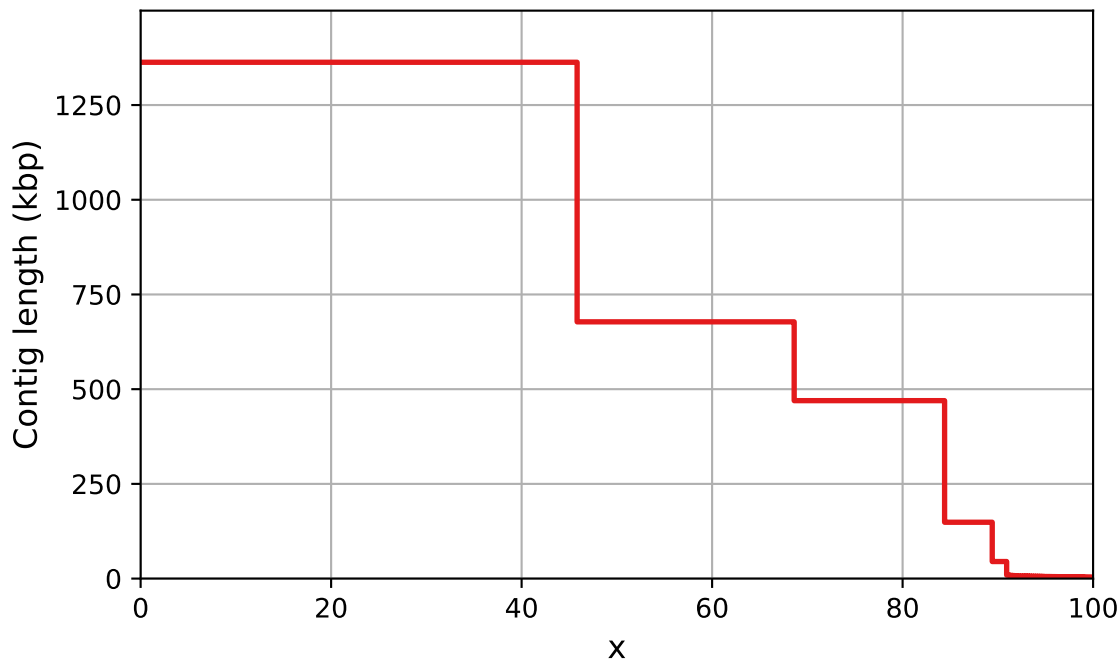


# Report

	genome.nextpolish
# contigs (>= 0 bp)	55
# contigs (>= 1000 bp)	55
# contigs (>= 5000 bp)	34
# contigs (>= 10000 bp)	5
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	4
Total length (>= 0 bp)	2974053
Total length (>= 1000 bp)	2974053
Total length (>= 5000 bp)	2884220
Total length (>= 10000 bp)	2704130
Total length (>= 25000 bp)	2704130
Total length (>= 50000 bp)	2659098
# contigs	55
Largest contig	1362914
Total length	2974053
GC (%)	34.64
N50	677759
N90	45032
auN	861853.5
L50	2
L90	5
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	12 + 10 part

