

Report

	scaffolds
# contigs (≥ 1000 bp)	150
# contigs (≥ 5000 bp)	108
# contigs (≥ 10000 bp)	82
# contigs (≥ 25000 bp)	40
# contigs (≥ 50000 bp)	21
Total length (≥ 1000 bp)	3521952
Total length (≥ 5000 bp)	3428883
Total length (≥ 10000 bp)	3229812
Total length (≥ 25000 bp)	2532410
Total length (≥ 50000 bp)	1861491
# contigs	182
Largest contig	186891
Total length	3541698
Reference length	4641652
GC (%)	50.88
Reference GC (%)	50.79
N50	52546
NG50	33321
N75	22200
NG75	2361
L50	20
LG50	33
L75	46
LG75	124
# misassemblies	1
# misassembled contigs	1
Misassembled contigs length	66116
# local misassemblies	2
# unaligned contigs	0 + 0 part
Unaligned length	0
Genome fraction (%)	75.323
Duplication ratio	1.013
# N's per 100 kbp	0.00
# mismatches per 100 kbp	1009.55
# indels per 100 kbp	0.72
Largest alignment	186891
NA50	51199
NGA50	31609
NA75	22200
NGA75	2361
LA50	20
LGA50	34
LA75	46
LGA75	125

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

	scaffolds
# misassemblies	1
# relocations	1
# translocations	0
# inversions	0
# possibly misassembled contigs	0
# misassembled contigs	1
Misassembled contigs length	66116
# local misassemblies	2
# mismatches	35296
# indels	25
# short indels	25
# long indels	0
Indels length	31

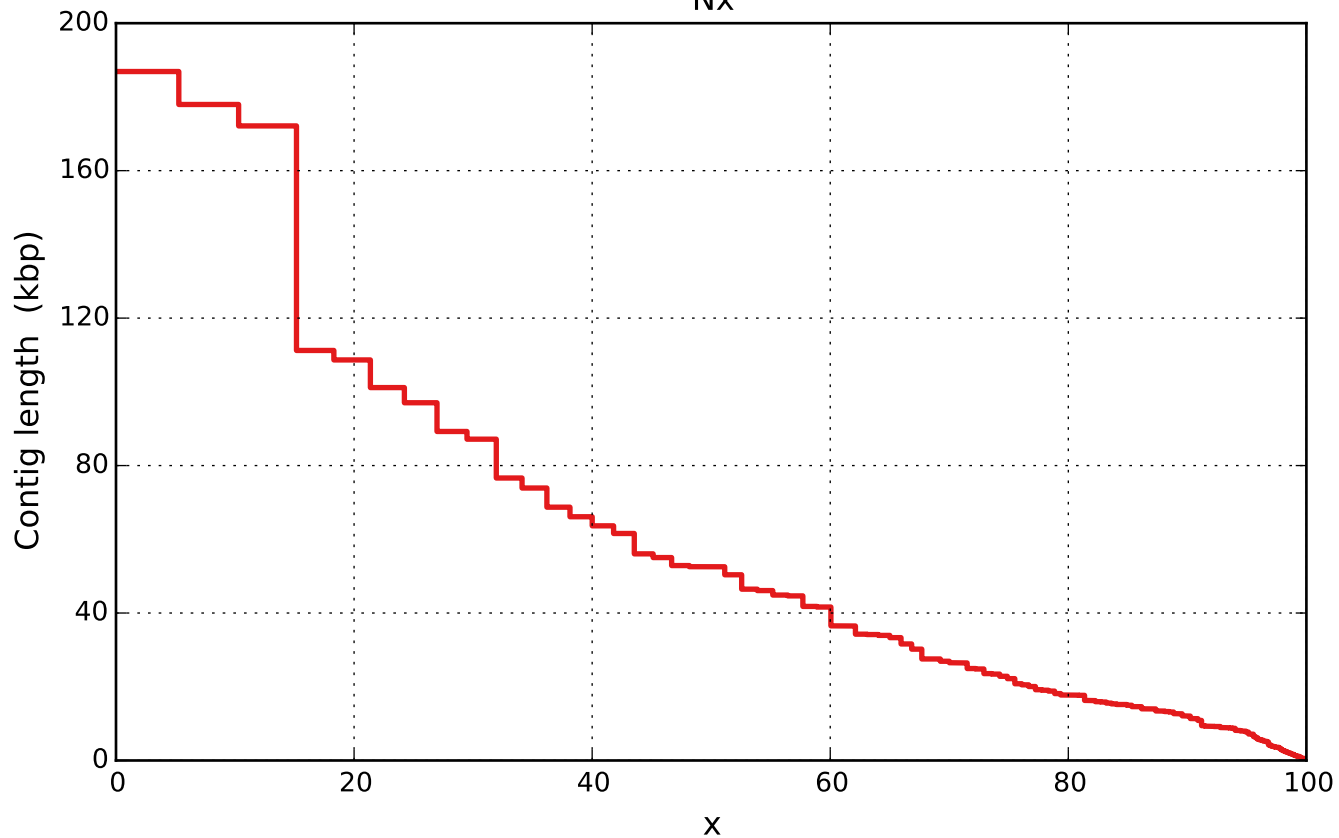
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

