

Report

	scaffolds
# contigs (≥ 1000 bp)	118
# contigs (≥ 5000 bp)	92
# contigs (≥ 10000 bp)	87
# contigs (≥ 25000 bp)	75
# contigs (≥ 50000 bp)	58
Total length (≥ 1000 bp)	9007220
Total length (≥ 5000 bp)	8960493
Total length (≥ 10000 bp)	8921146
Total length (≥ 25000 bp)	8704633
Total length (≥ 50000 bp)	8130674
# contigs	137
Largest contig	527156
Total length	9019919
Reference length	4641652
GC (%)	50.78
Reference GC (%)	50.78
N50	132337
NG50	263375
N75	86976
NG75	200796
L50	19
LG50	7
L75	39
LG75	13
# misassemblies	268
# misassembled contigs	42
Misassembled contigs length	4251762
# local misassemblies	4
# unaligned contigs	0 + 2 part
Unaligned length	7180
Genome fraction (%)	98.671
Duplication ratio	1.968
# N's per 100 kbp	0.00
# mismatches per 100 kbp	853.00
# indels per 100 kbp	1.24
Largest alignment	527156
NA50	61543
NGA50	197358
NA75	20909
NGA75	89865
LA50	33
LGA50	8
LA75	98
LGA75	19

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	268
# relocations	222
# translocations	0
# inversions	46
# possibly misassembled contigs	13
# misassembled contigs	42
Misassembled contigs length	4251762
# local misassemblies	4
# mismatches	39067
# indels	57
# short indels	57
# long indels	0
Indels length	58

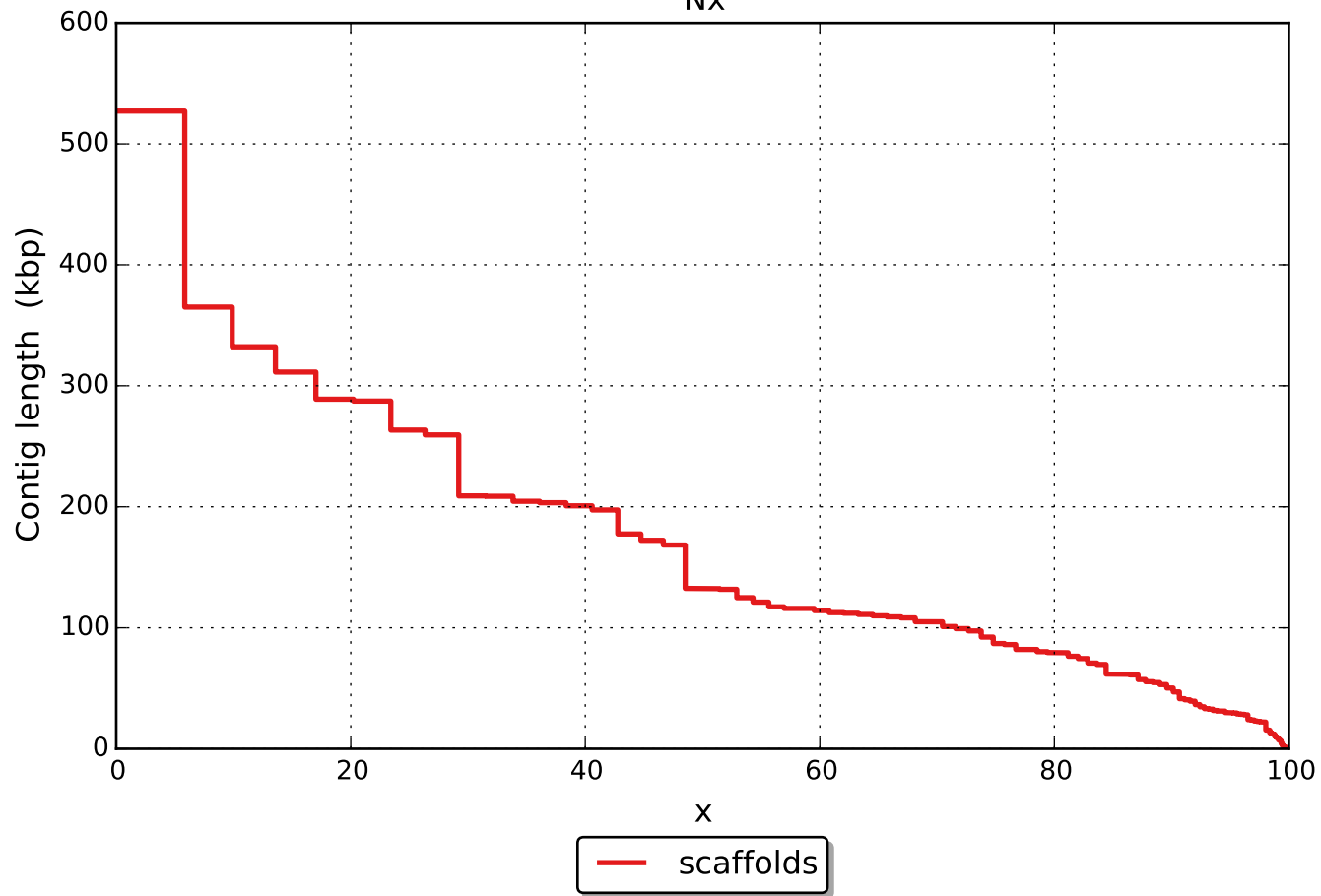
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

