

Report

	contigs
# contigs (≥ 0 bp)	118
# contigs (≥ 1000 bp)	72
# contigs (≥ 5000 bp)	52
# contigs (≥ 10000 bp)	48
# contigs (≥ 25000 bp)	42
# contigs (≥ 50000 bp)	27
Total length (≥ 0 bp)	4577280
Total length (≥ 1000 bp)	4561376
Total length (≥ 5000 bp)	4516669
Total length (≥ 10000 bp)	4486780
Total length (≥ 25000 bp)	4396467
Total length (≥ 50000 bp)	3869162
# contigs	83
Largest contig	327173
Total length	4569160
Reference length	4641652
GC (%)	50.74
Reference GC (%)	50.79
N50	173566
NG50	133063
N75	78670
NG75	78670
L50	10
LG50	11
L75	21
LG75	21
# misassemblies	0
# misassembled contigs	0
Misassembled contigs length	0
# local misassemblies	0
# unaligned contigs	0 + 0 part
Unaligned length	0
Genome fraction (%)	98.365
Duplication ratio	1.001
# N's per 100 kbp	0.00
# mismatches per 100 kbp	1.25
# indels per 100 kbp	0.13
Largest alignment	327173
NA50	172028
NGA50	133063
NA75	78670
NGA75	78670
LA50	10
LGA50	11
LA75	21
LGA75	21

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

	contigs
# misassemblies	0
# relocations	0
# translocations	0
# inversions	0
# misassembled contigs	0
Misassembled contigs length	0
# local misassemblies	0
# mismatches	57
# indels	6
# short indels	6
# long indels	0
Indels length	7

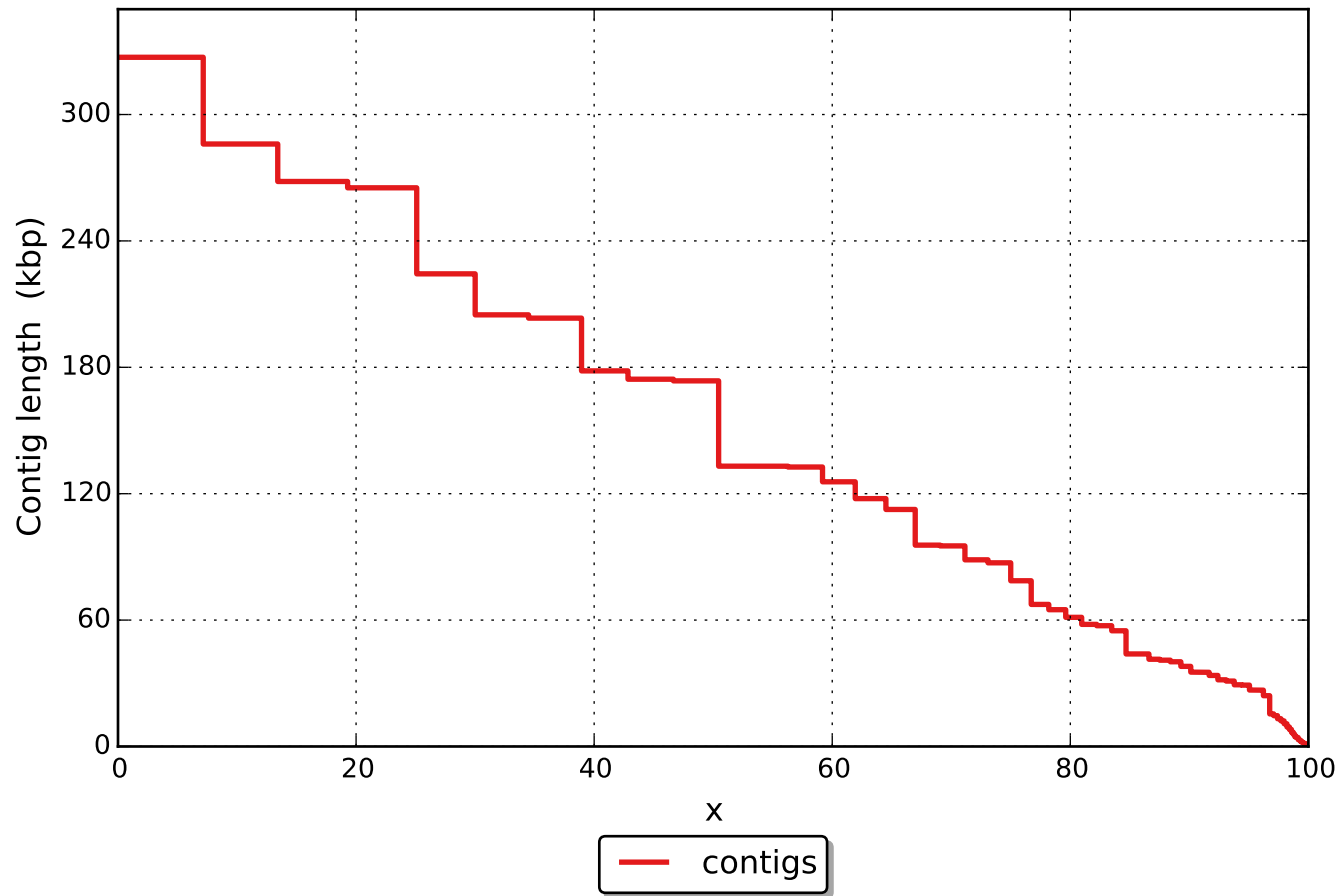
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

