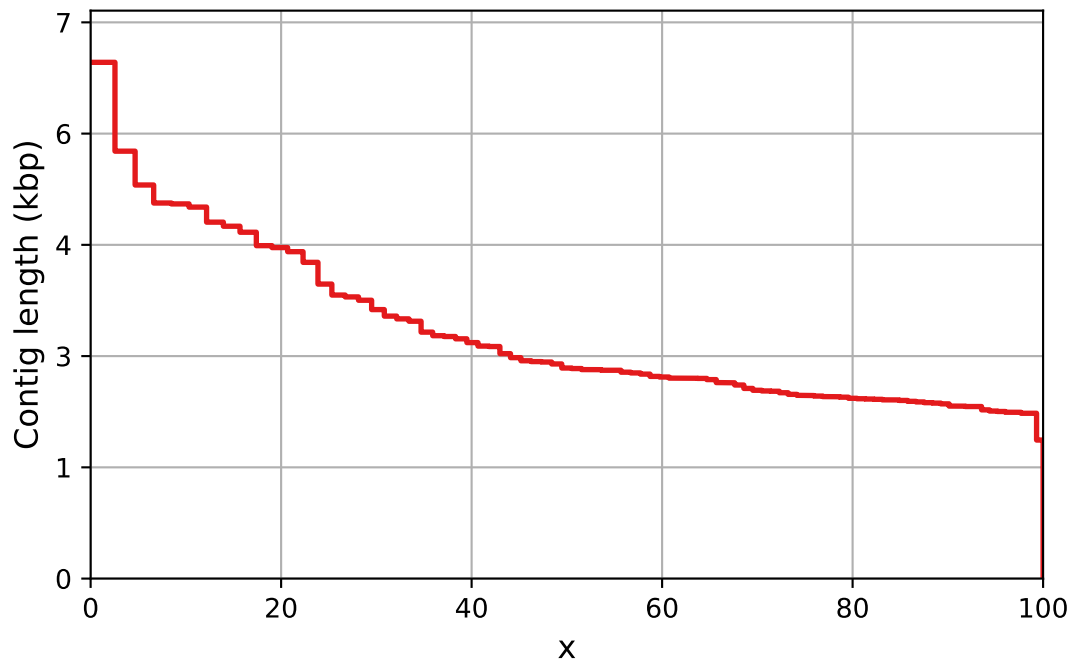


# Report

	genome.nextpolish
# contigs (>= 0 bp)	89
# contigs (>= 1000 bp)	89
# contigs (>= 5000 bp)	6
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	272353
Total length (>= 1000 bp)	272353
Total length (>= 5000 bp)	33156
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	89
Largest contig	6961
Total length	272353
GC (%)	41.17
N50	2839
N90	2356
auN	3347.2
L50	35
L90	77
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 2 part

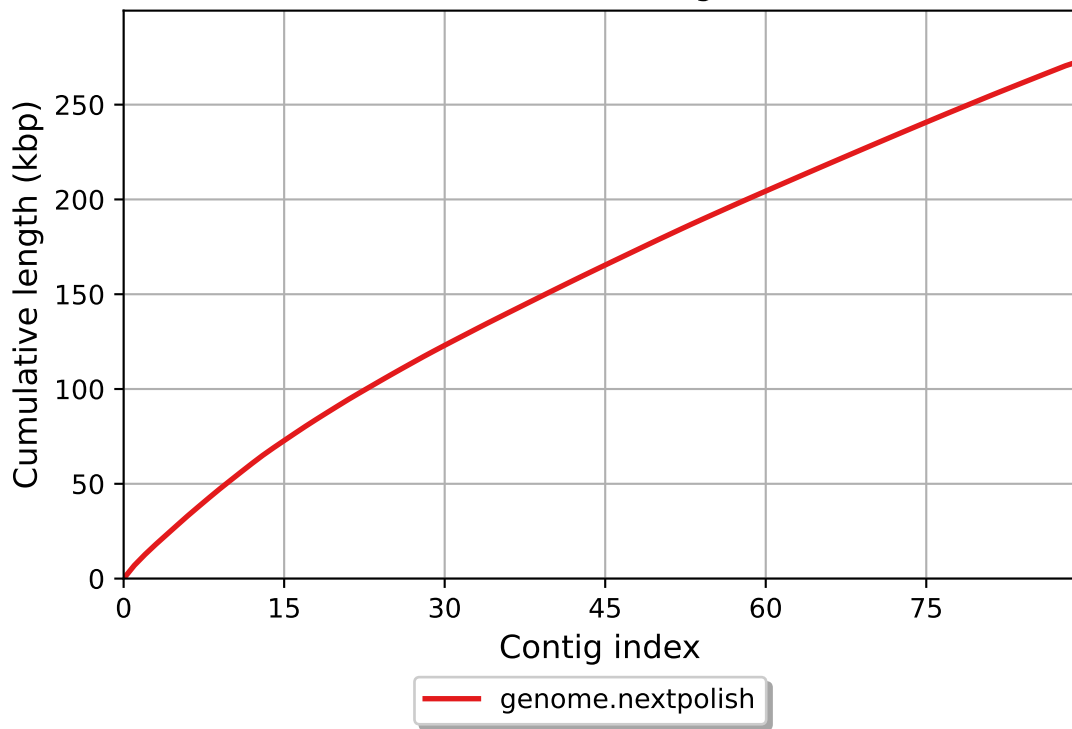
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

