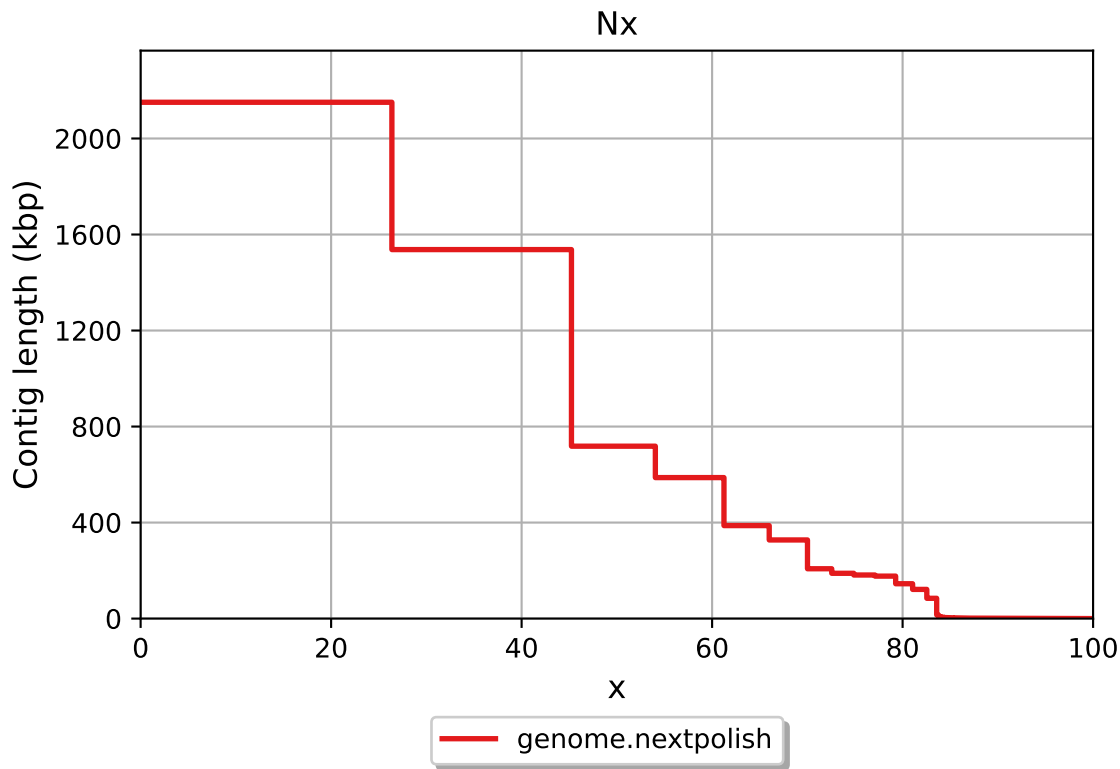


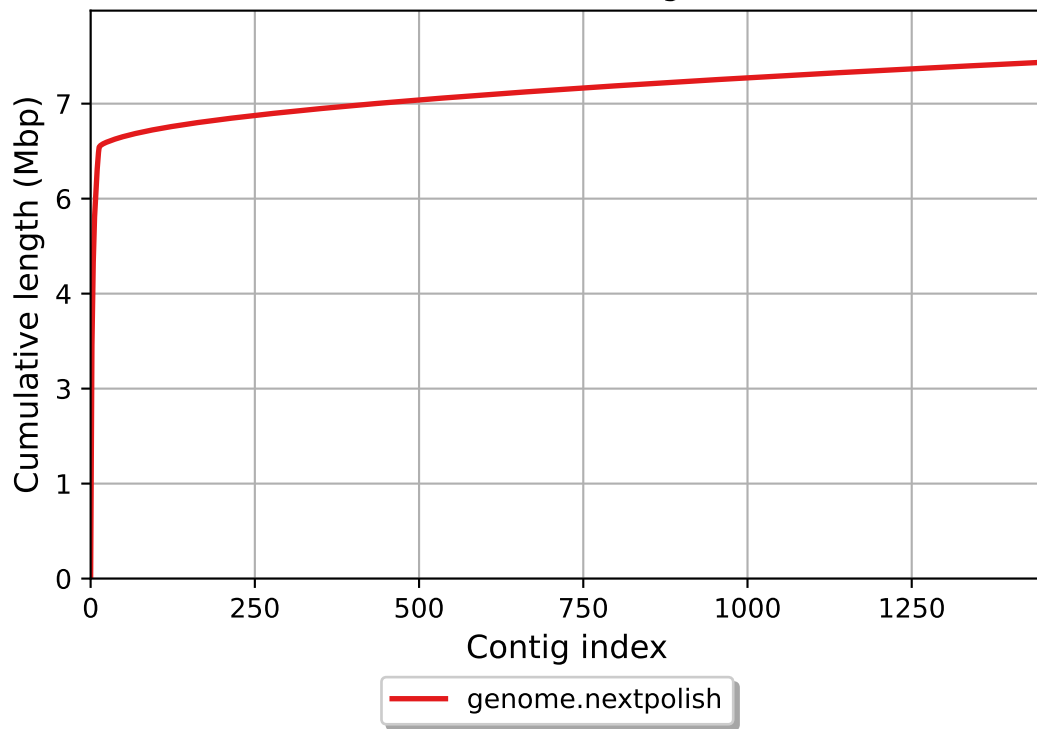
Report

	genome.nextpolish
# contigs (>= 0 bp)	74988
# contigs (>= 1000 bp)	352
# contigs (>= 5000 bp)	21
# contigs (>= 10000 bp)	14
# contigs (>= 25000 bp)	13
# contigs (>= 50000 bp)	13
Total length (>= 0 bp)	26556646
Total length (>= 1000 bp)	7423896
Total length (>= 5000 bp)	6876777
