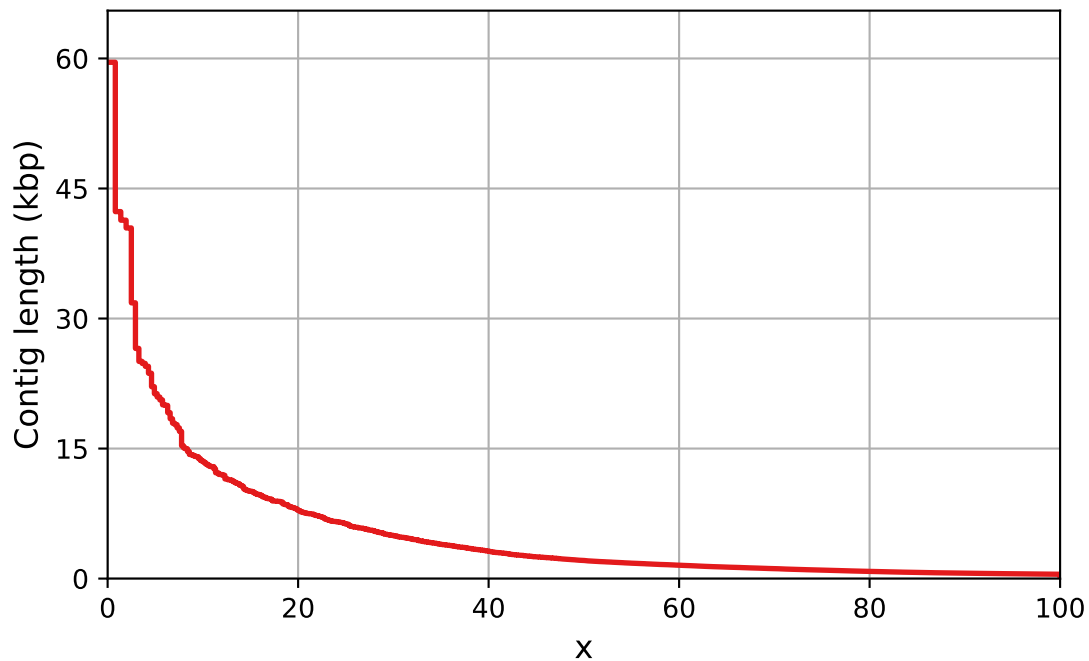


Report

	genome.nextpolish
# contigs (>= 0 bp)	69696
# contigs (>= 1000 bp)	1977
# contigs (>= 5000 bp)	222
# contigs (>= 10000 bp)	67
# contigs (>= 25000 bp)	7
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	24082943
Total length (>= 1000 bp)	5482153
Total length (>= 5000 bp)	2197851
Total length (>= 10000 bp)	1124056
Total length (>= 25000 bp)	267104
Total length (>= 50000 bp)	59560
# contigs	4802
Largest contig	59560
Total length	7377175
GC (%)	61.01
N50	2087
N90	627
auN	5534.8
L50	706
L90	3475
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 4 part

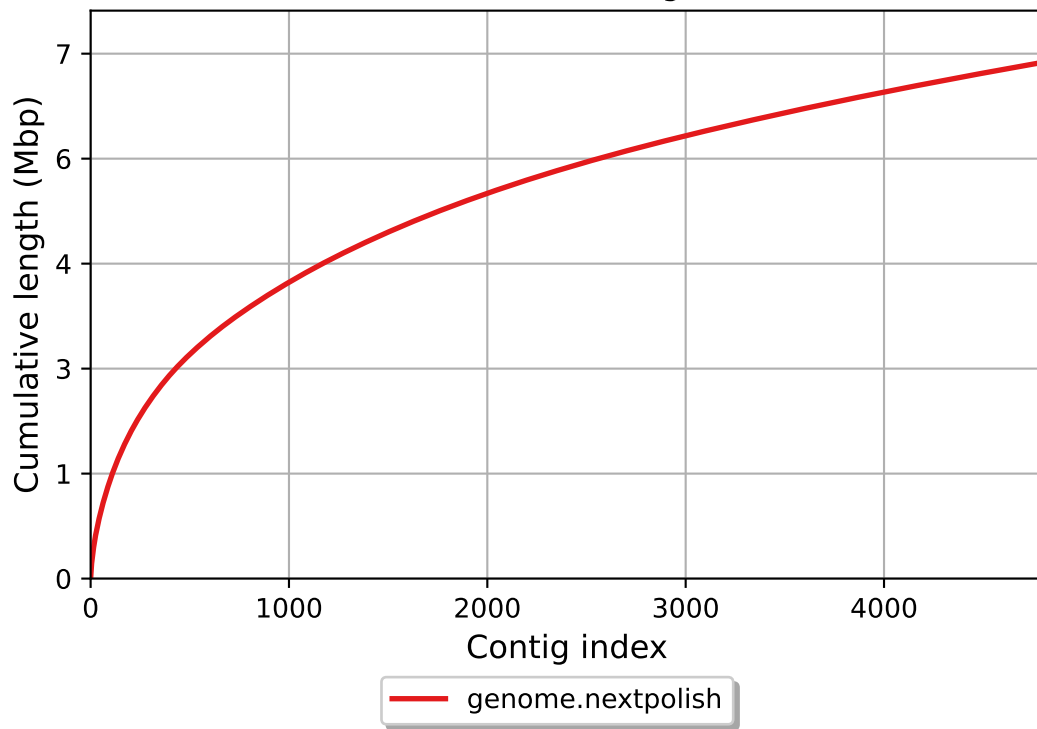
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