Unaligned report

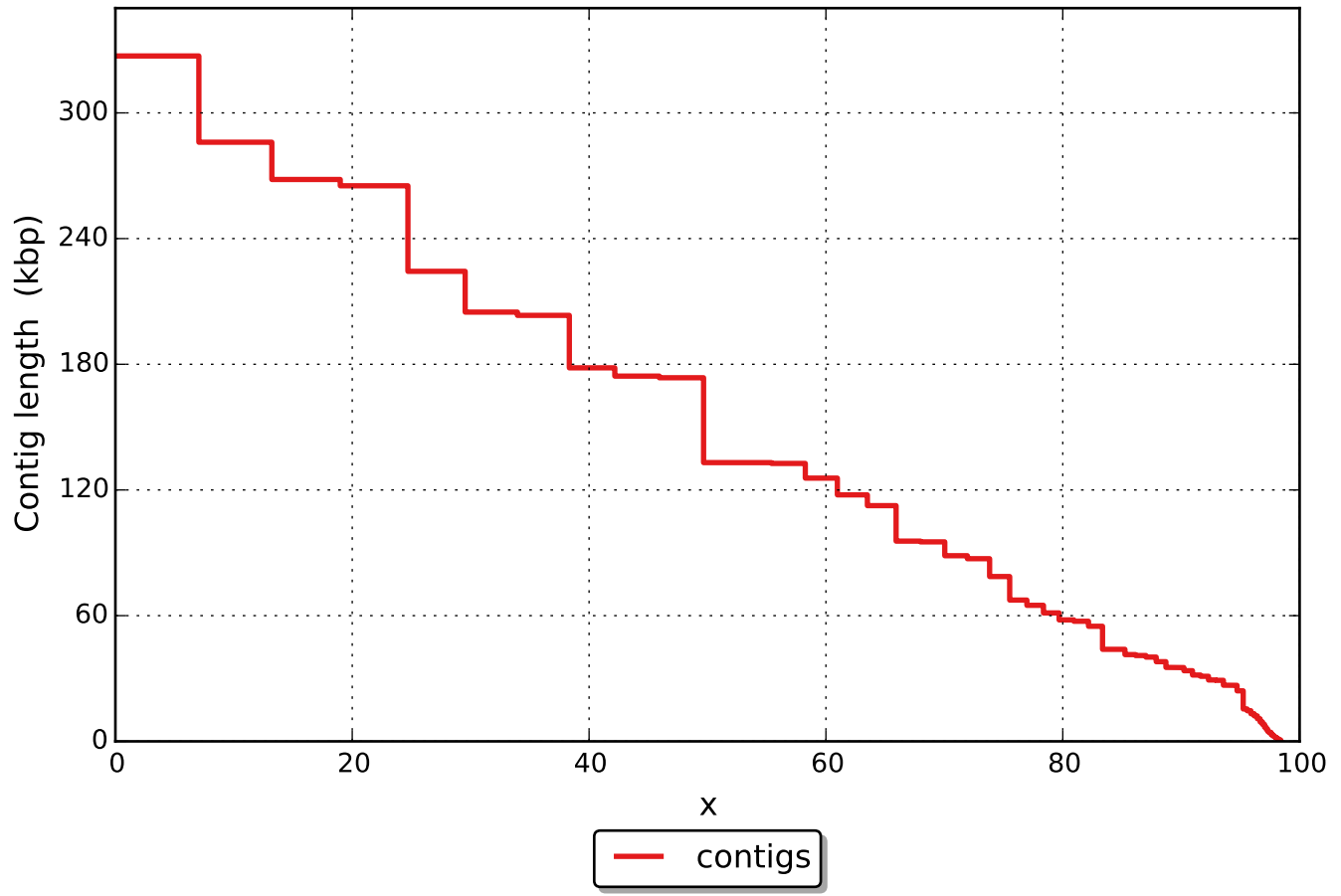
	contigs
# fully unaligned contigs	0
Fully unaligned length	0
# partially unaligned contigs	0
# with misassembly	0
# both parts are significant	0
Partially unaligned length	0
# N's	0

