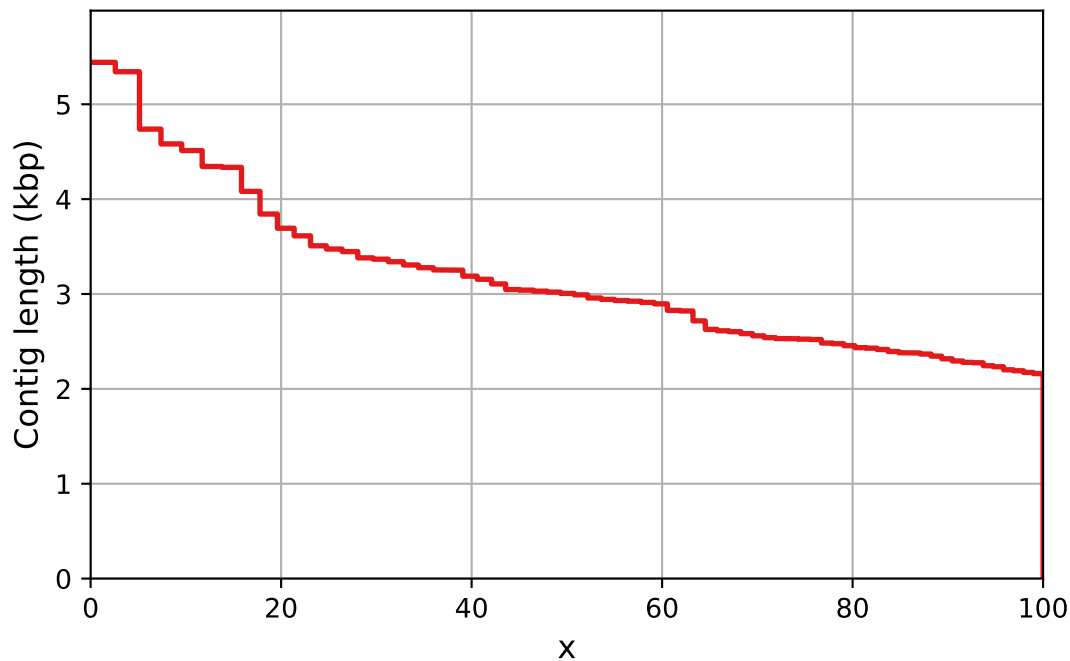


# Report

	genome.nextpolish
# contigs (>= 0 bp)	70
# contigs (>= 1000 bp)	70
# contigs (>= 5000 bp)	2
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	210194
Total length (>= 1000 bp)	210194
Total length (>= 5000 bp)	10785
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	70
Largest contig	5442
Total length	210194
GC (%)	41.89
N50	3006
N90	2317
auN	3187.3
L50	29
L90	61
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

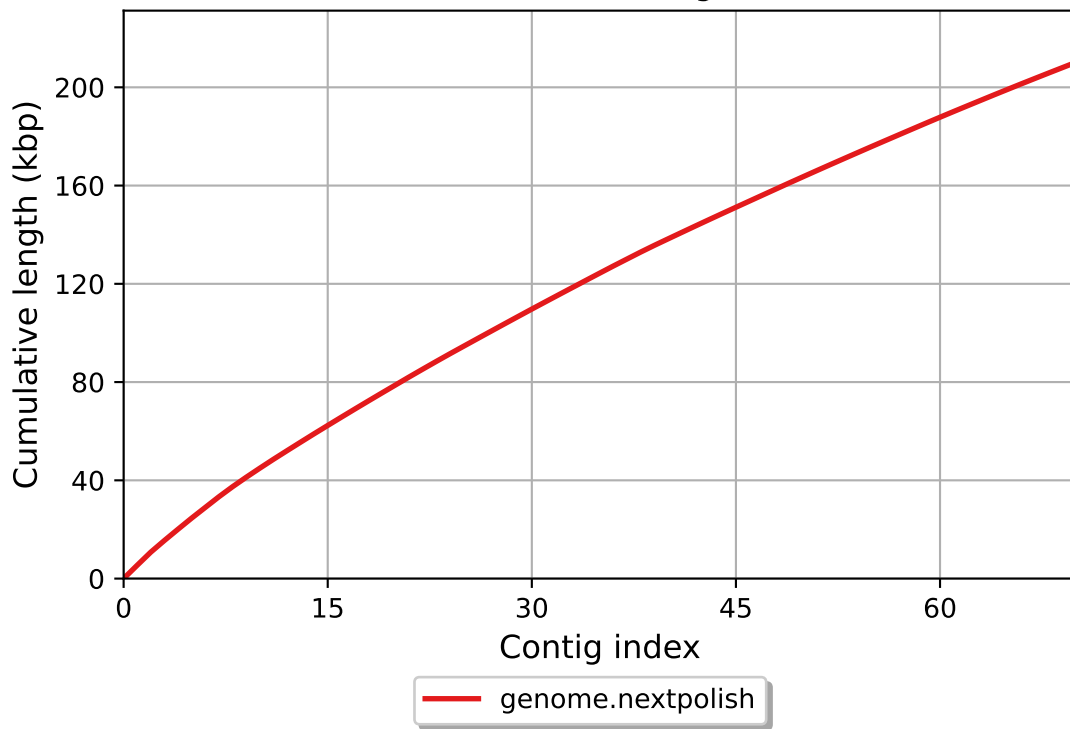
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

