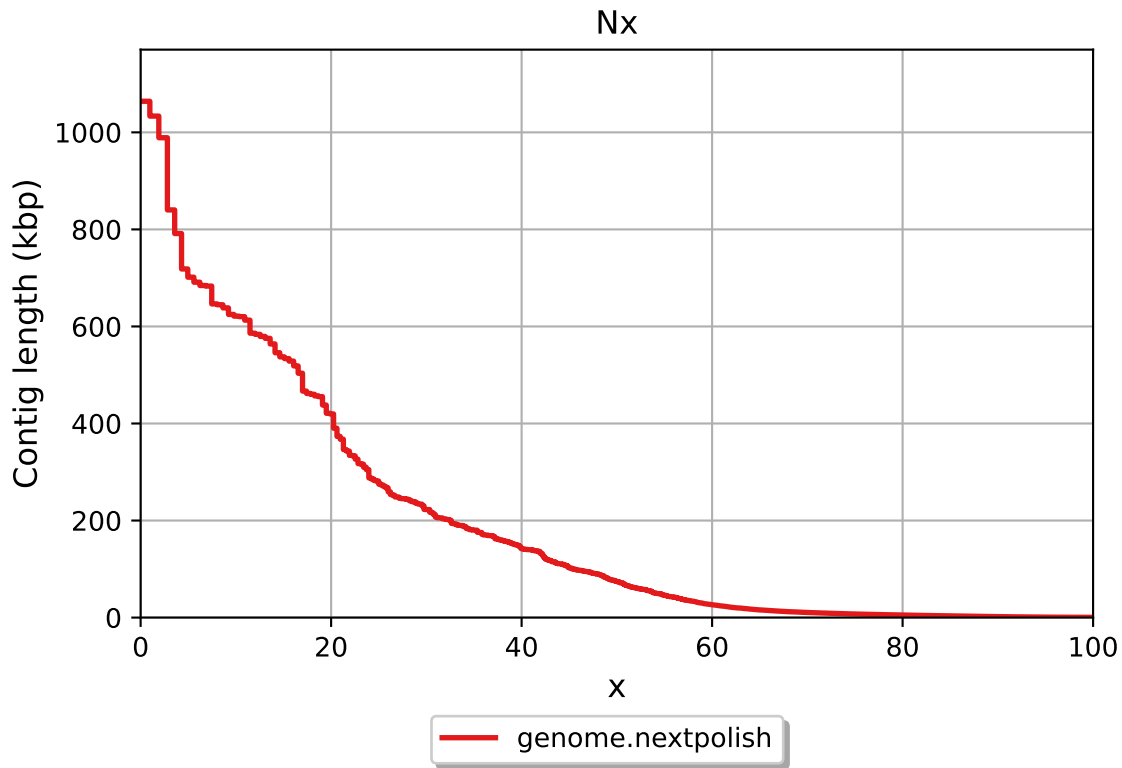


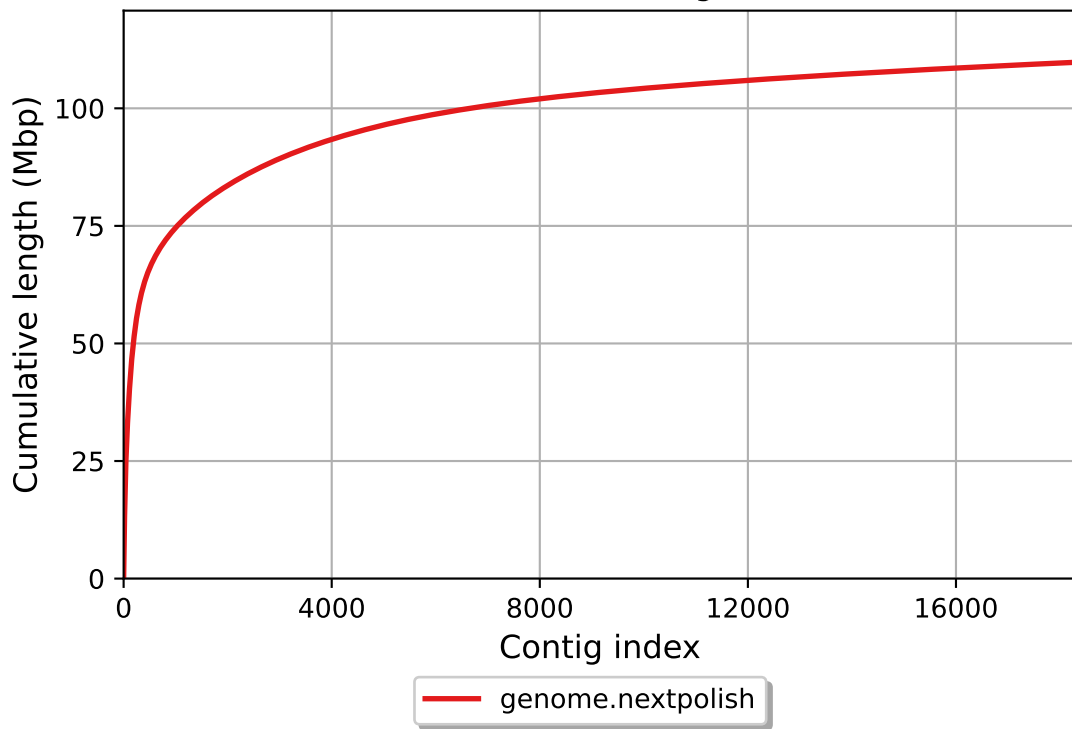
Report

	genome.nextpolish
# contigs (>= 0 bp)	82795
# contigs (>= 1000 bp)	9633
# contigs (>= 5000 bp)	2855
# contigs (>= 10000 bp)	1272
# contigs (>= 25000 bp)	520
# contigs (>= 50000 bp)	317
Total length (>= 0 bp)	127525458
Total length (>= 1000 bp)	103887287
Total length (>= 5000 bp)	88611101