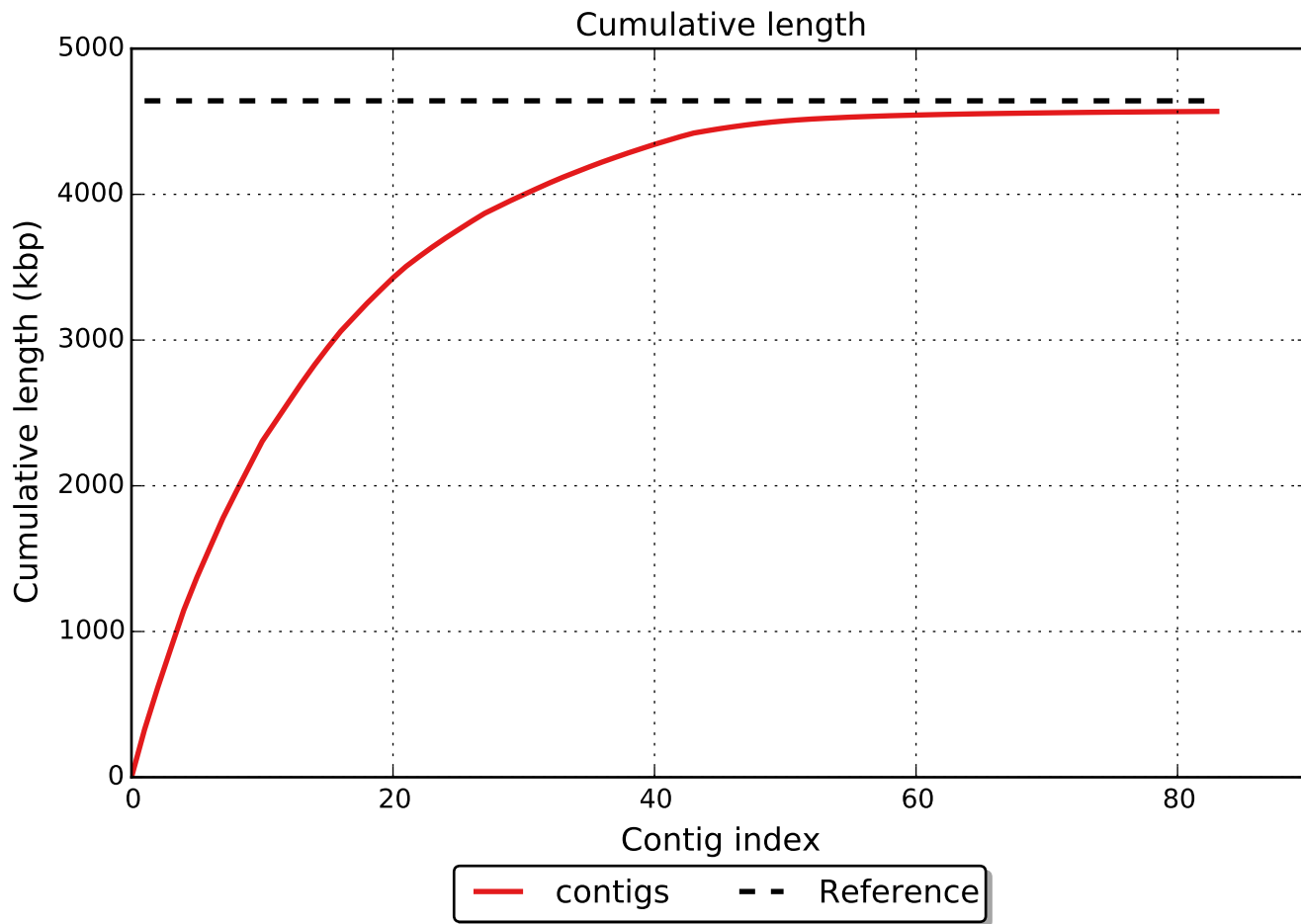
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