Unaligned report

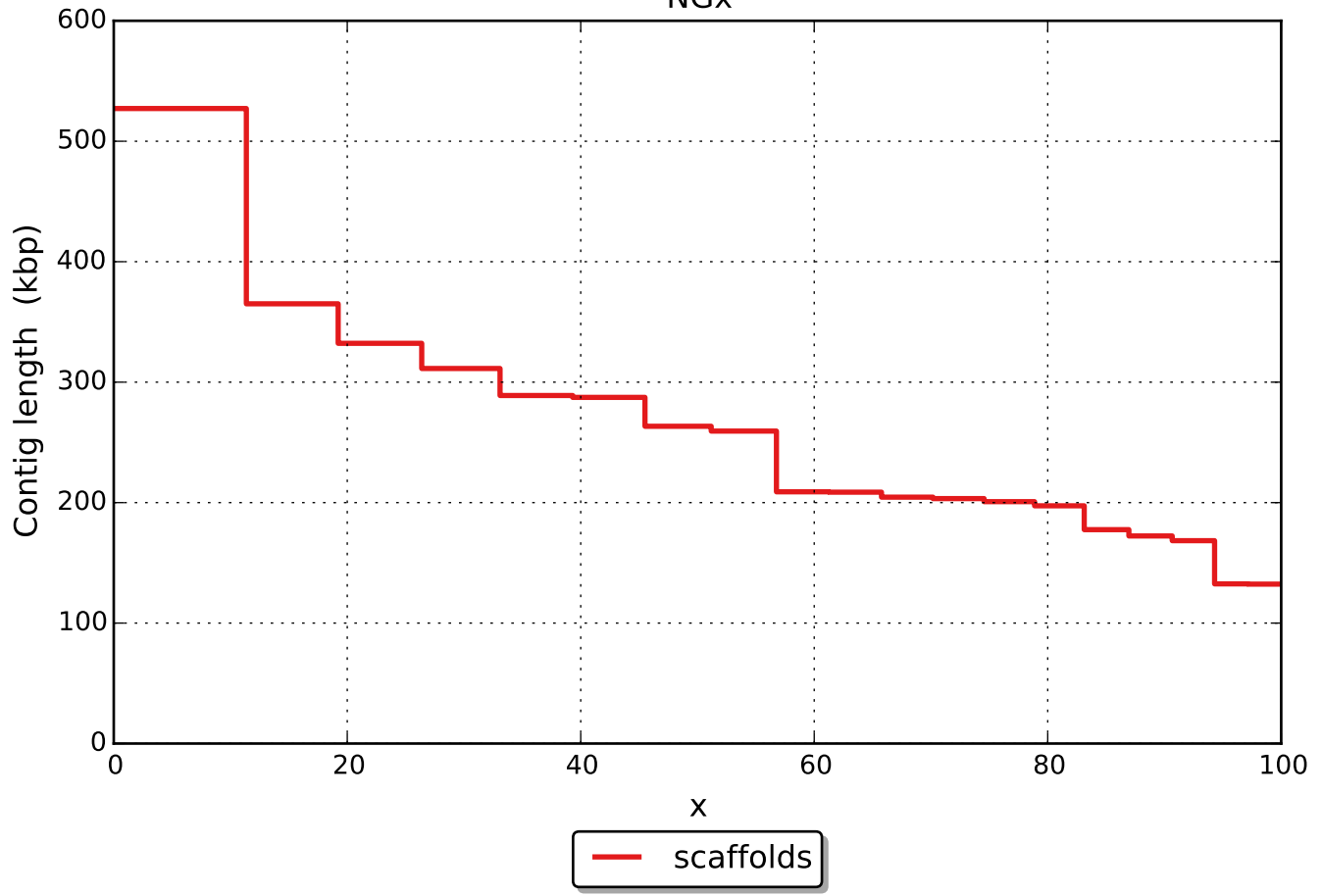
	scaffolds
# fully unaligned contigs	0
Fully unaligned length	0
# partially unaligned contigs	2
# with misassembly	0
# both parts are significant	1
Partially unaligned length	7180
# N's	0