Total length (>= 10000 bp)	77630237
Total length (>= 25000 bp)	66546843
Total length (>= 50000 bp)	59475757
# contigs	18309
Largest contig	1064071
Total length	109791400
GC (%)	43.44
N50	74541
N90	2002
auN	197932.3
L50	242
L90	6018
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	70 + 15 part

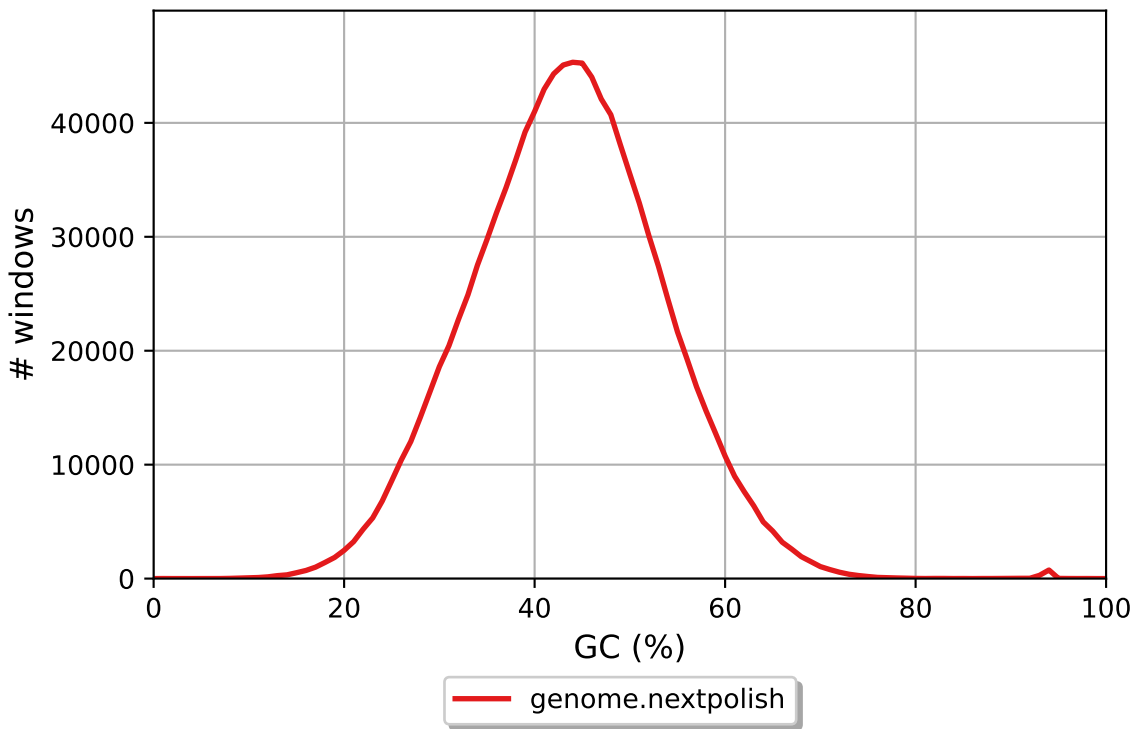
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