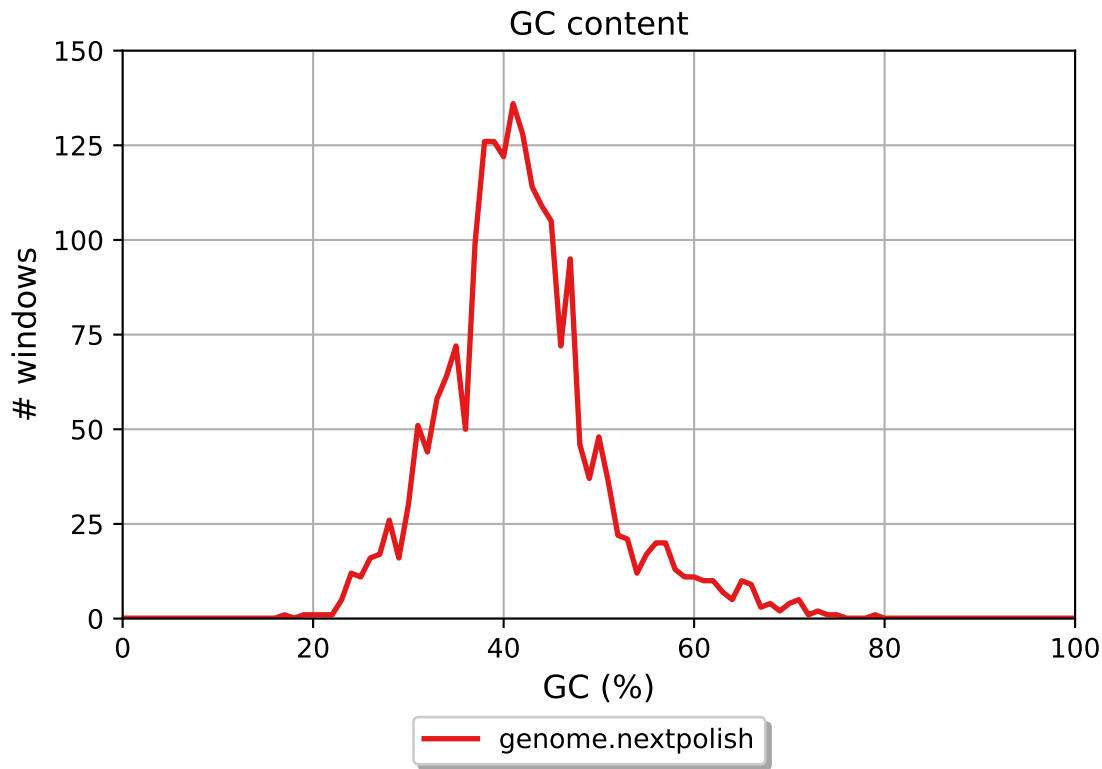
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