Unaligned report

	scaffolds
# 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

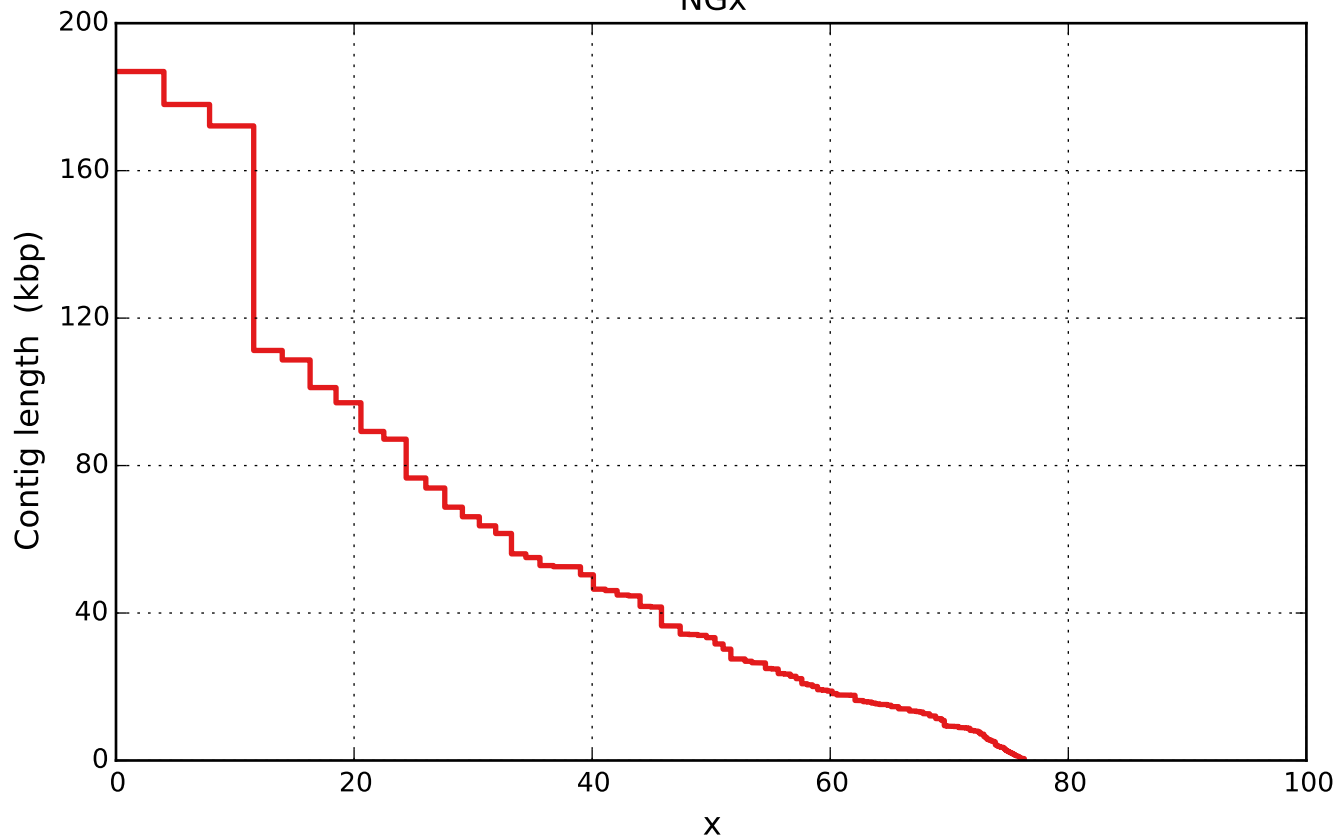
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

