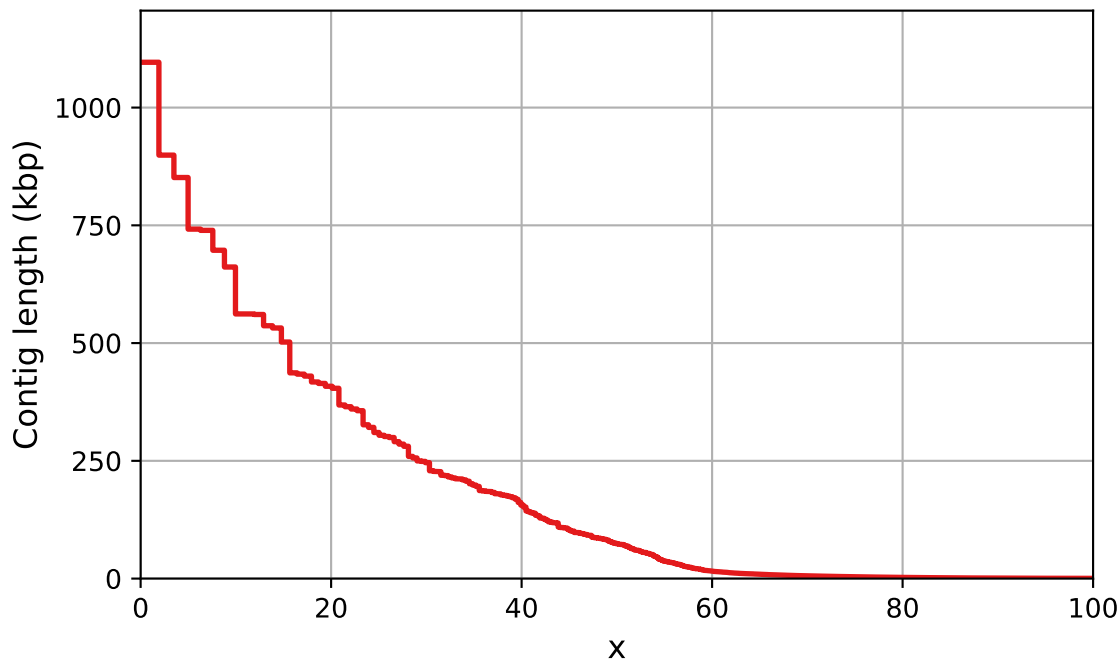


Report

	genome.nextpolish
# contigs (>= 0 bp)	86421
# contigs (>= 1000 bp)	6205
# contigs (>= 5000 bp)	1133
# contigs (>= 10000 bp)	486
# contigs (>= 25000 bp)	216
# contigs (>= 50000 bp)	157
Total length (>= 0 bp)	77214760
Total length (>= 1000 bp)	51460602
Total length (>= 5000 bp)	41057273
Total length (>= 10000 bp)	36618831
Total length (>= 25000 bp)	32749257
Total length (>= 50000 bp)	30731526
# contigs	14630
Largest contig	1096204
Total length	57105645
GC (%)	46.77
N50	74098
N90	1010
auN	200791.3
L50	122
L90	6140
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	44 + 13 part

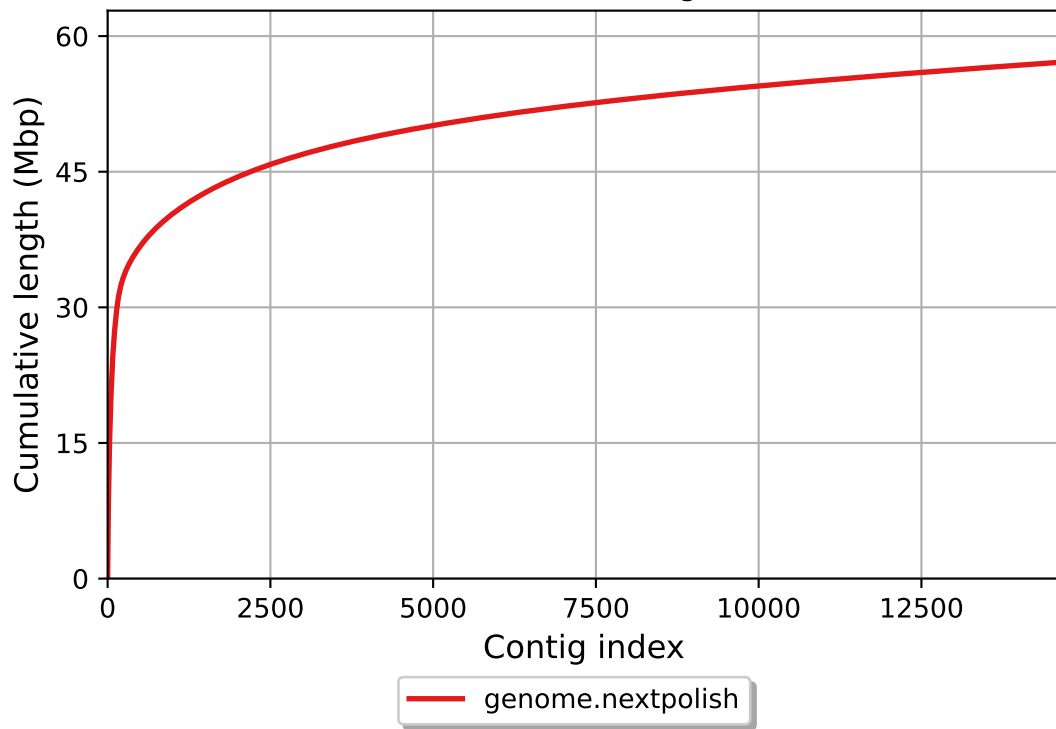
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

