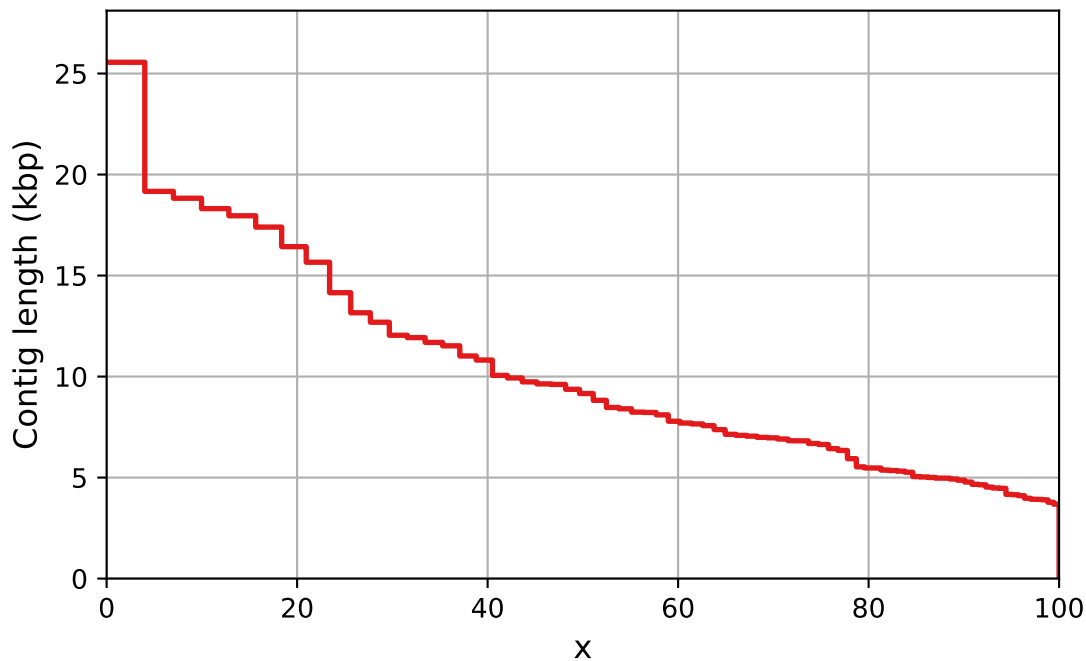


Report

	genome.nextpolish
# contigs (>= 0 bp)	77
# contigs (>= 1000 bp)	77
# contigs (>= 5000 bp)	58
# contigs (>= 10000 bp)	18
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	637807
Total length (>= 1000 bp)	637807
Total length (>= 5000 bp)	554881
Total length (>= 10000 bp)	268391
Total length (>= 25000 bp)	25557
Total length (>= 50000 bp)	0
# contigs	77
Largest contig	25557
Total length	637807
GC (%)	41.19
N50	9161
N90	4871
auN	10635.7
L50	24
L90	62
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 2 part