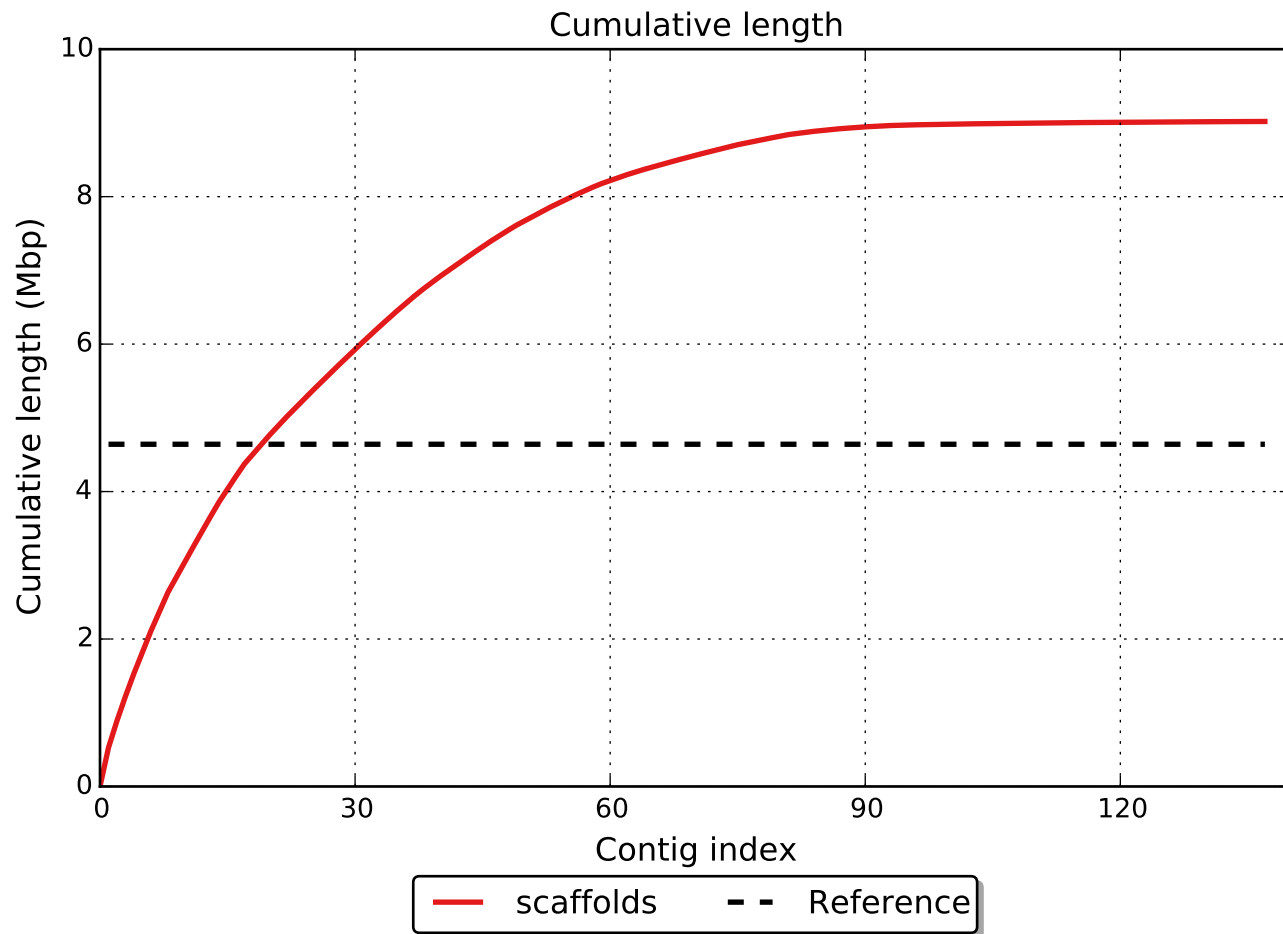
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

