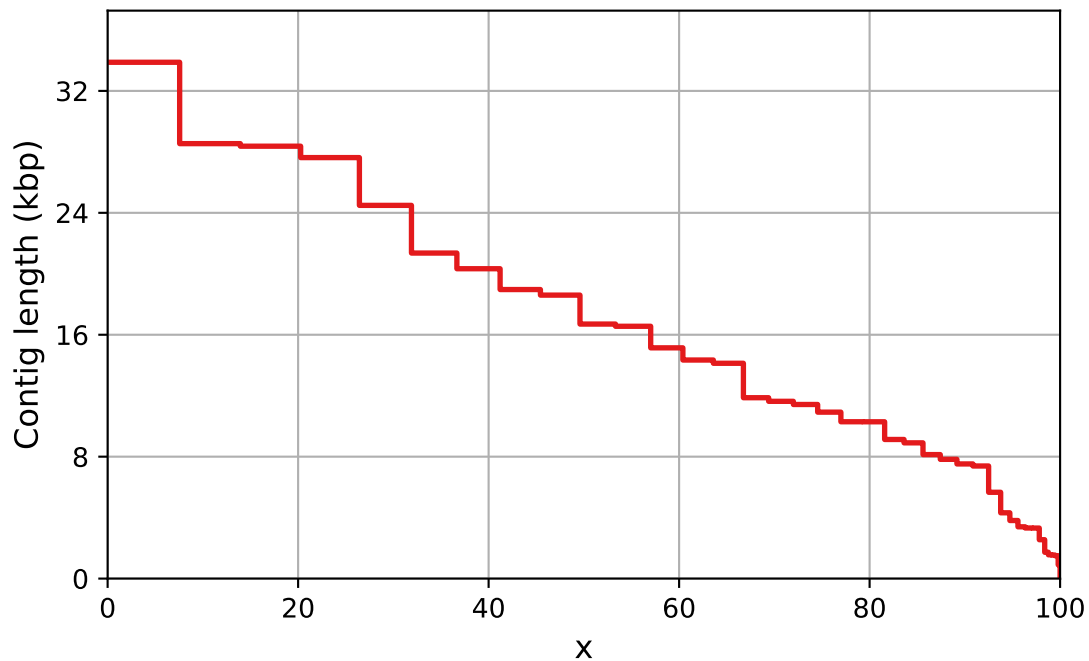


# Report

	genome.nextpolish
# contigs (>= 0 bp)	38
# contigs (>= 1000 bp)	37
# contigs (>= 5000 bp)	27
# contigs (>= 10000 bp)	20
# contigs (>= 25000 bp)	4
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	447975
Total length (>= 1000 bp)	447072
Total length (>= 5000 bp)	420024
Total length (>= 10000 bp)	365454
Total length (>= 25000 bp)	118415
Total length (>= 50000 bp)	0
# contigs	38
Largest contig	33879
Total length	447975
GC (%)	37.54
N50	16701
N90	7523
auN	18188.7
L50	10
L90	25
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 0 part