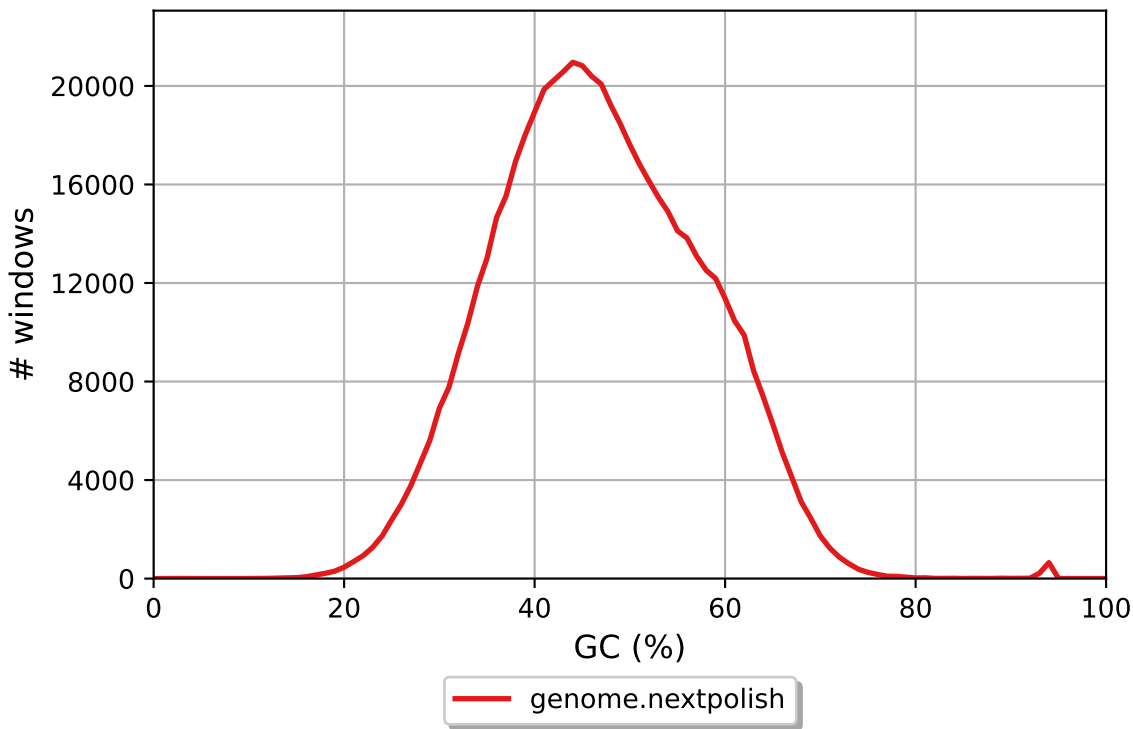


— genome.nextpolish

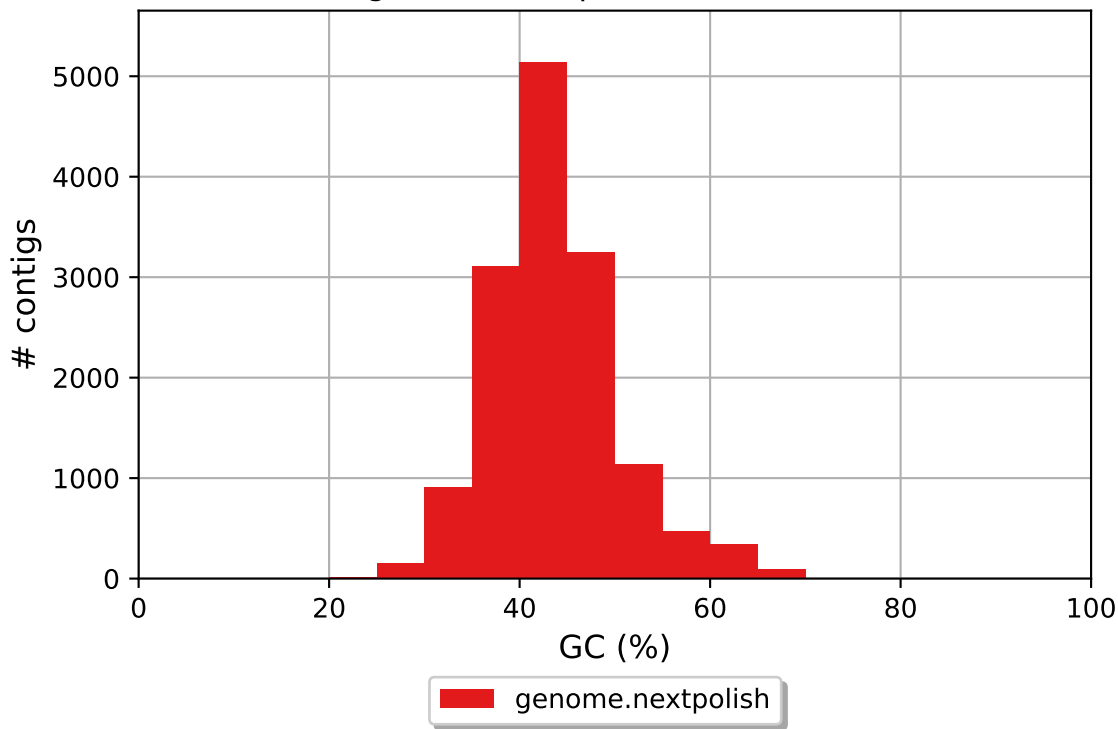
Cumulative length