Nx

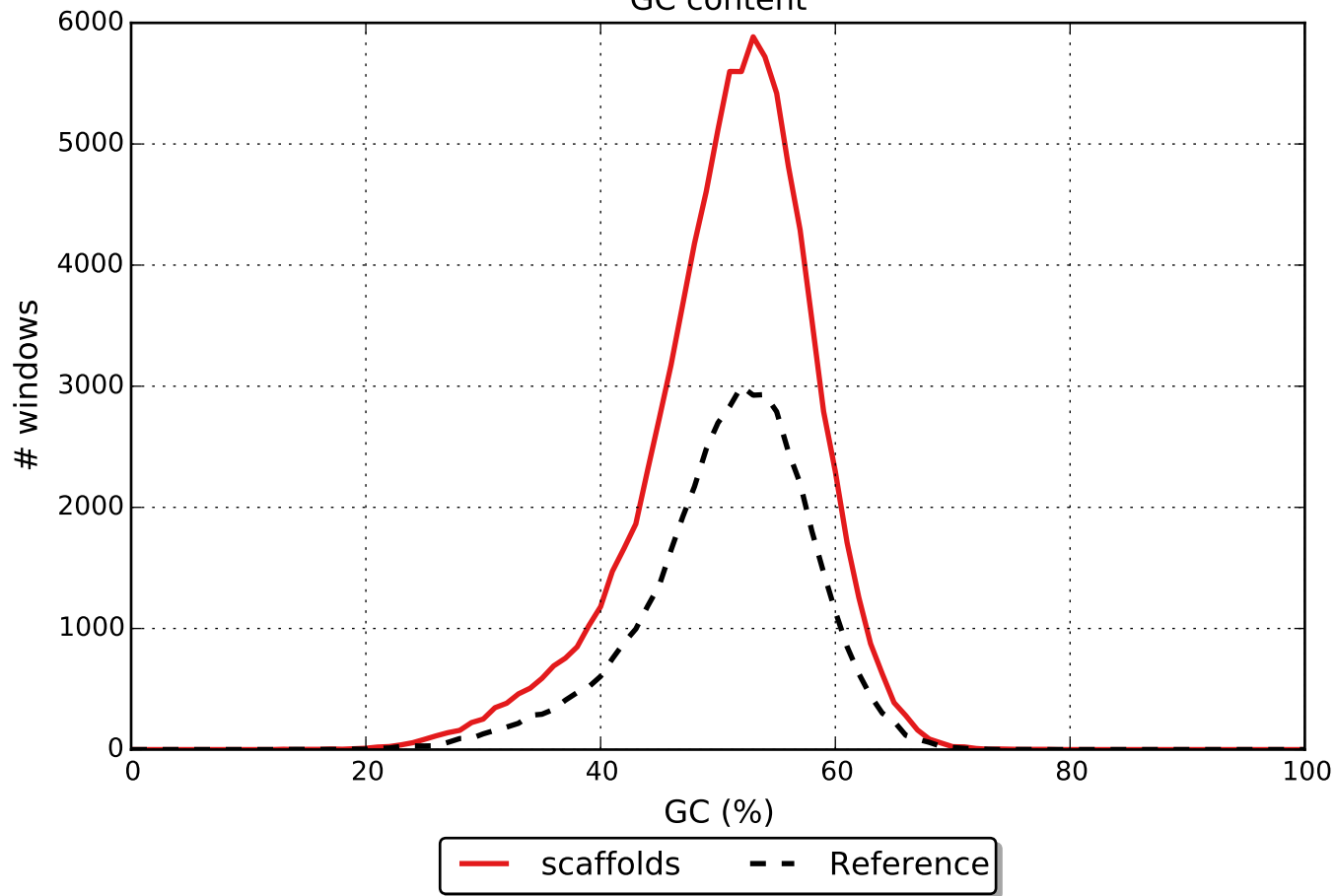


NGx