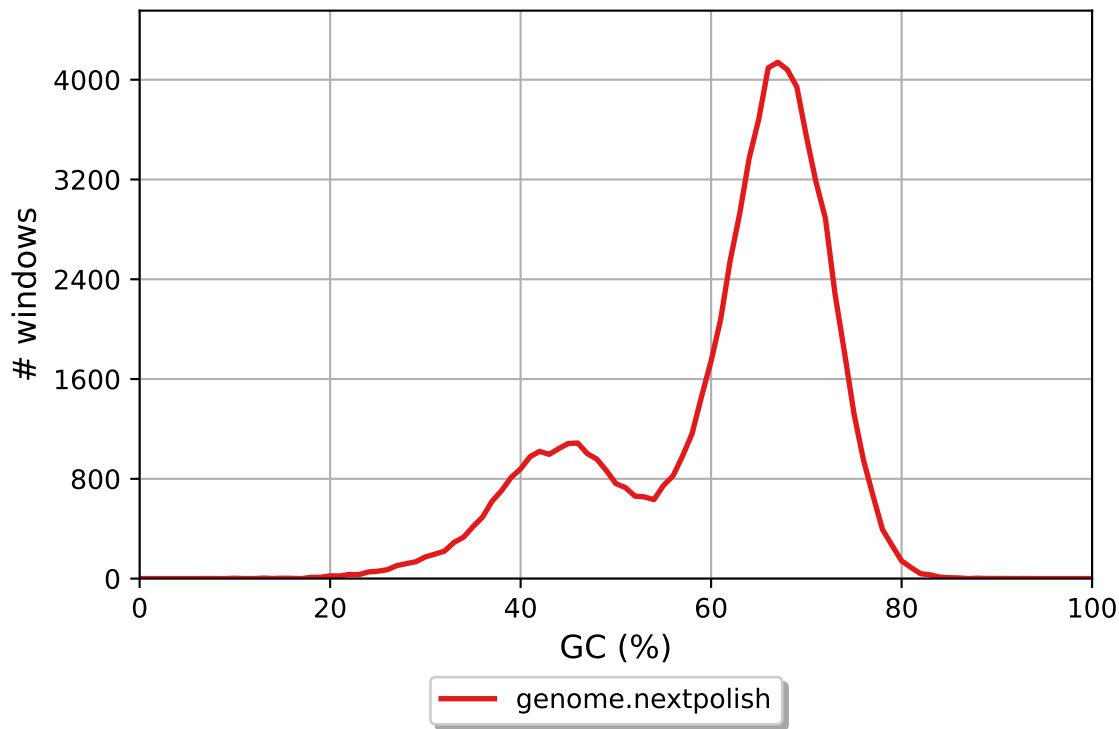


— genome.nextpolish

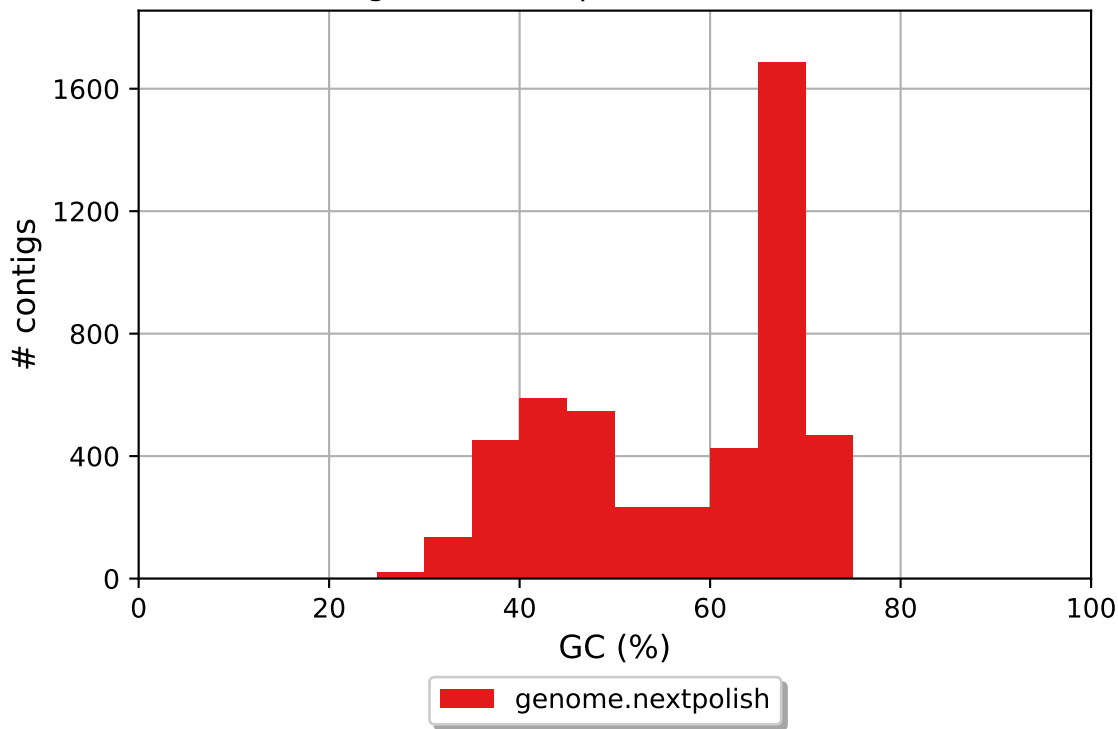
Cumulative length