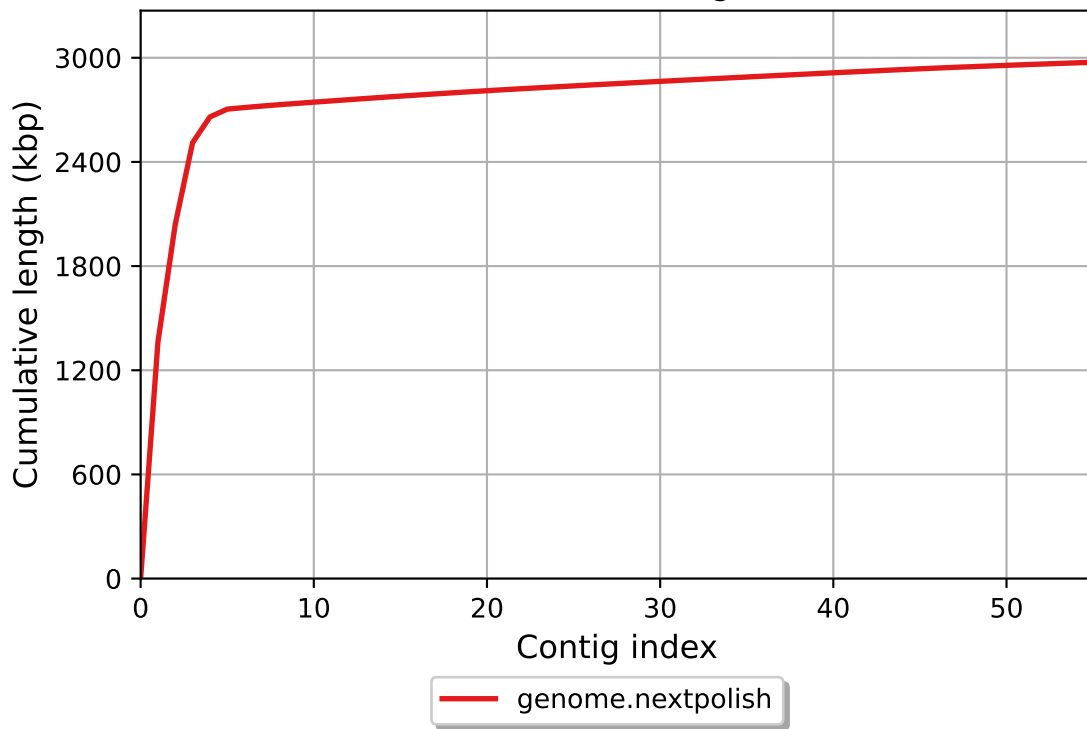
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

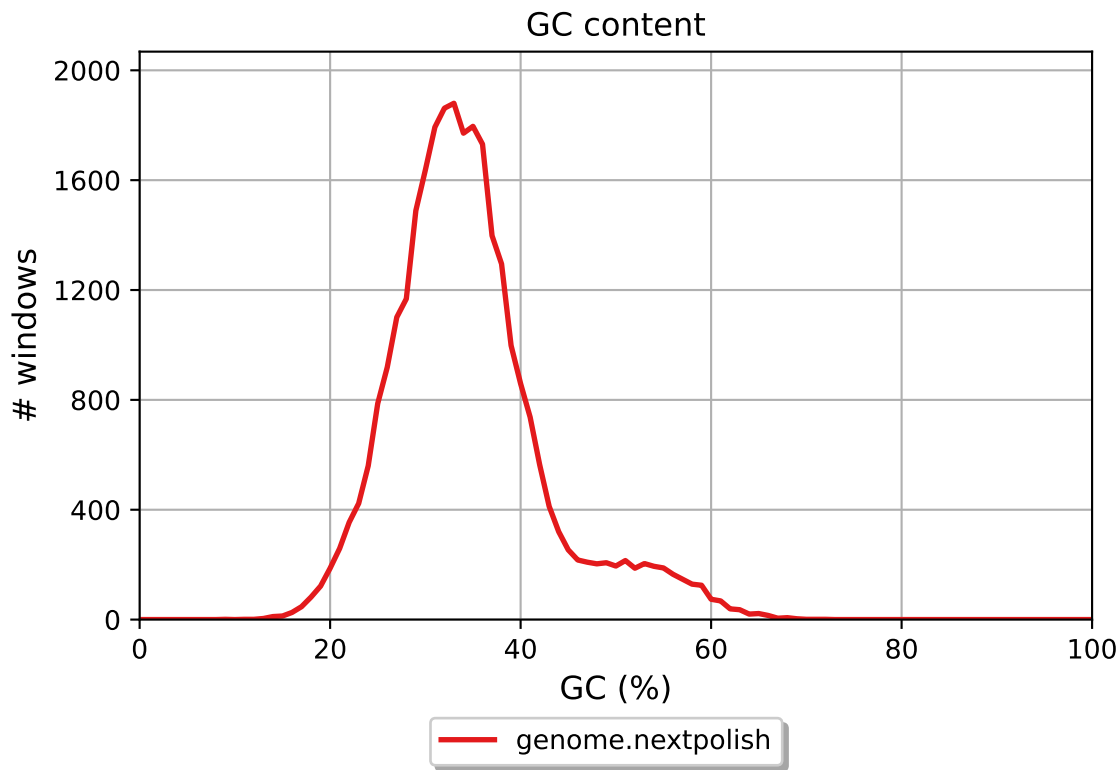
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

