

## Report

	final.contigs
# contigs ( $\geq 1000$ bp)	664
# contigs ( $\geq 5000$ bp)	102
# contigs ( $\geq 10000$ bp)	9
# contigs ( $\geq 25000$ bp)	0
# contigs ( $\geq 50000$ bp)	0
Total length ( $\geq 1000$ bp)	2018796
Total length ( $\geq 5000$ bp)	710807
Total length ( $\geq 10000$ bp)	102572
Total length ( $\geq 25000$ bp)	0
Total length ( $\geq 50000$ bp)	0
# contigs	1241
Largest contig	16503
Total length	2402300
Reference length	4641652
GC (‰)	50.84
Reference GC (‰)	50.79
N50	3176
NG50	554
N75	1568
L50	228
LG50	1087
L75	492
# misassemblies	0
# misassembled contigs	0
Misassembled contigs length	0
# local misassemblies	0
# unaligned contigs	0 + 0 part
Unaligned length	0
Genome fraction (‰)	50.583
Duplication ratio	1.023
# N's per 100 kbp	0.00
# mismatches per 100 kbp	418.97
# indels per 100 kbp	0.09
Largest alignment	16503
NA50	3176
NGA50	554
NA75	1568
LA50	228
LGA50	1087
LA75	492

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

## Misassemblies report

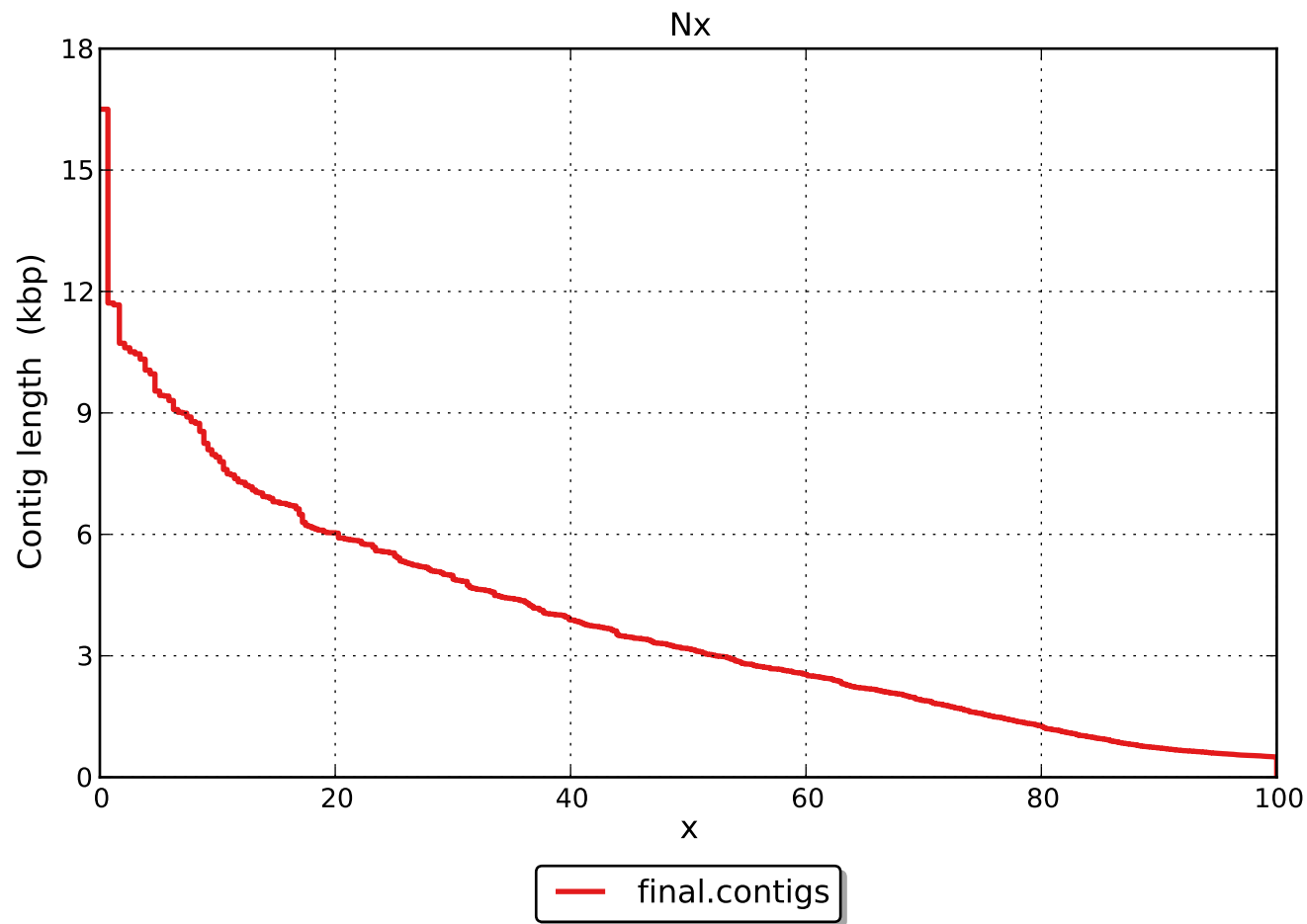
	final.contigs
# misassemblies	0
# relocations	0
# translocations	0
# inversions	0
# possibly misassembled contigs	0
# misassembled contigs	0
Misassembled contigs length	0
# local misassemblies	0
# mismatches	9837
# indels	2
# short indels	2
# long indels	0
Indels length	2

