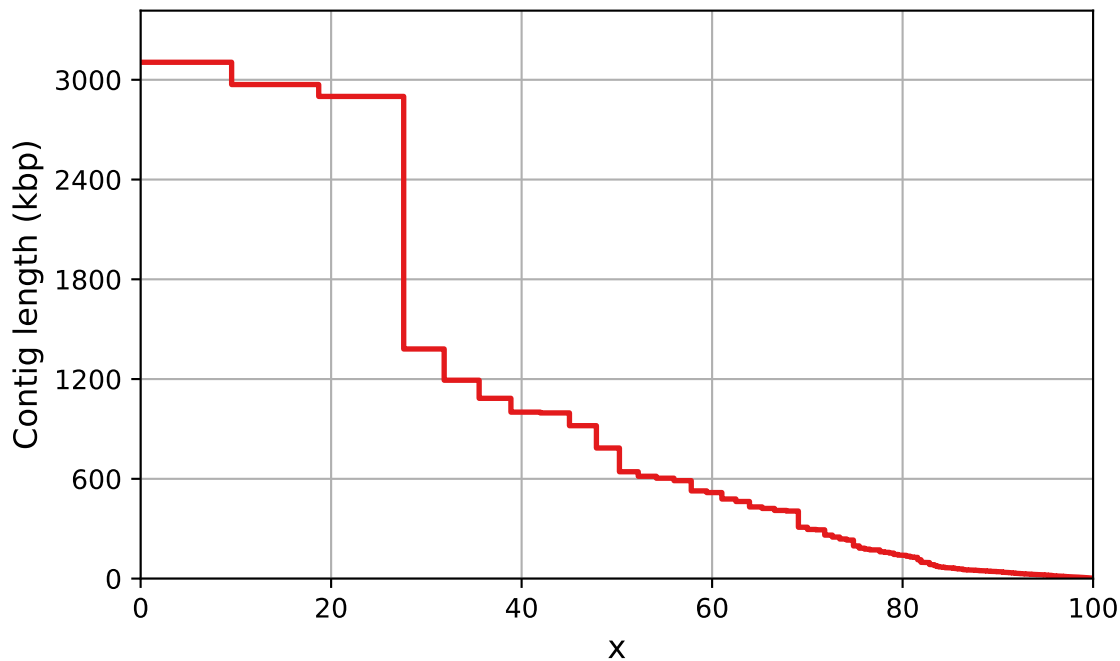


Report

	genome.nextpolish
# contigs (>= 0 bp)	338
# contigs (>= 1000 bp)	334
# contigs (>= 5000 bp)	279
# contigs (>= 10000 bp)	209
# contigs (>= 25000 bp)	129
# contigs (>= 50000 bp)	73
Total length (>= 0 bp)	32499563
Total length (>= 1000 bp)	32496669
Total length (>= 5000 bp)	32326010
Total length (>= 10000 bp)	31809897
Total length (>= 25000 bp)	30517087
Total length (>= 50000 bp)	28527683
# contigs	338
Largest contig	3105315
Total length	32499563
GC (%)	41.94
N50	785343
N90	41637
auN	1204246.7
L50	10
L90	89
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	128 + 9 part

