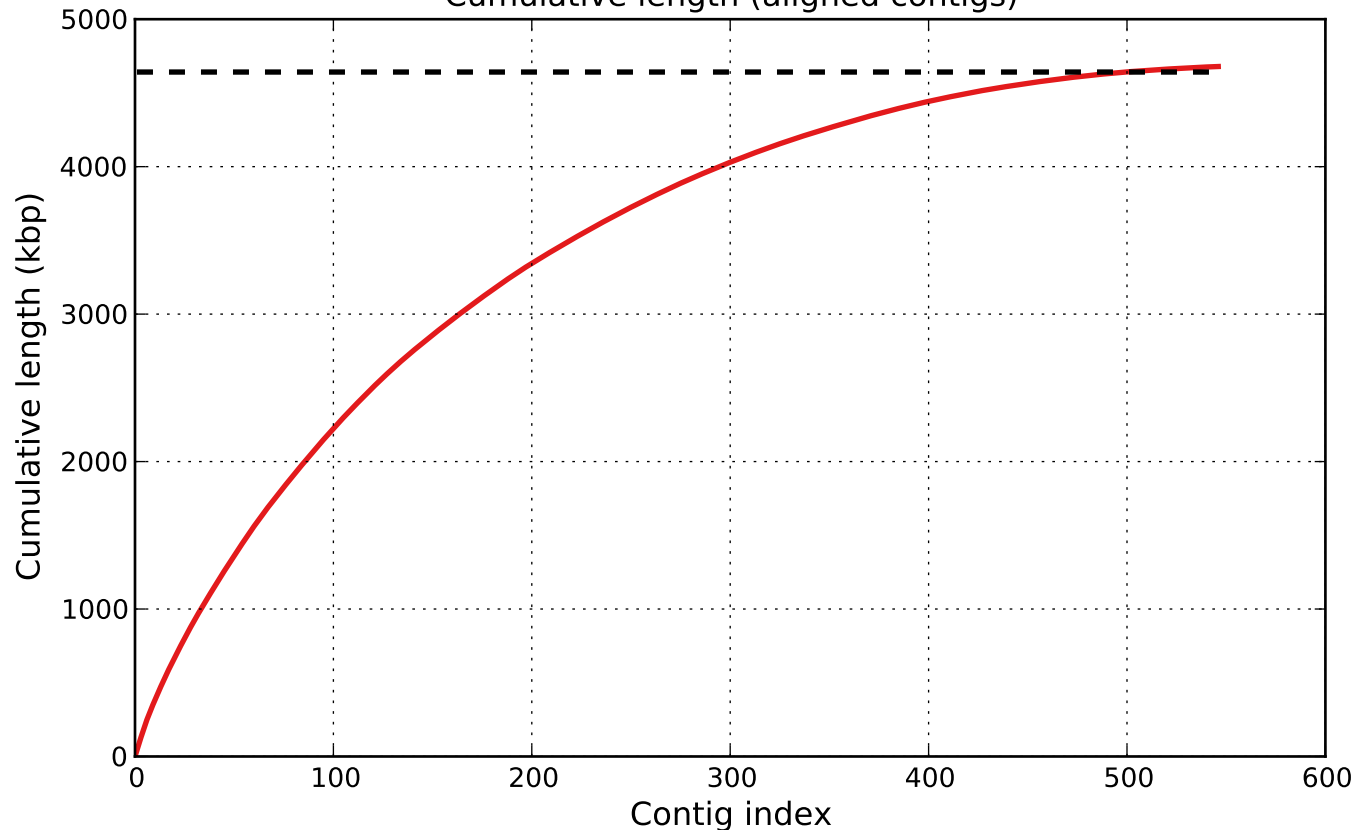
GC content



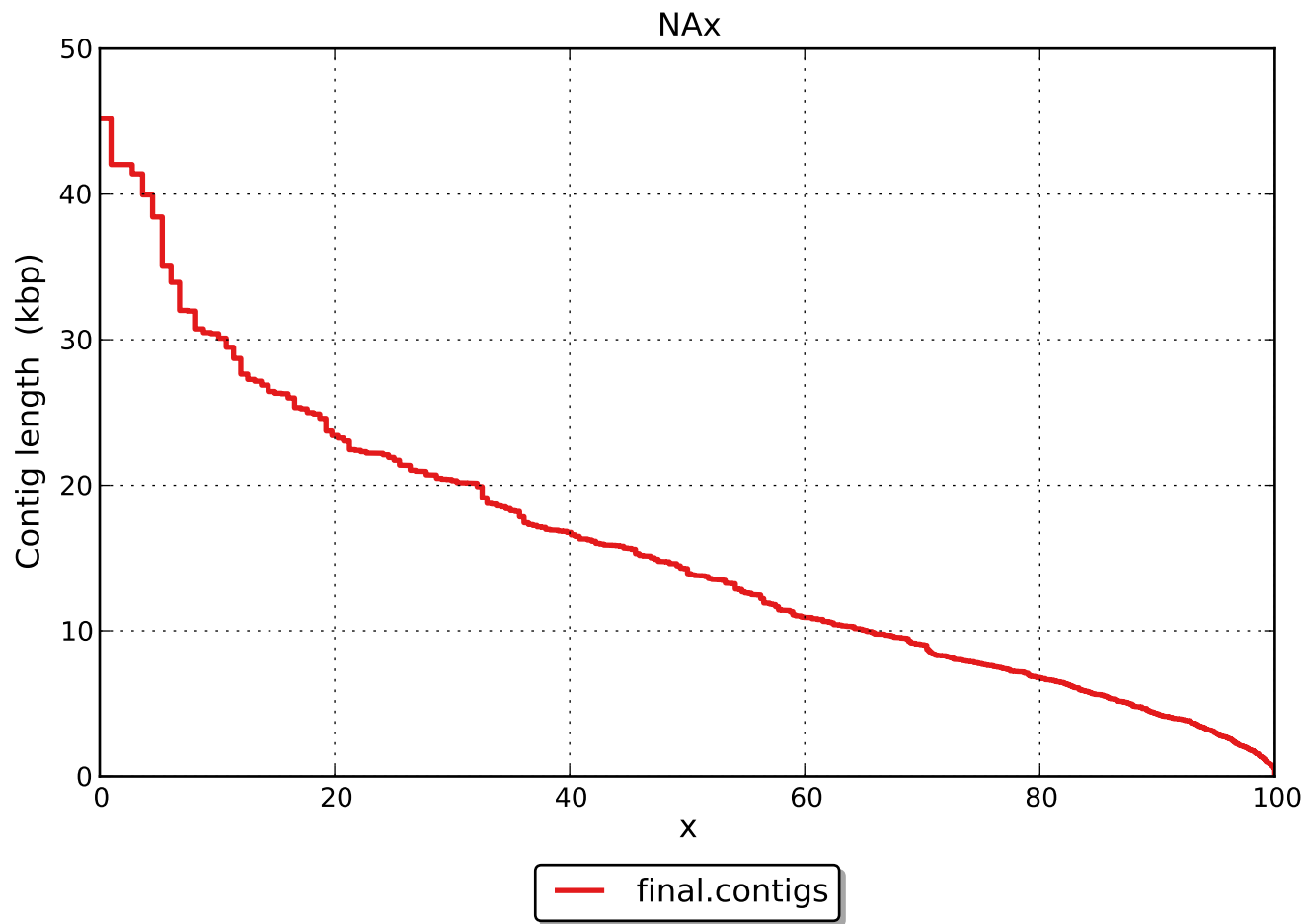
Misassemblies



Cumulative length (aligned contigs)



— final.contigs - - Reference



NGAx