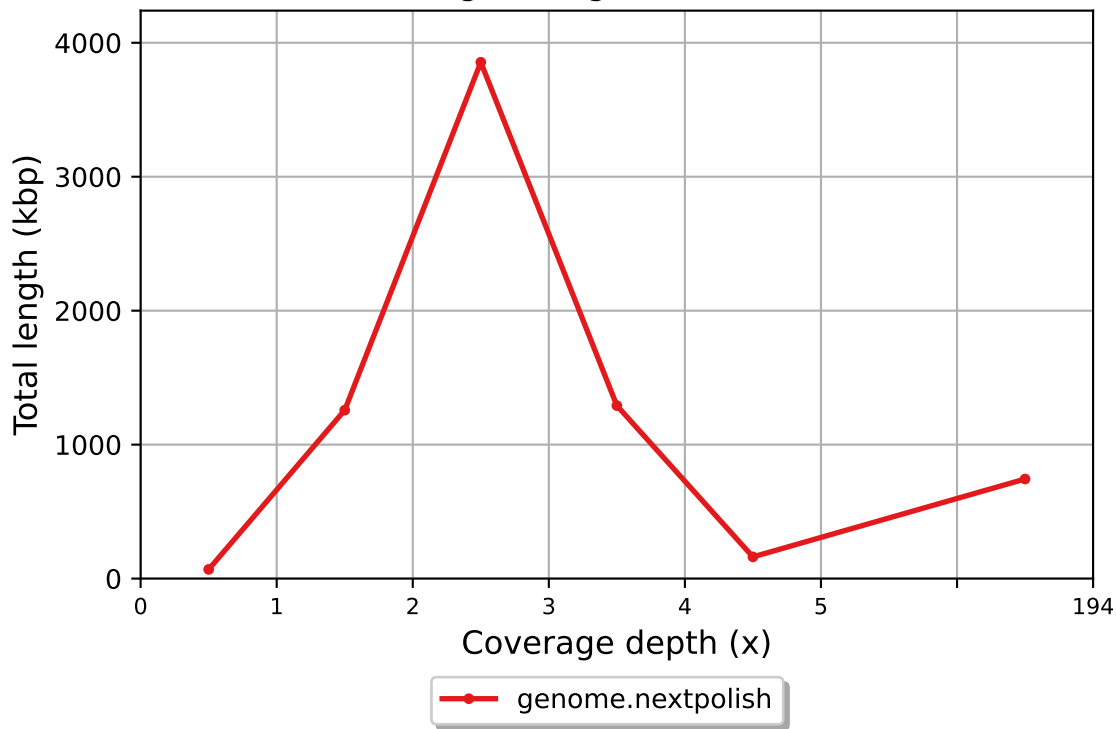
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



genome.nextpolish coverage histogram (bin size: 1x)