Nx

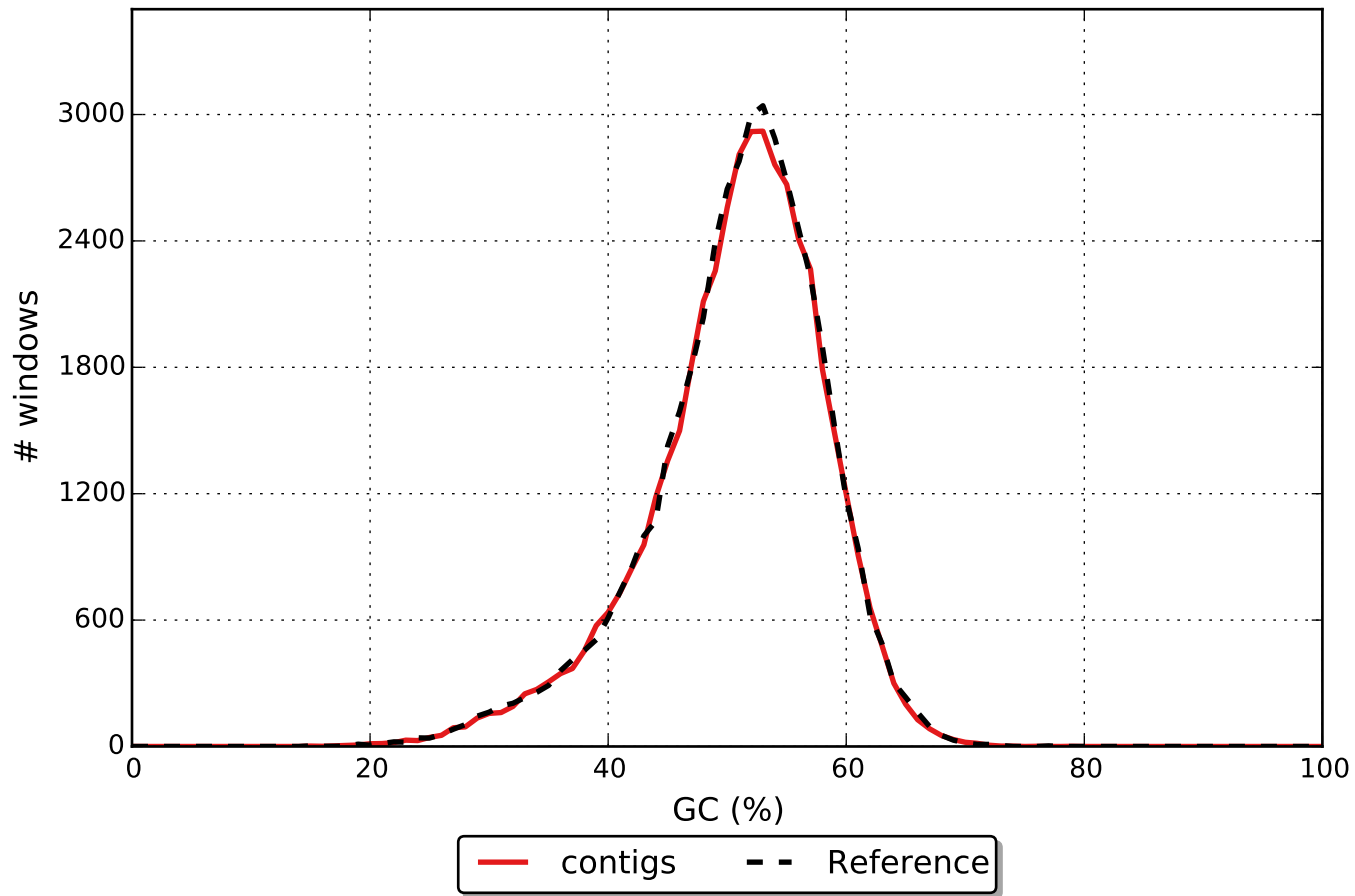


NGx