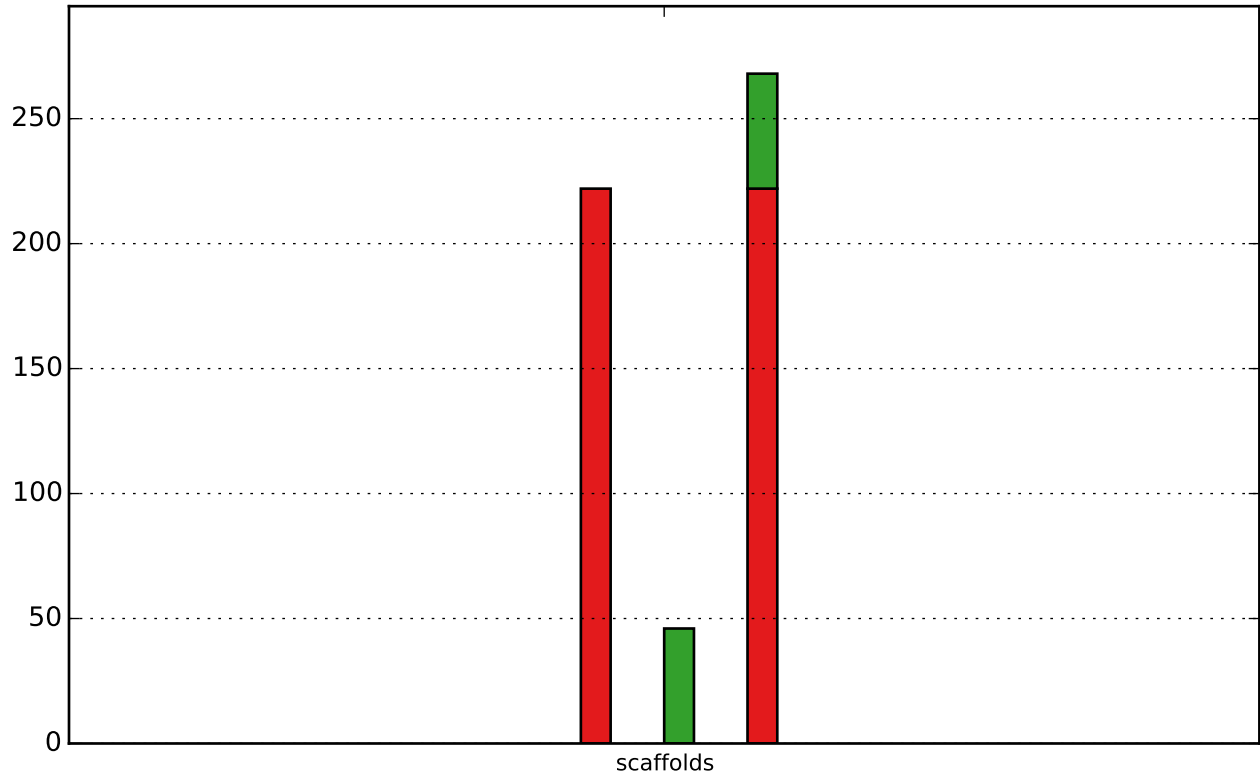


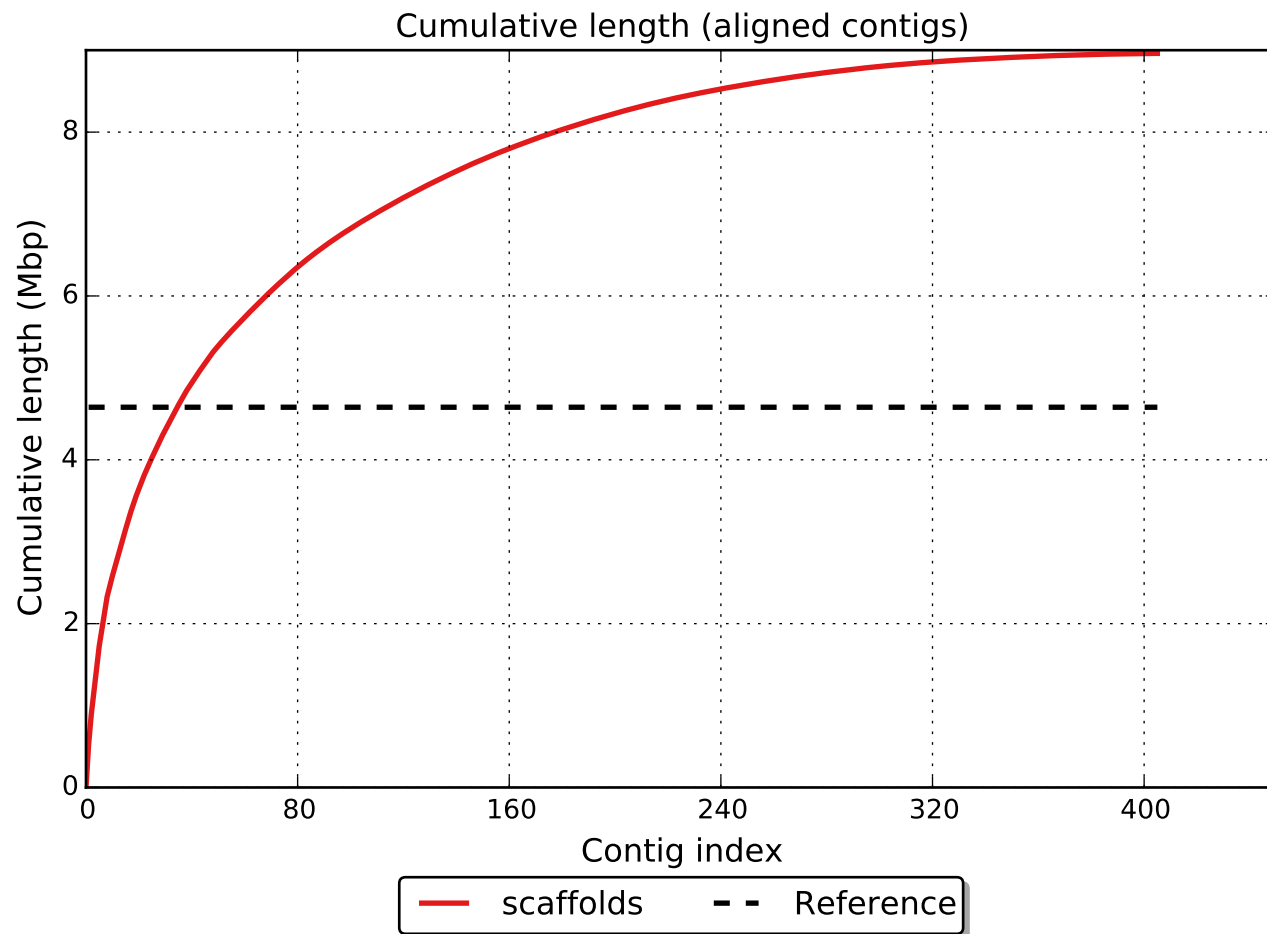