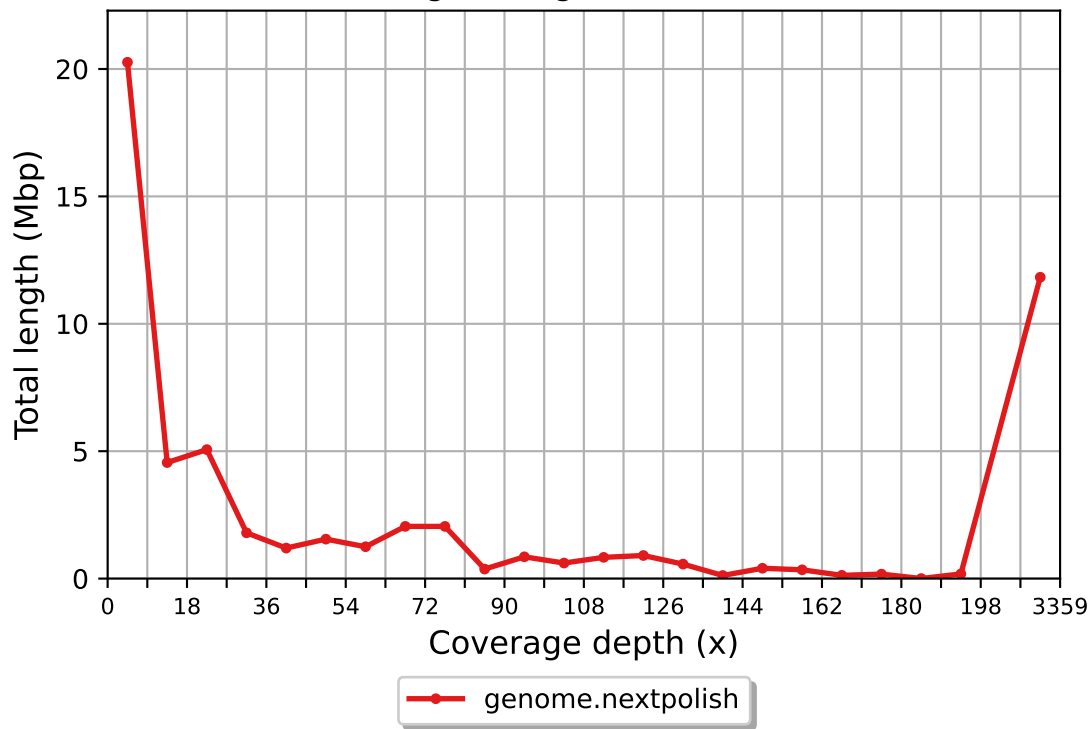
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 9x)



genome.nextpolish coverage histogram (bin size: 9x)