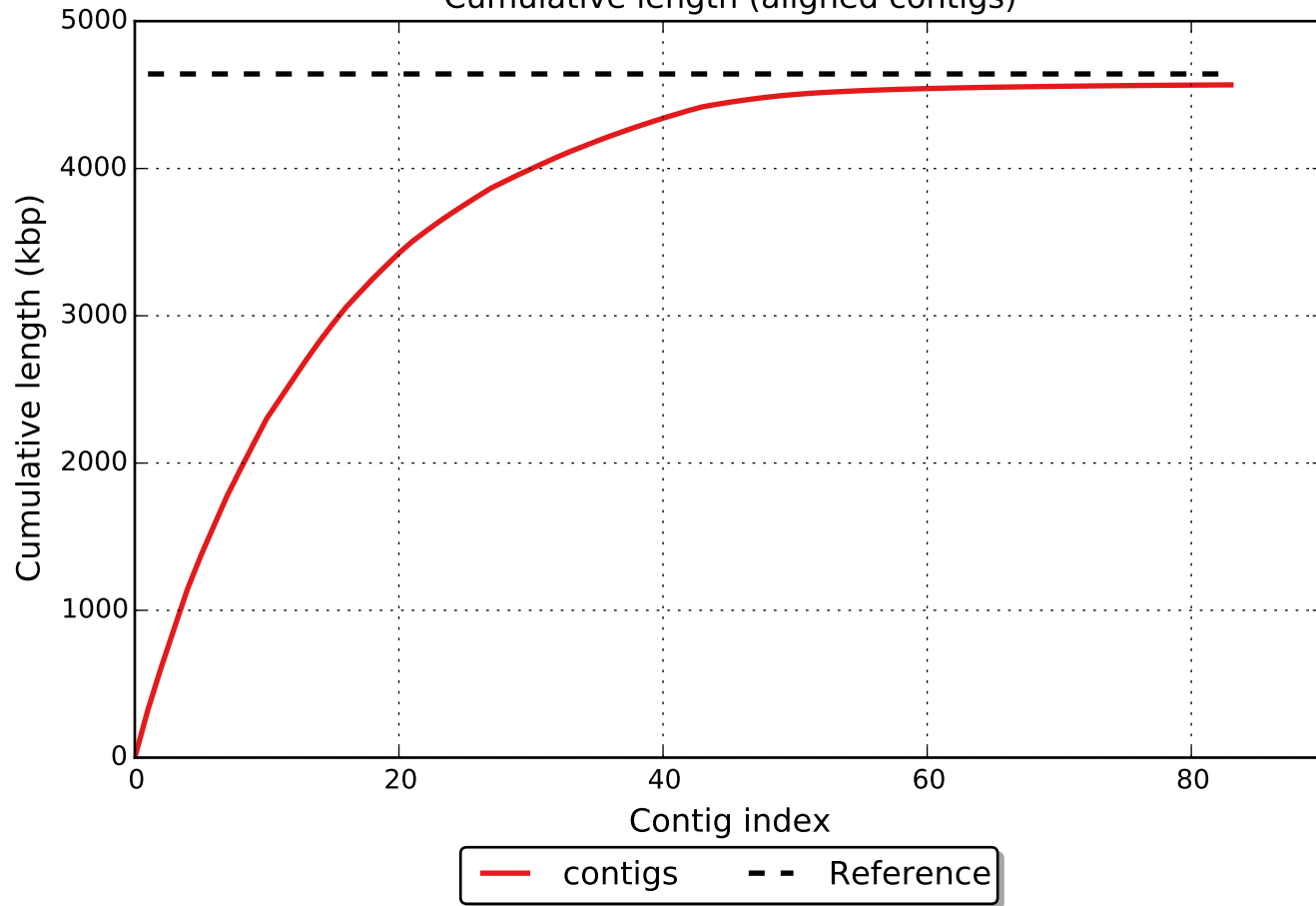
GC content



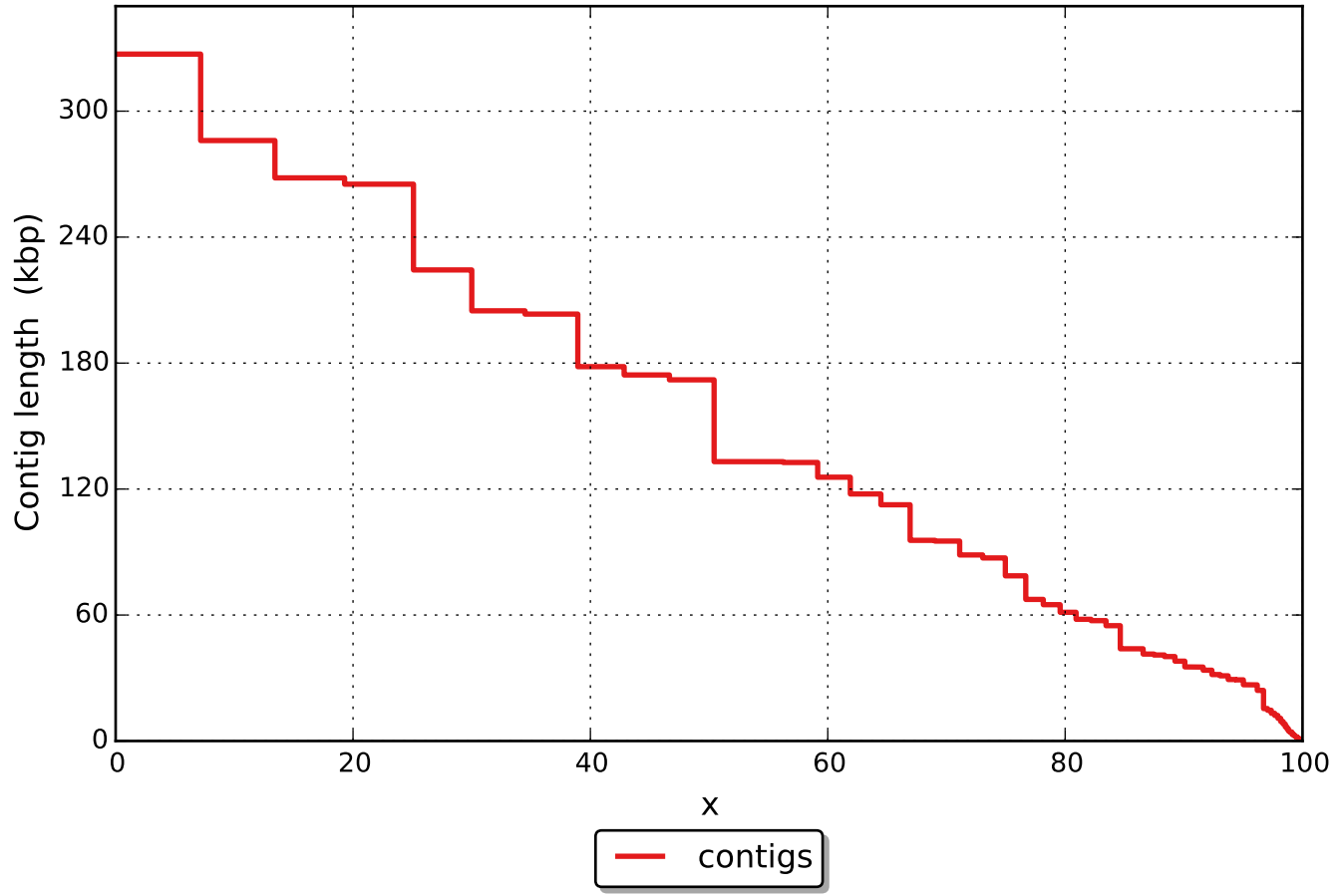
Misassemblies



Cumulative length (aligned contigs)



NAx



NGAx