GC content

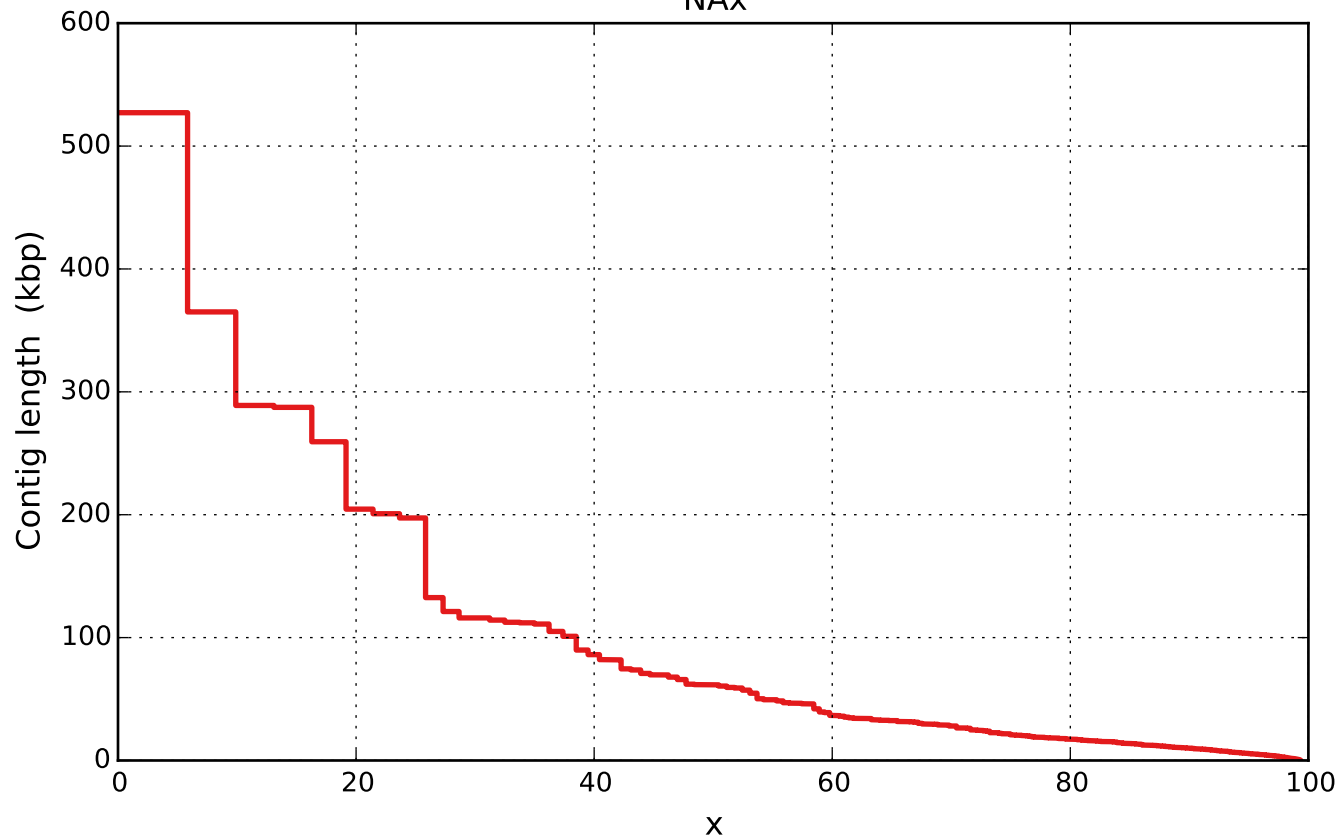


Misassemblies





NAx



— scaffolds

NGAx