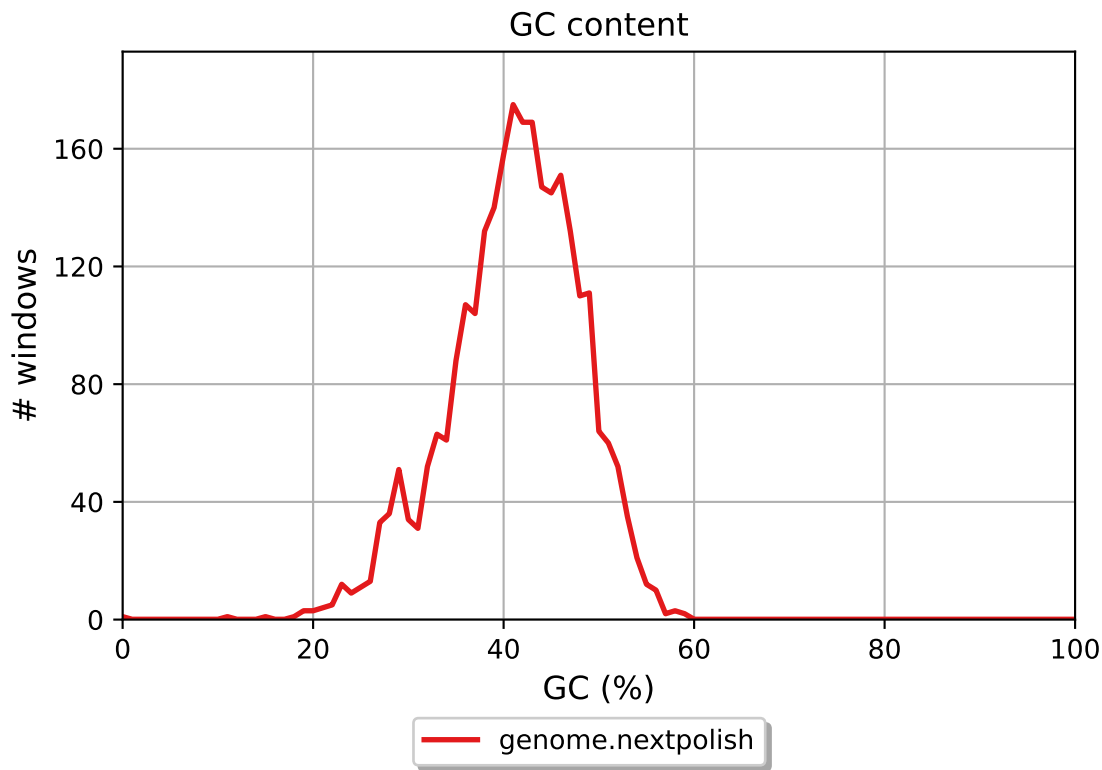
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