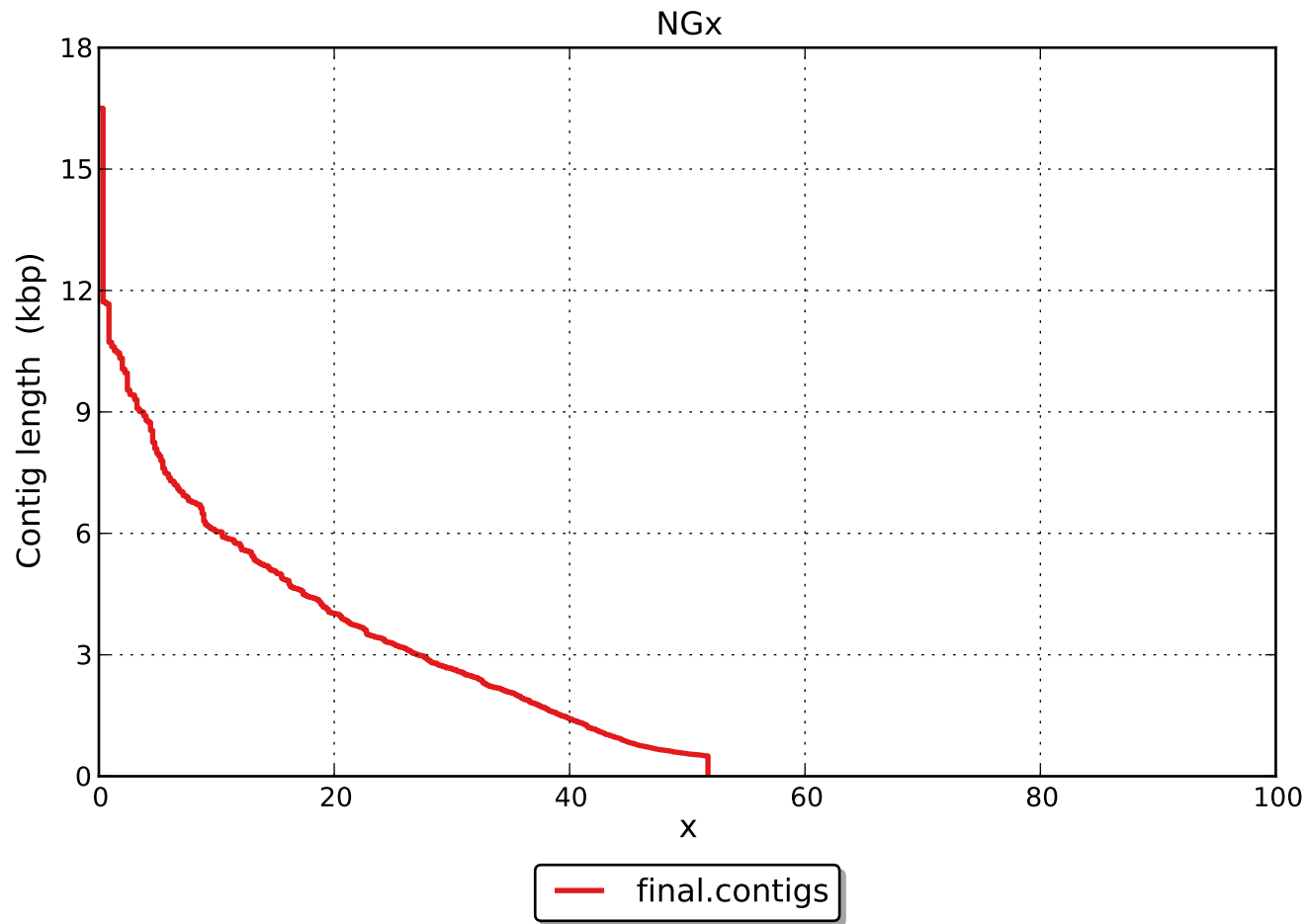
All statistics are based on contigs of size  $\geq 500$  bp, unless otherwise noted (e.g., "# contigs ( $\geq 0$  bp)" and "Total length ( $\geq 0$  bp)" include all contigs).