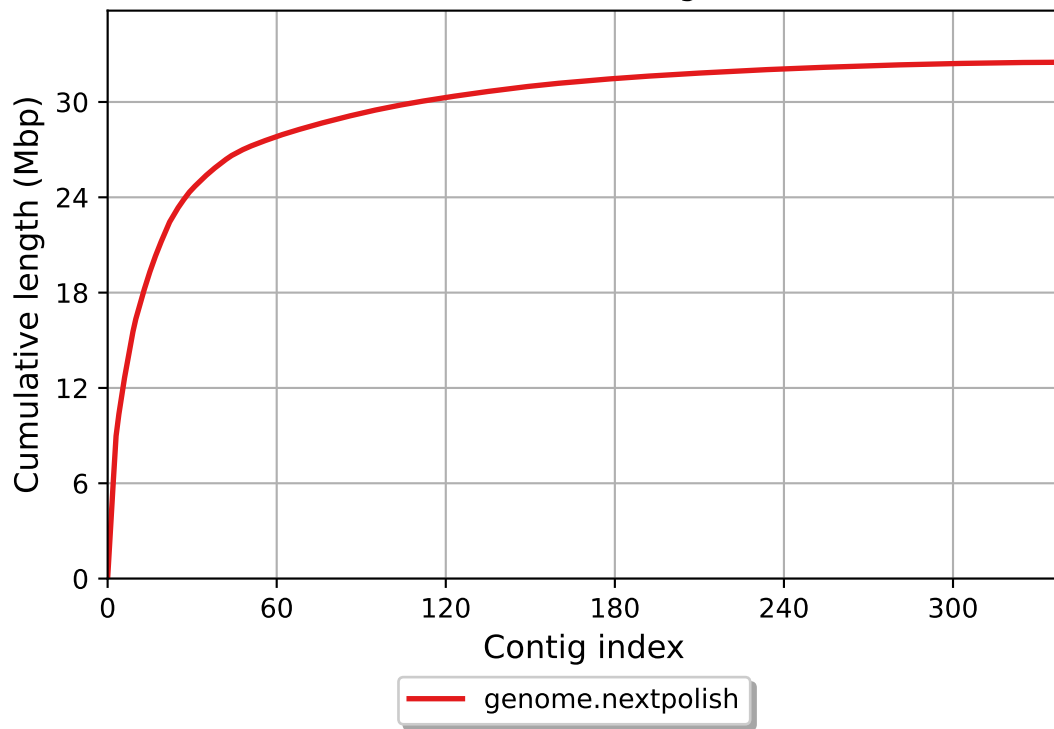
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

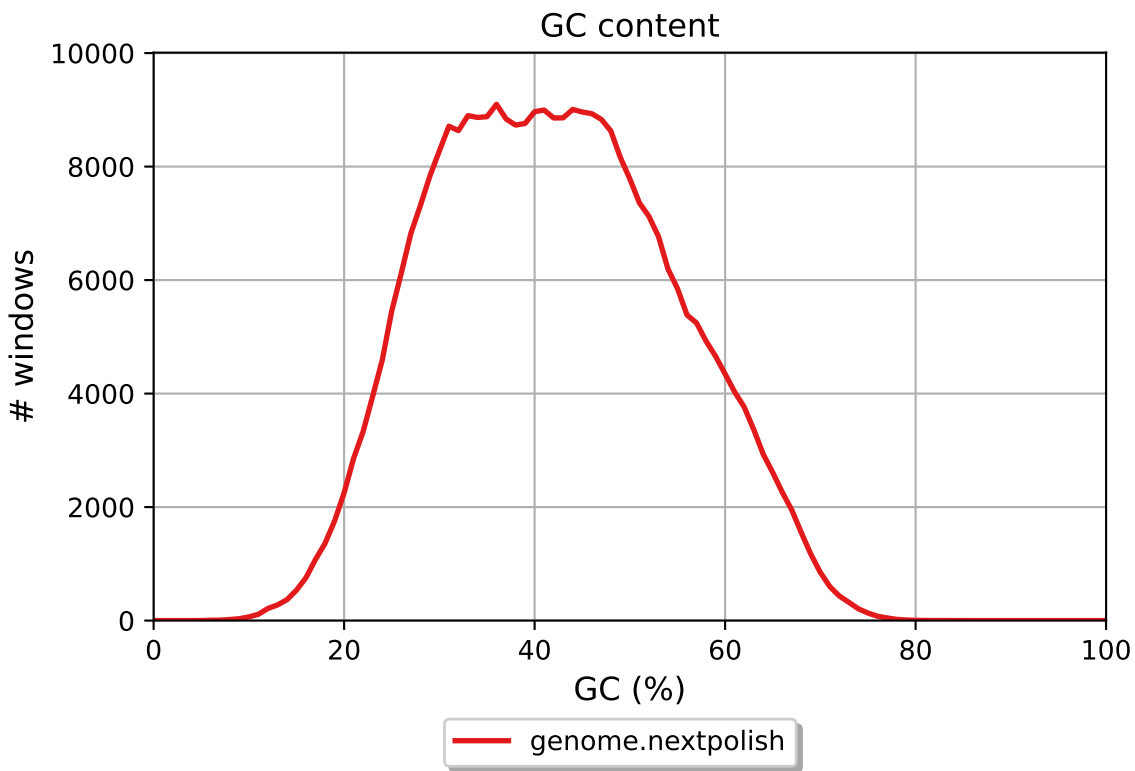
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