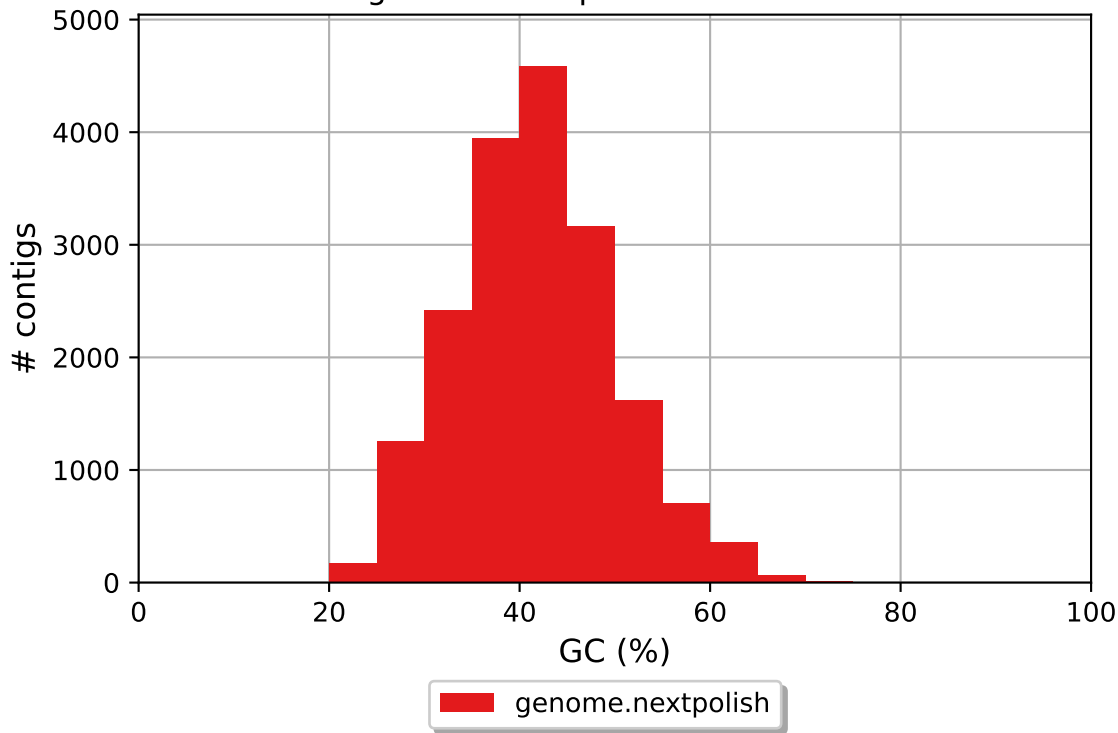
Cumulative length



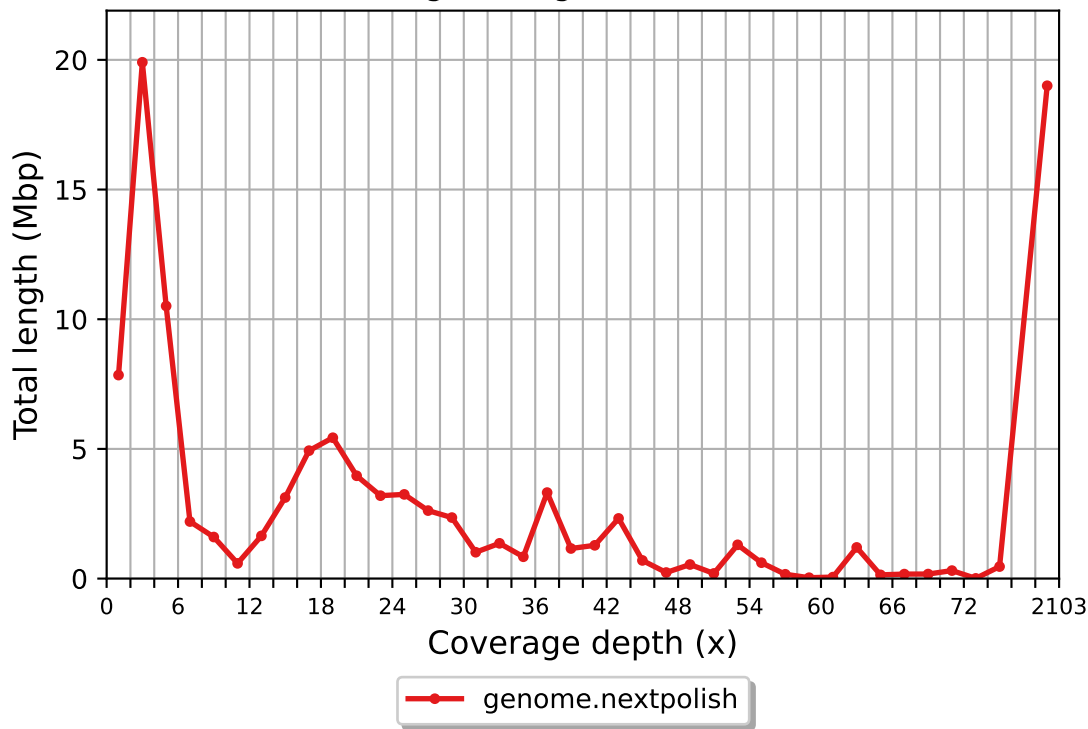
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 2x)



genome.nextpolish coverage histogram (bin size: 2x)