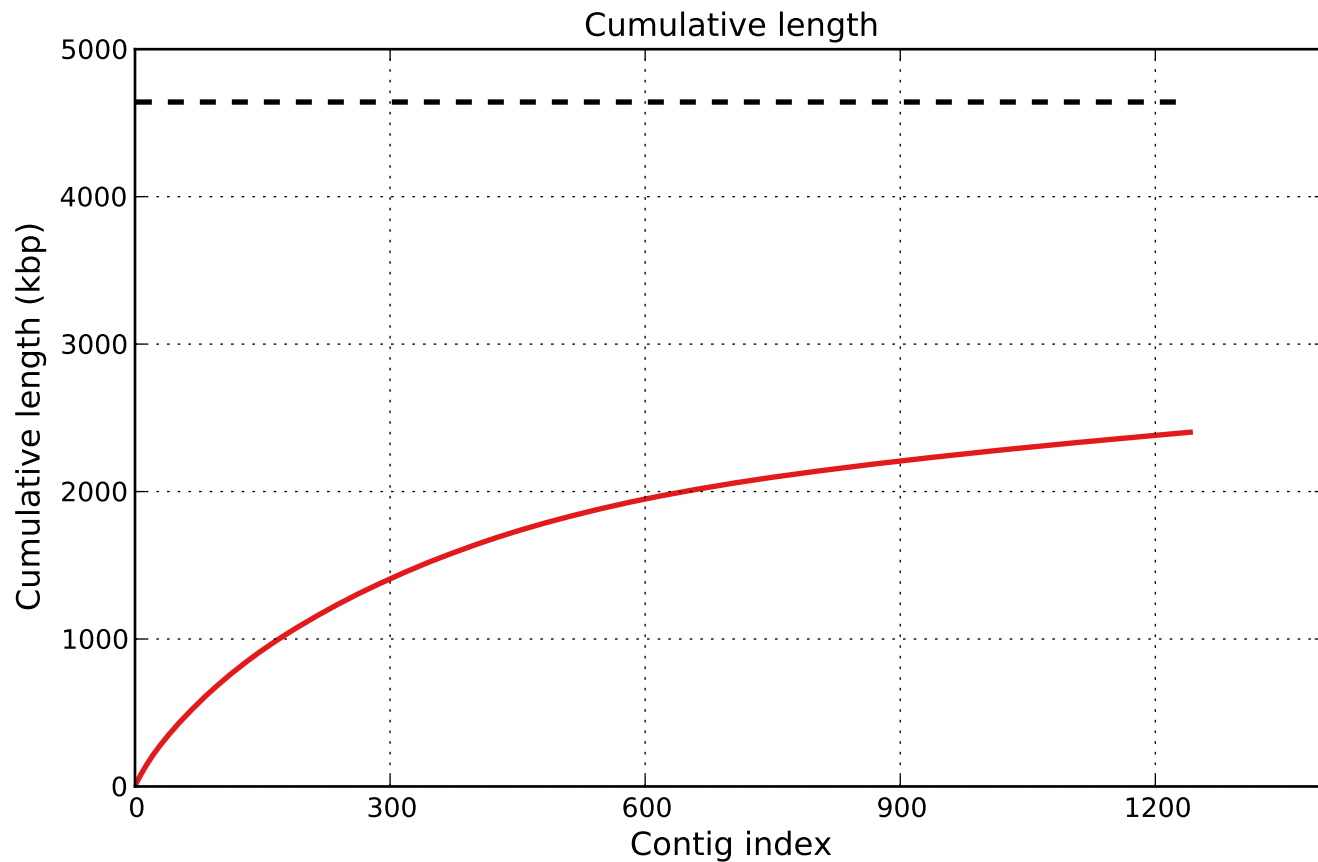
## Unaligned report

	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

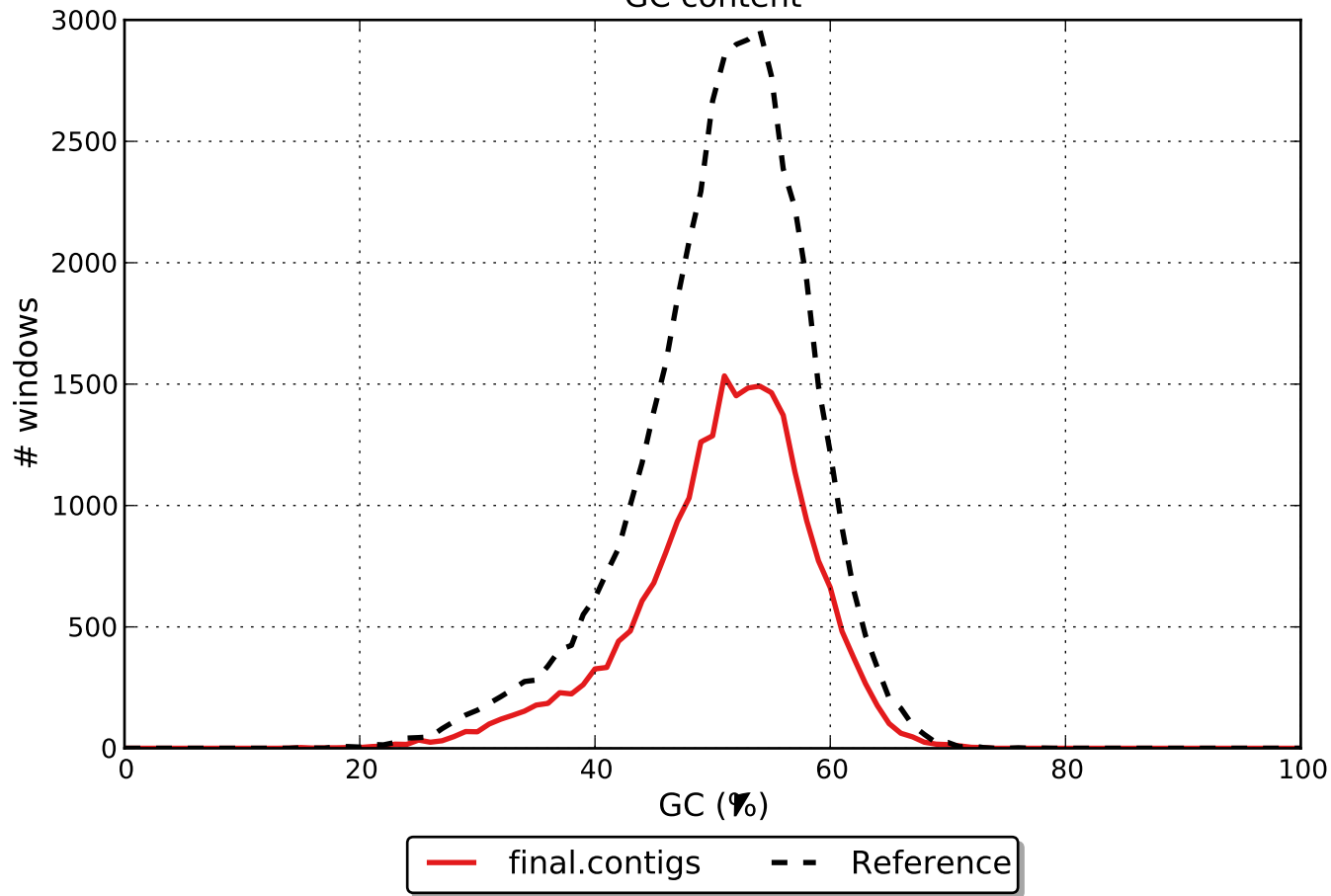
All statistics are based on contigs of size  $\geq 500$  bp, unless otherwise noted (e.g., "# contigs ( $\geq 0$  bp)" and "Total length ( $\geq 0$  bp)" include all contigs).







