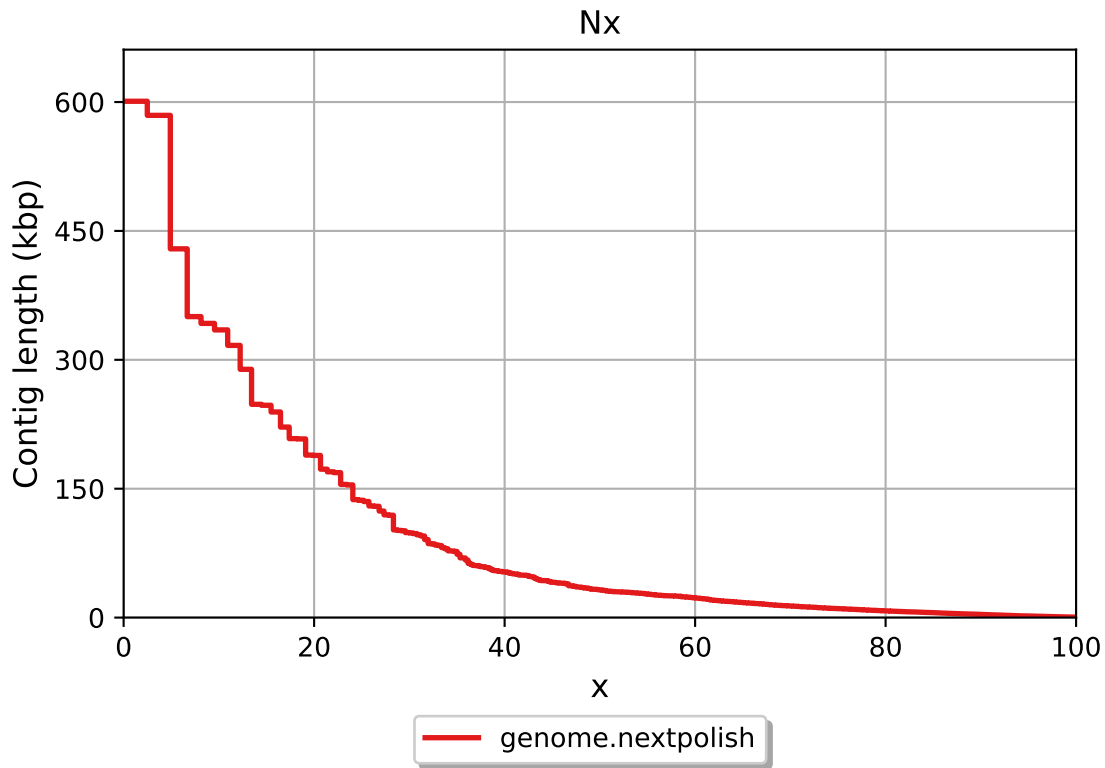


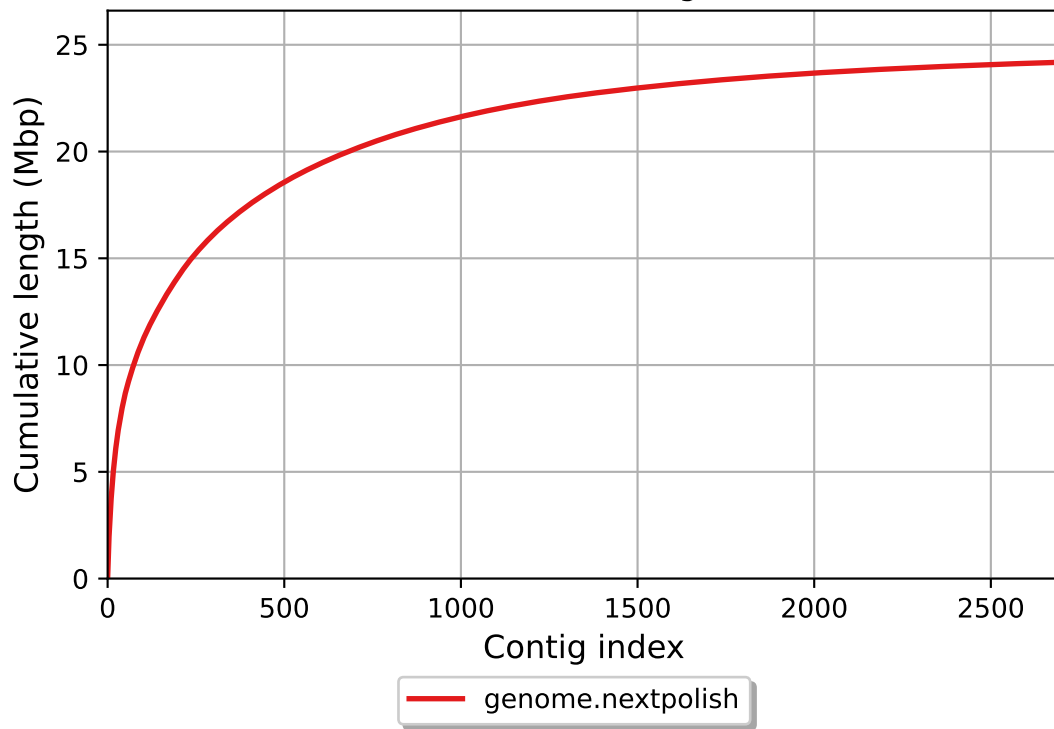
Report

	genome.nextpolish
# contigs (>= 0 bp)	8088
# contigs (>= 1000 bp)	2026
# contigs (>= 5000 bp)	842
# contigs (>= 10000 bp)	476
# contigs (>= 25000 bp)	196
# contigs (>= 50000 bp)	74
Total length (>= 0 bp)	25025729
Total length (>= 1000 bp)	23701307
Total length (>= 5000 bp)	20932757