Total length (>= 10000 bp)	6829775
Total length (>= 25000 bp)	6814600
Total length (>= 50000 bp)	6814600
# contigs	1450
Largest contig	2150911
Total length	8153543
GC (%)	62.23
N50	718167
N90	1142
auN	1017430.1
L50	3
L90	271
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 1 part

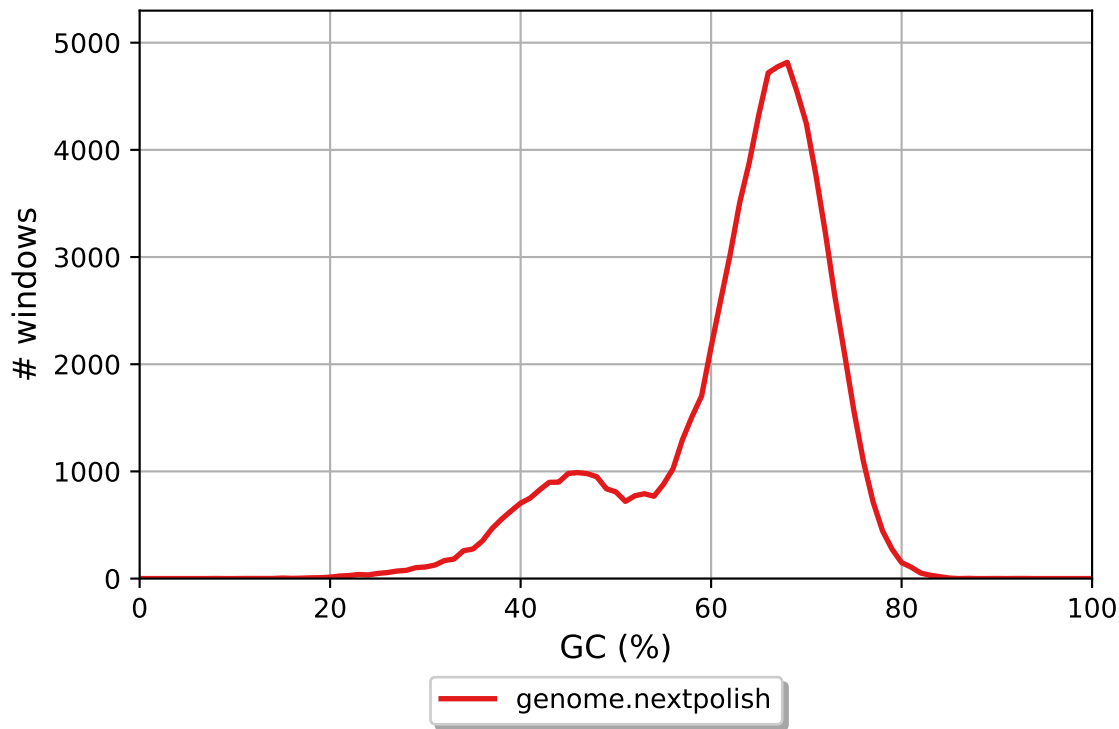
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