GC content

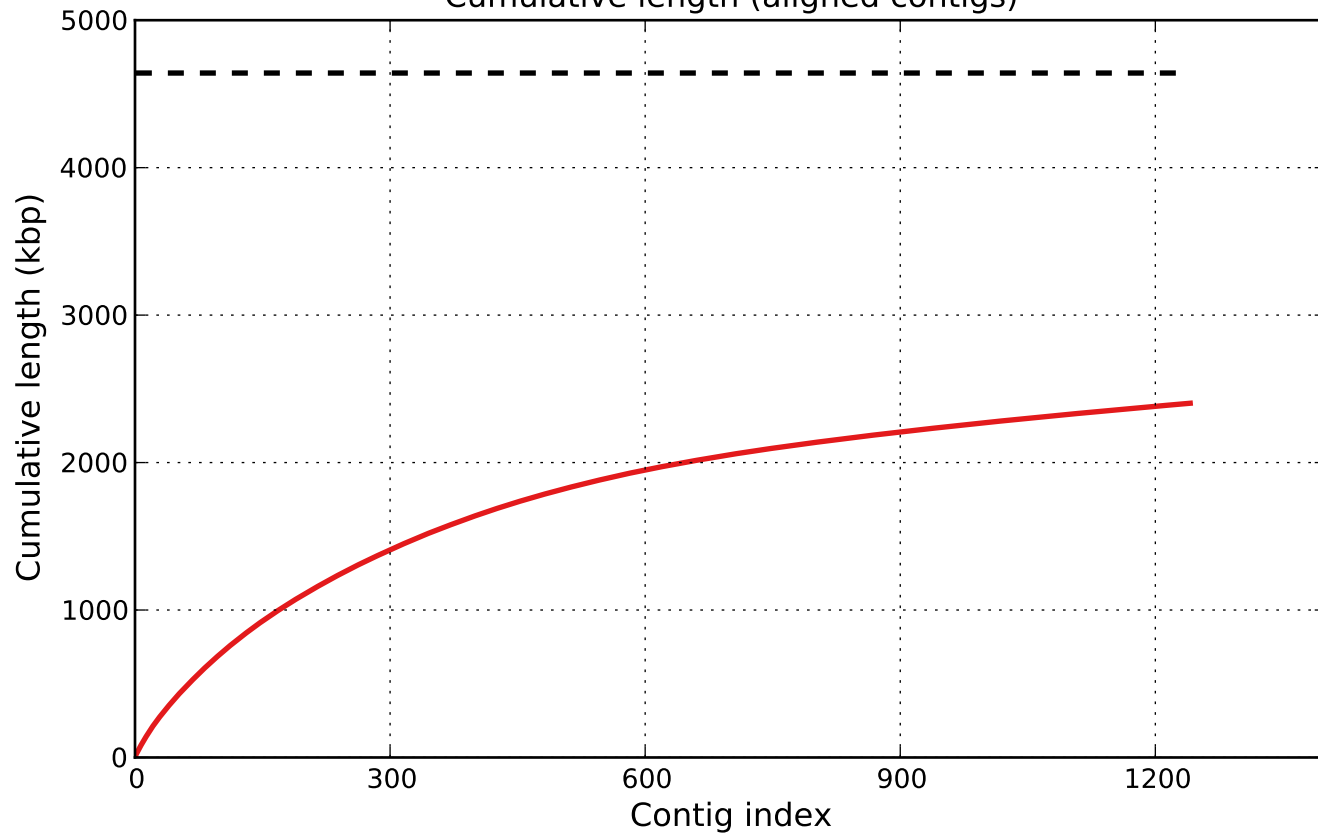


# Misassemblies





Cumulative length (aligned contigs)



— final.contigs    - - Reference