Total length (>= 10000 bp)	18333491
Total length (>= 25000 bp)	14057338
Total length (>= 50000 bp)	10028363
# contigs	2696
Largest contig	600893
Total length	24180644
GC (%)	43.06
N50	32186
N90	3710
auN	105551.0
L50	126
L90	1035
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 2 part

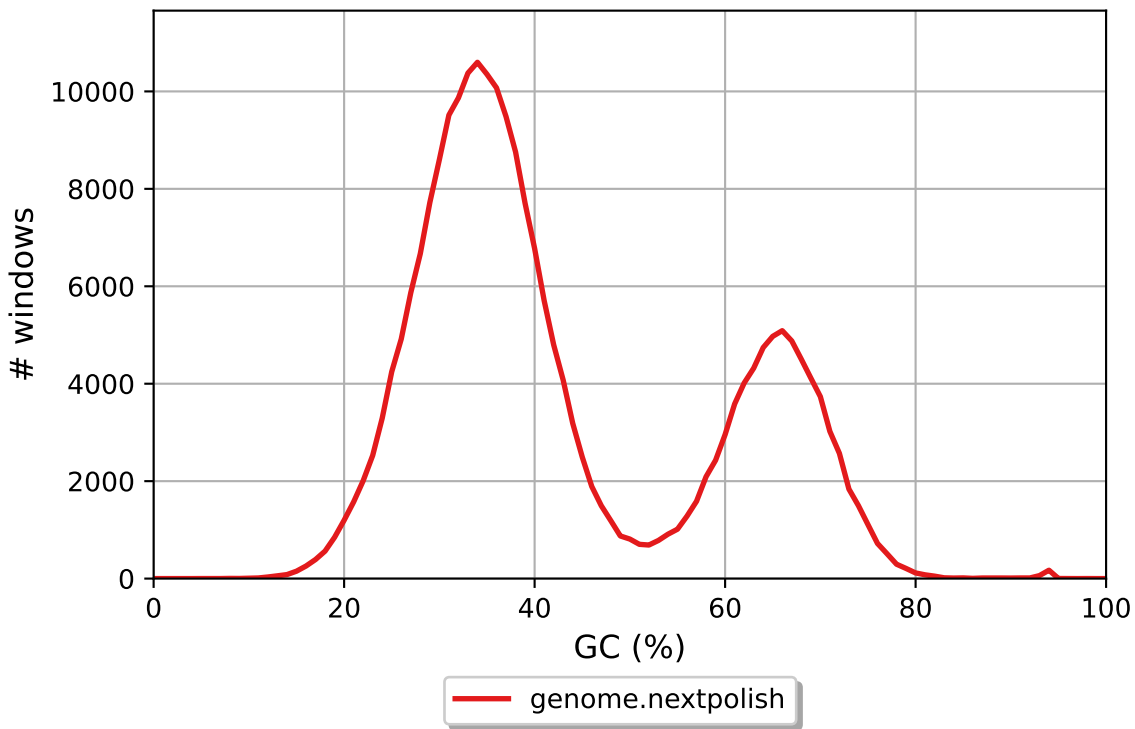
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



