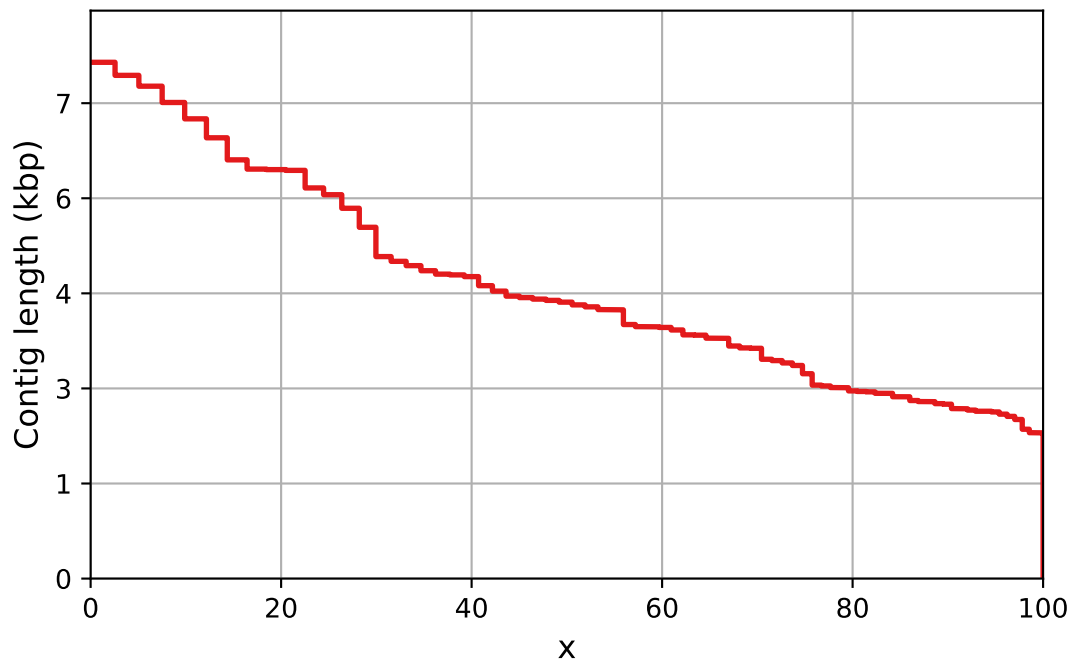


Report

	genome.nextpolish
# contigs (>= 0 bp)	77
# contigs (>= 1000 bp)	77
# contigs (>= 5000 bp)	16
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	317613
Total length (>= 1000 bp)	317613
Total length (>= 5000 bp)	105216
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	77
Largest contig	8145
Total length	317613
GC (%)	41.42
N50	4361
N90	2751
auN	4665.6
L50	28
L90	65
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 2 part

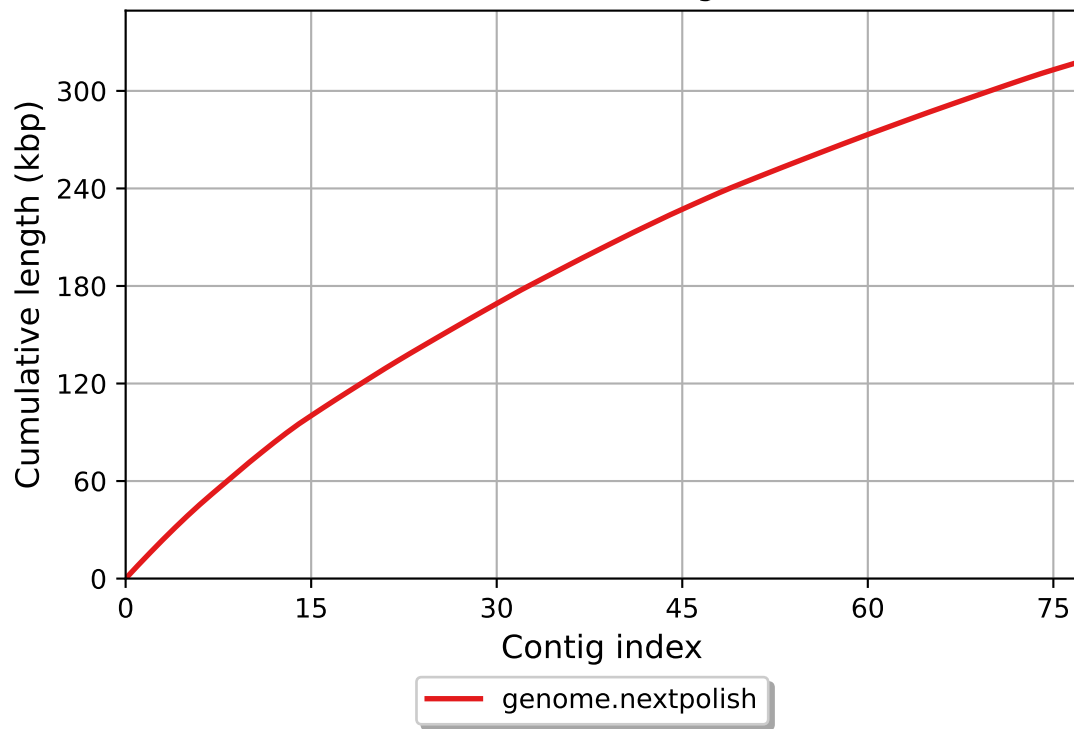
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