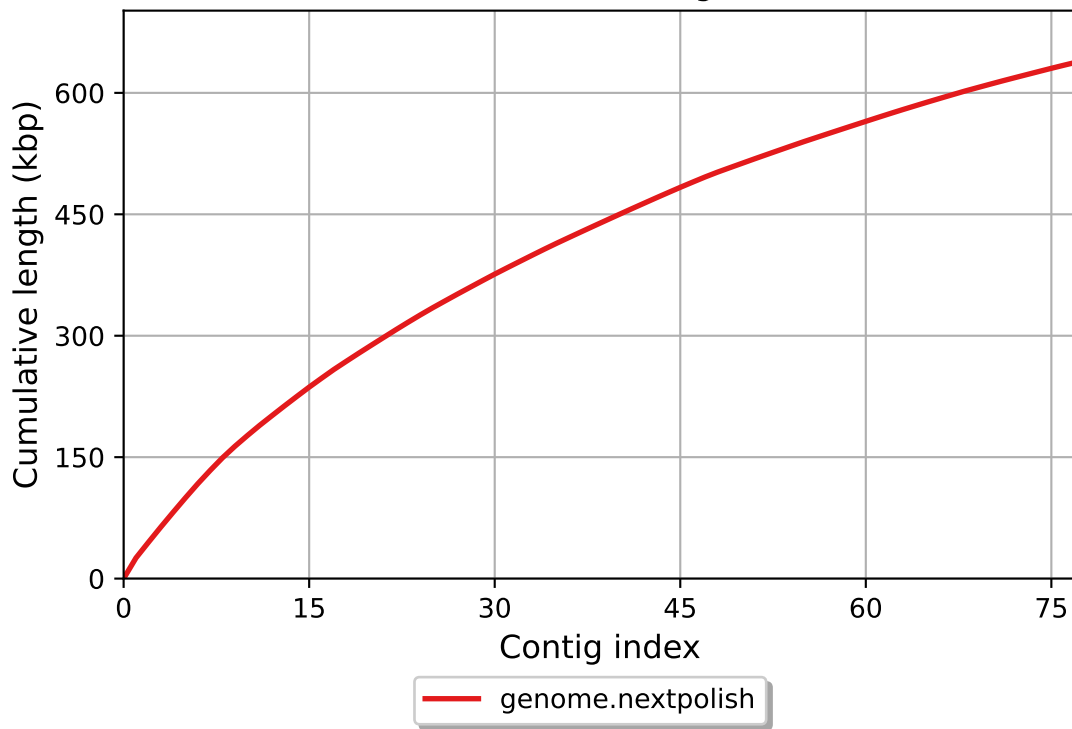
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

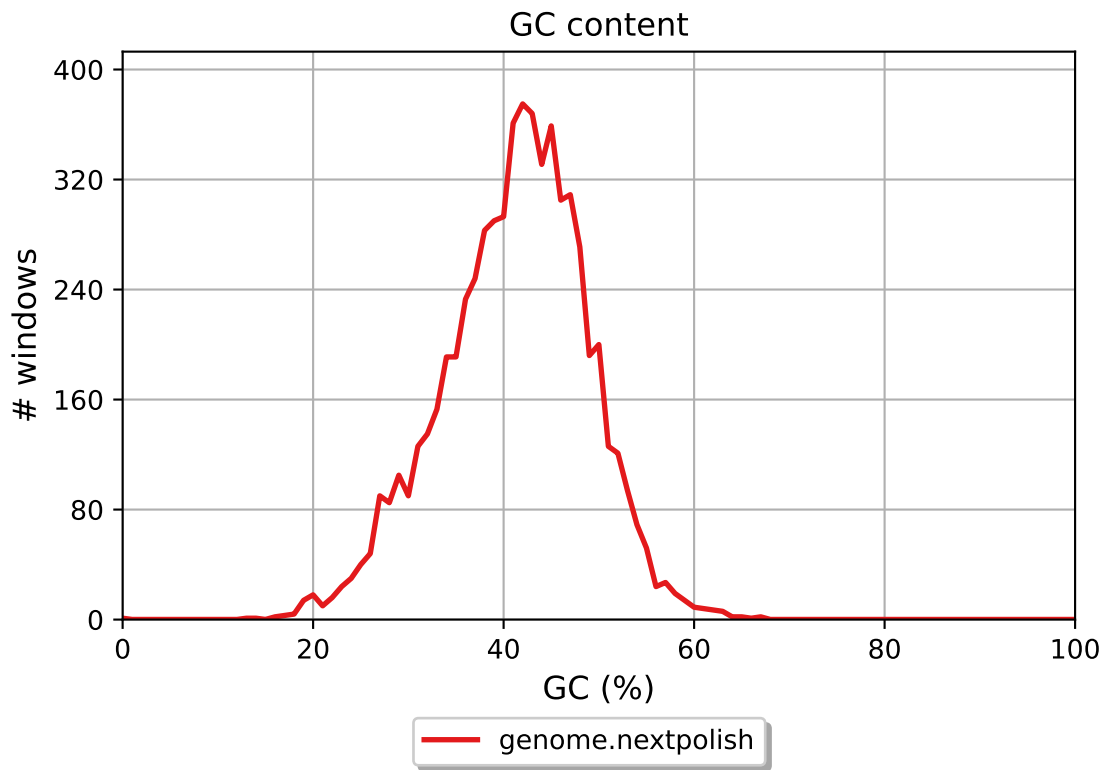
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