Nx

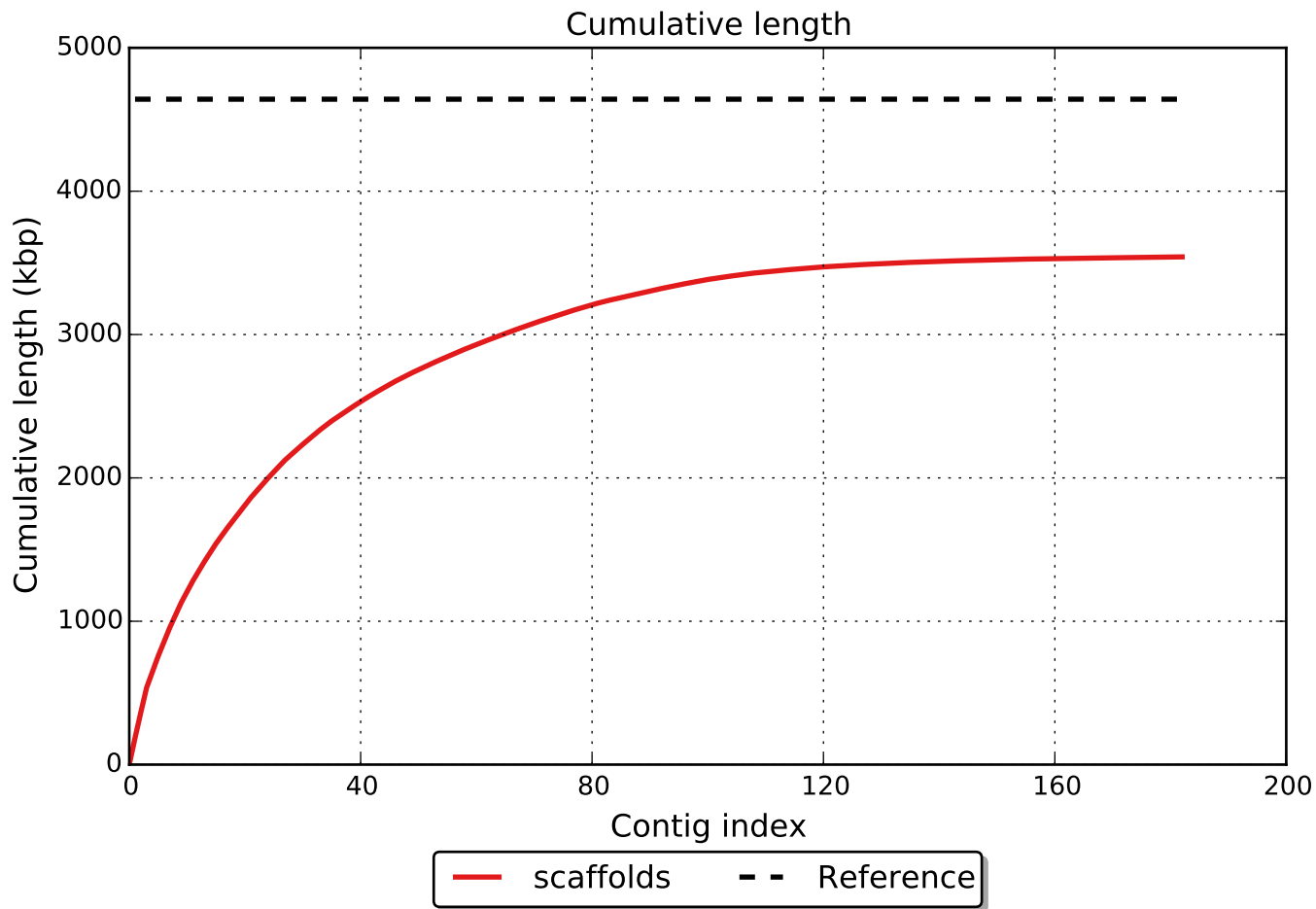


scaffolds