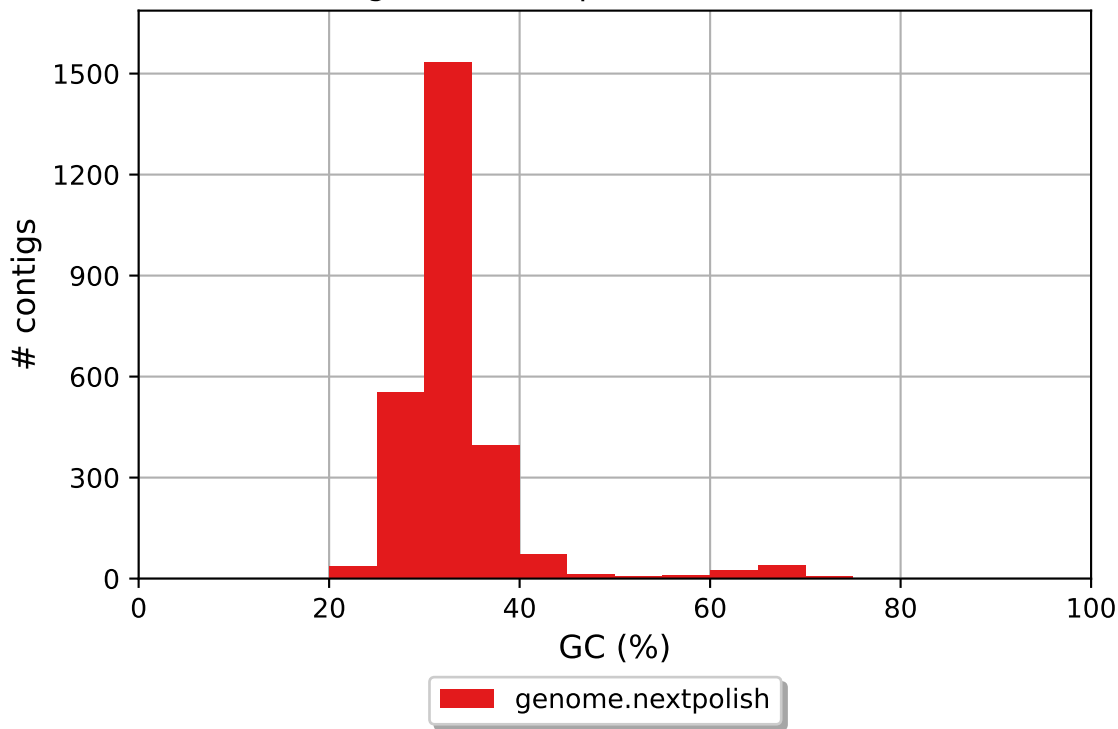
Cumulative length



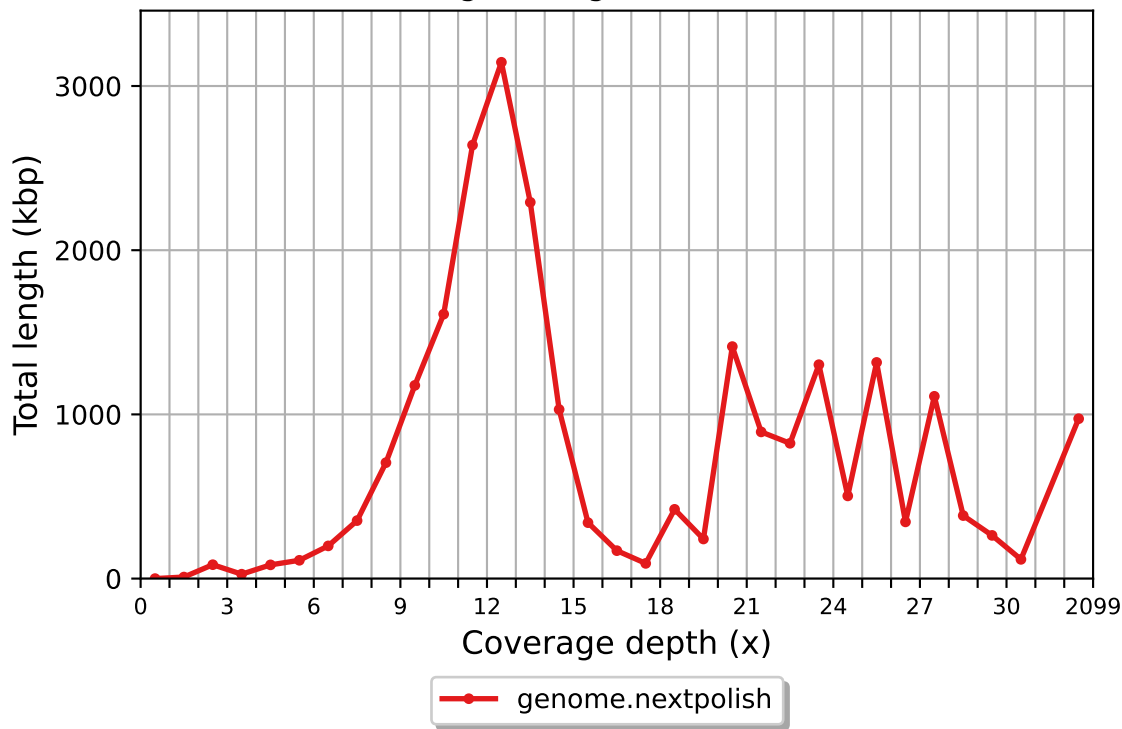
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