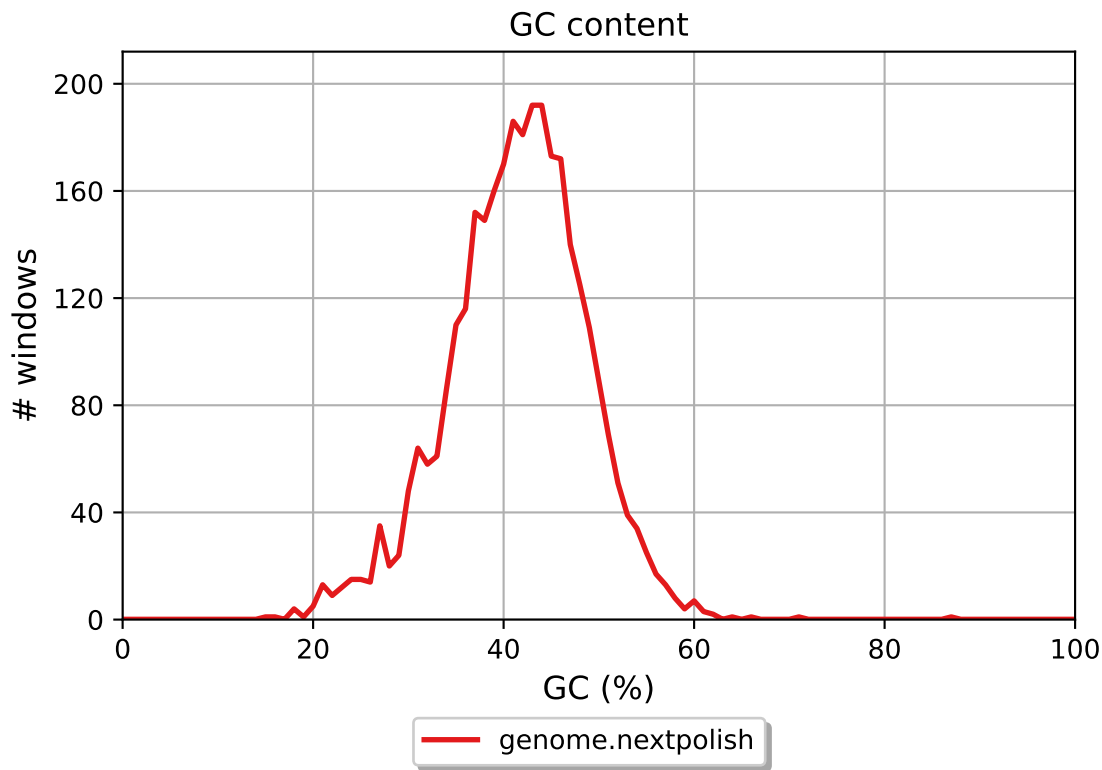
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