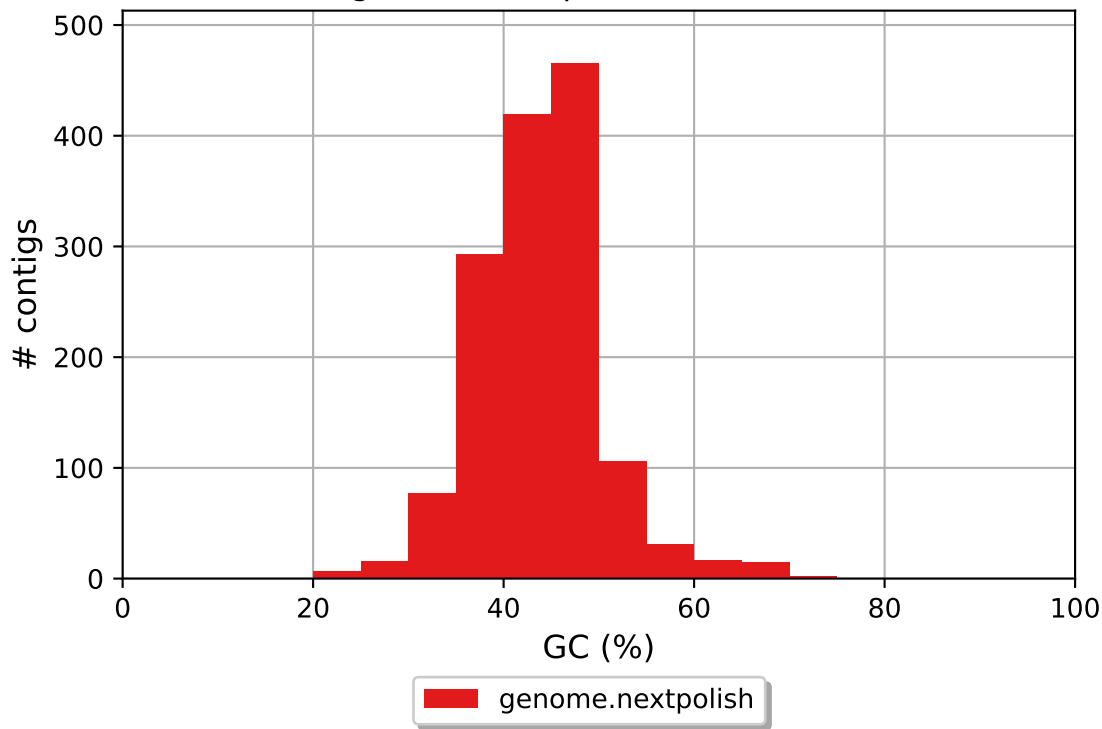
Cumulative length



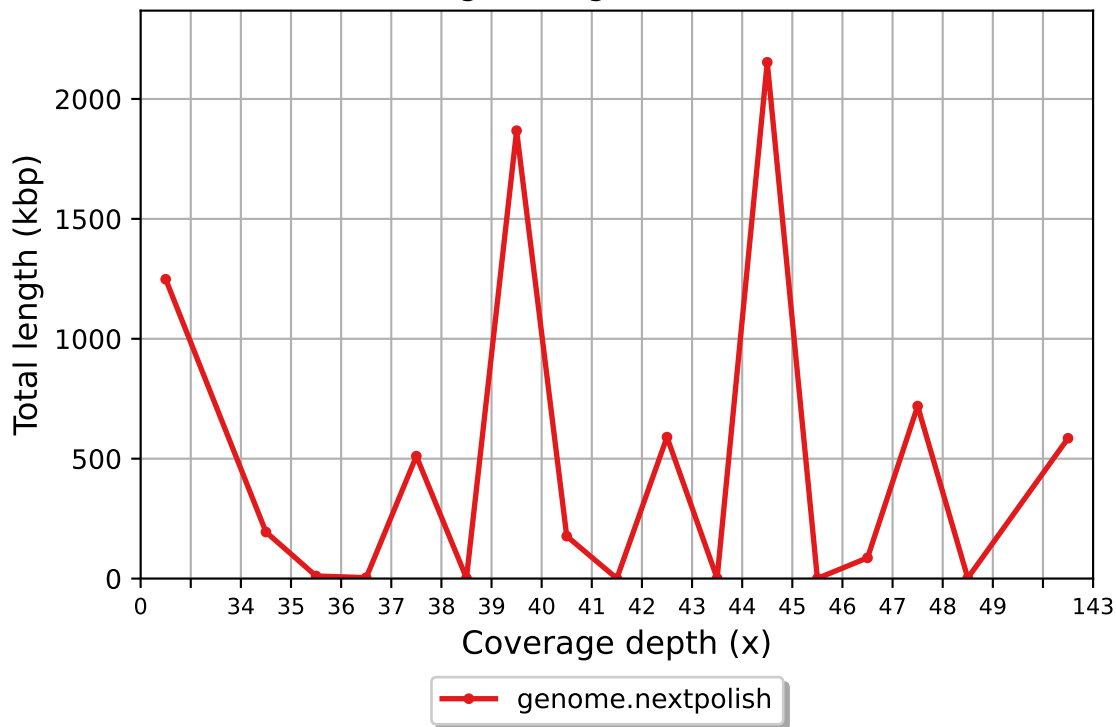
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