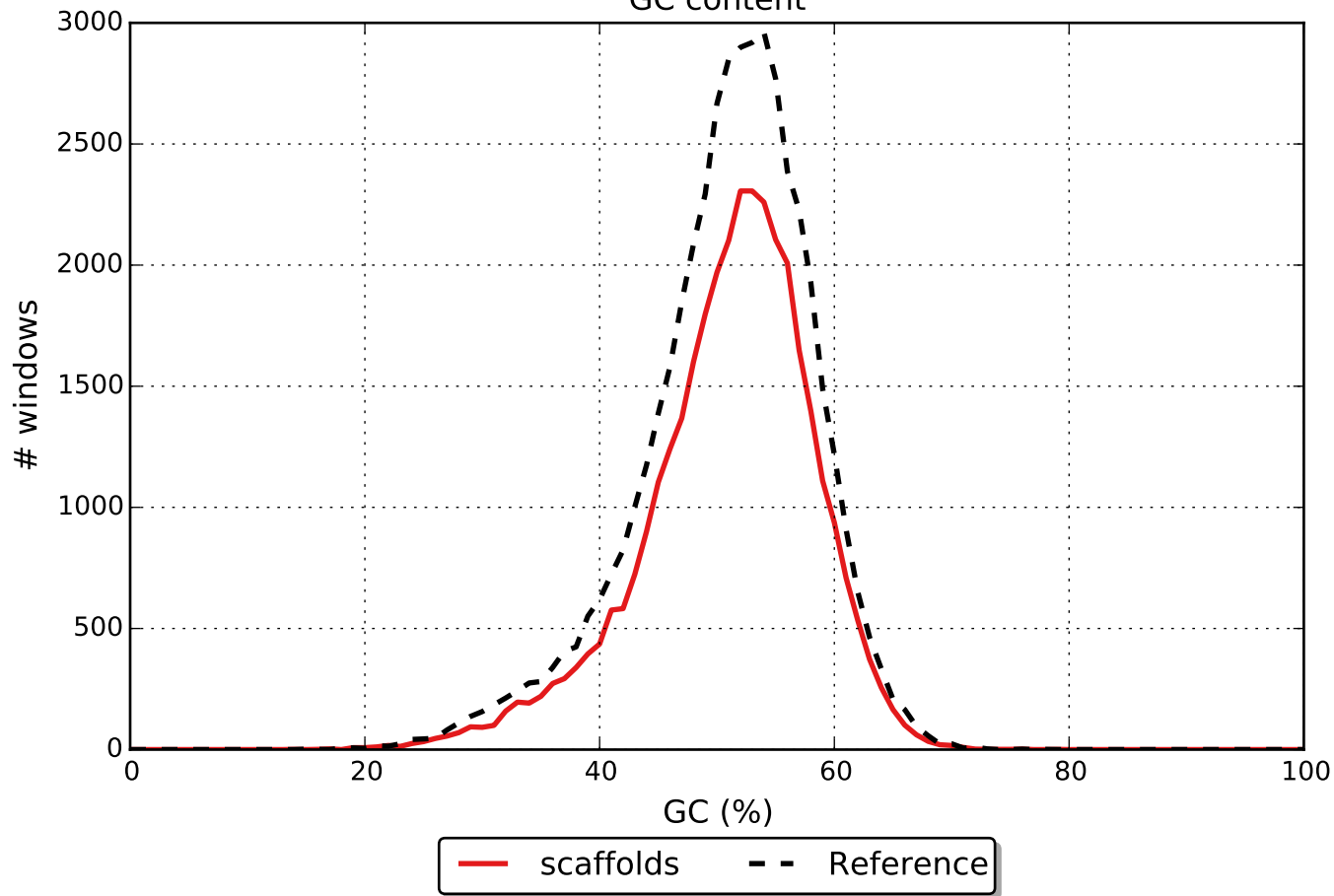
NGx



— scaffolds