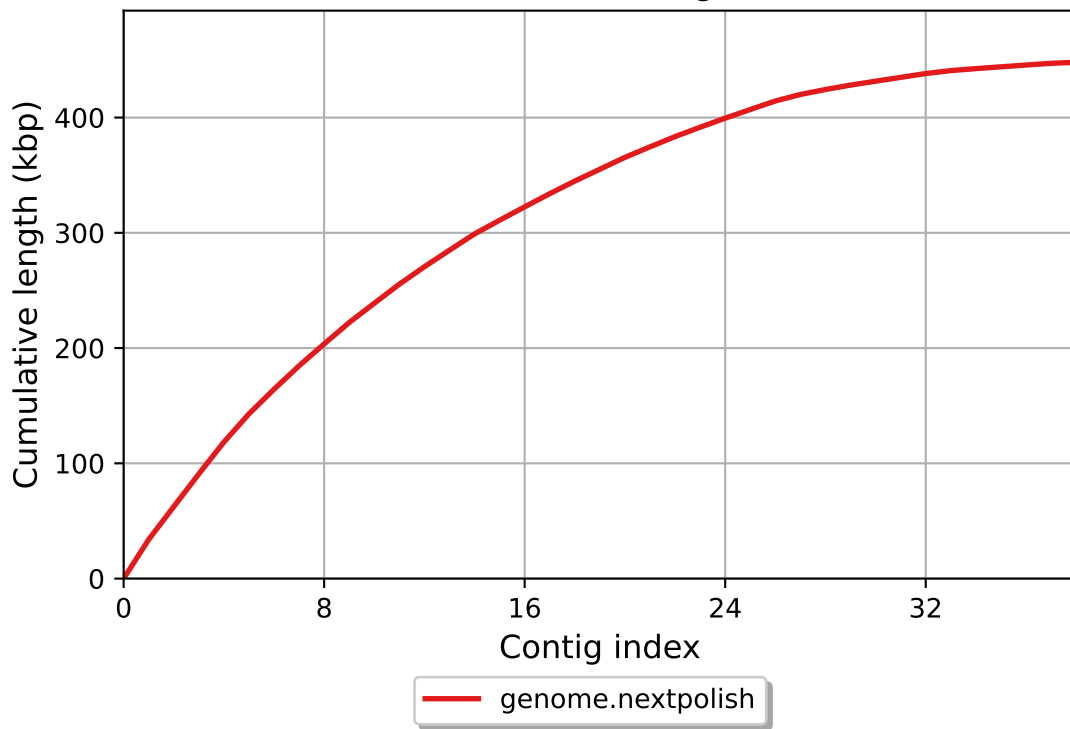
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

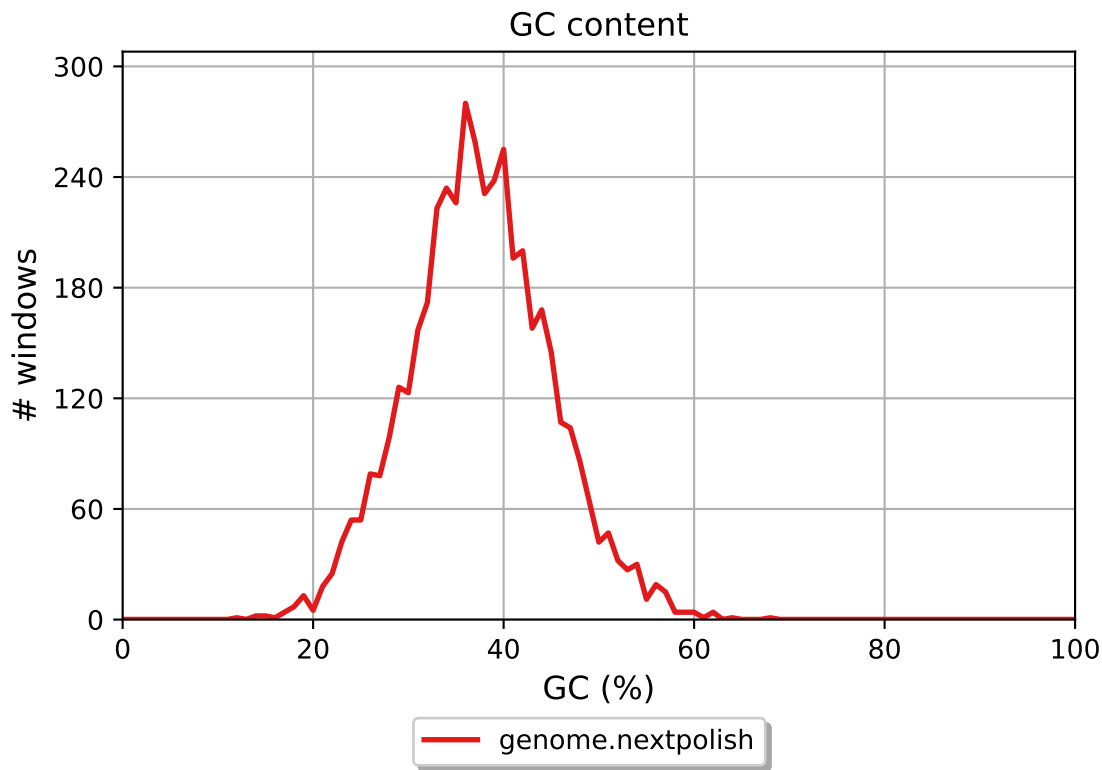
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