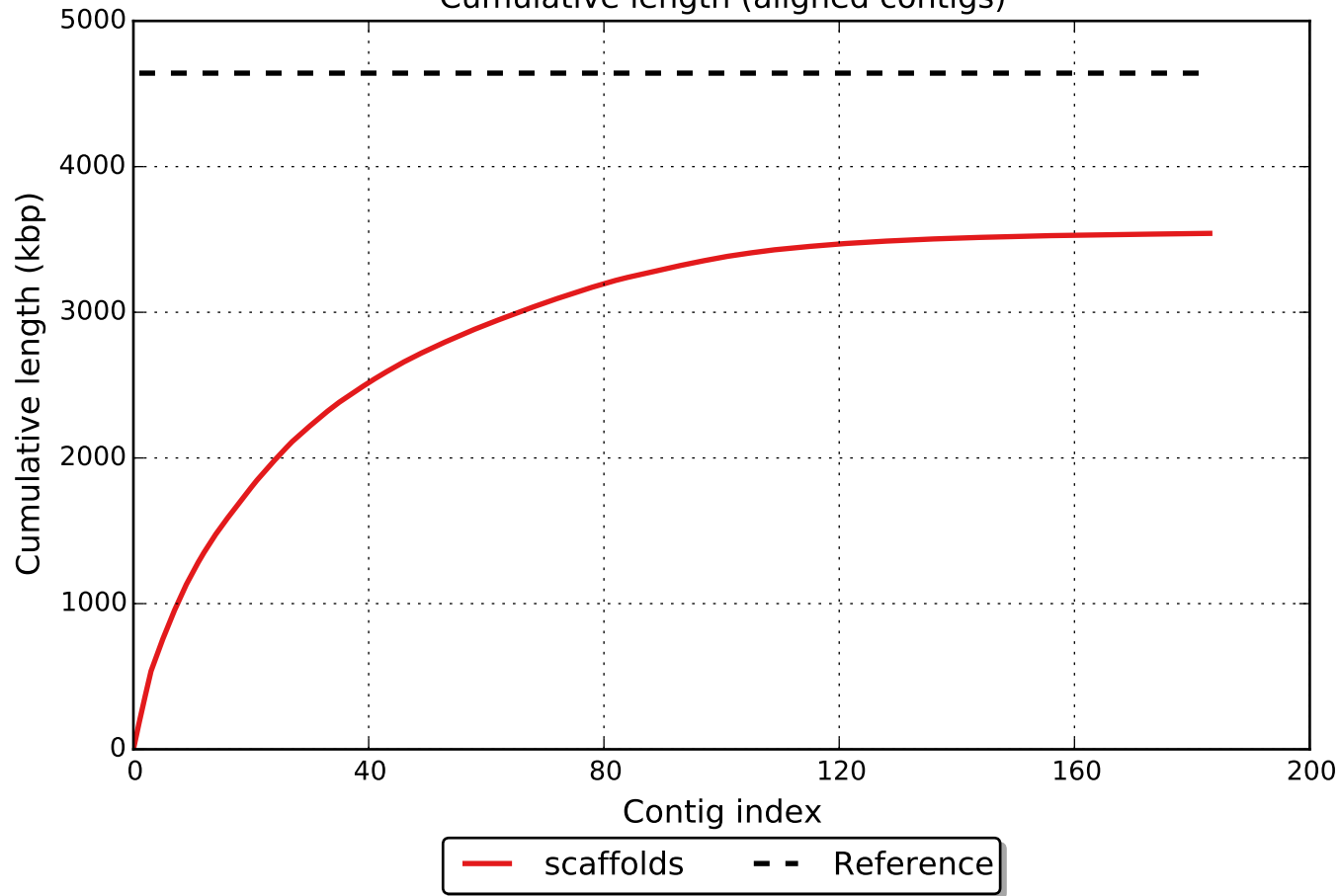


GC content

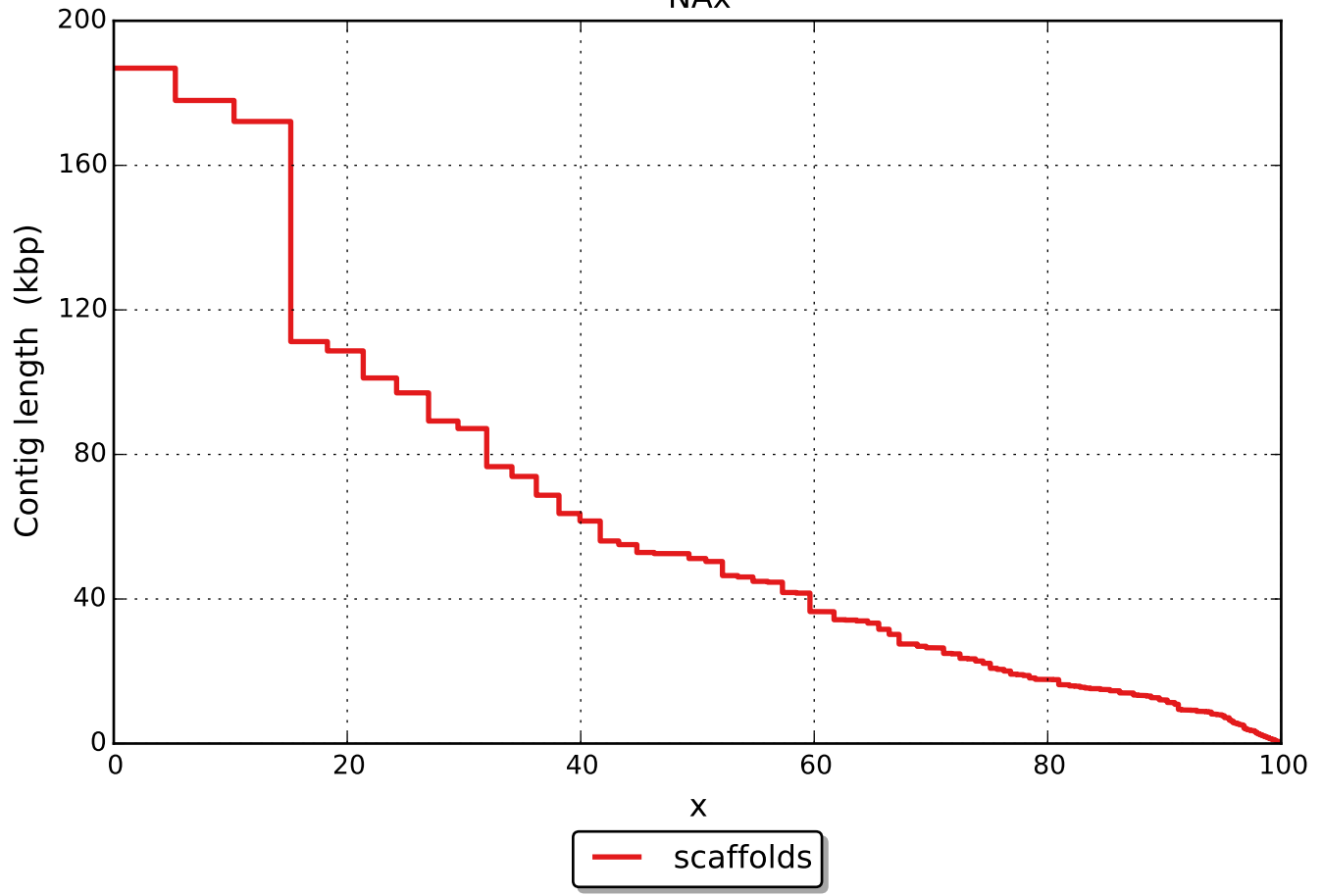




Cumulative length (aligned contigs)



NAx



NGAx

