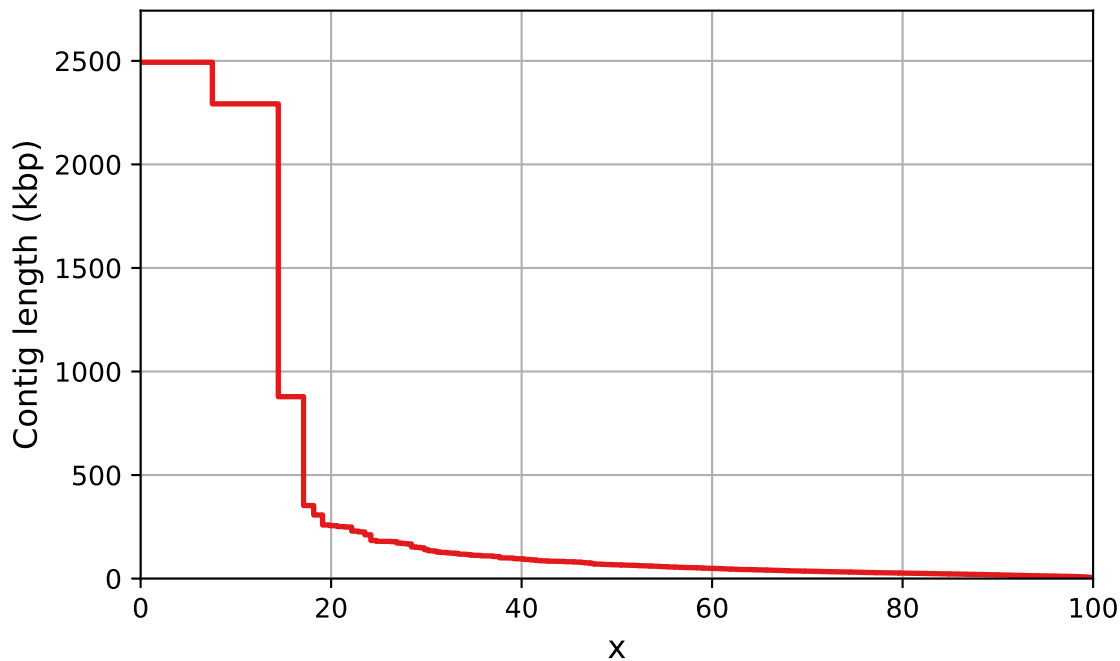


Report

	genome.nextpolish
# contigs (>= 0 bp)	773
# contigs (>= 1000 bp)	772
# contigs (>= 5000 bp)	765
# contigs (>= 10000 bp)	688
# contigs (>= 25000 bp)	367
# contigs (>= 50000 bp)	149
Total length (>= 0 bp)	33113095
Total length (>= 1000 bp)	33112217
Total length (>= 5000 bp)	33082905
Total length (>= 10000 bp)	32466392
Total length (>= 25000 bp)	27196331
Total length (>= 50000 bp)	19703458
# contigs	773
Largest contig	2493248
Total length	33113095
GC (%)	48.12
N50	66602
N90	17819
auN	434452.4
L50	95
L90	490
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	126 + 2 part

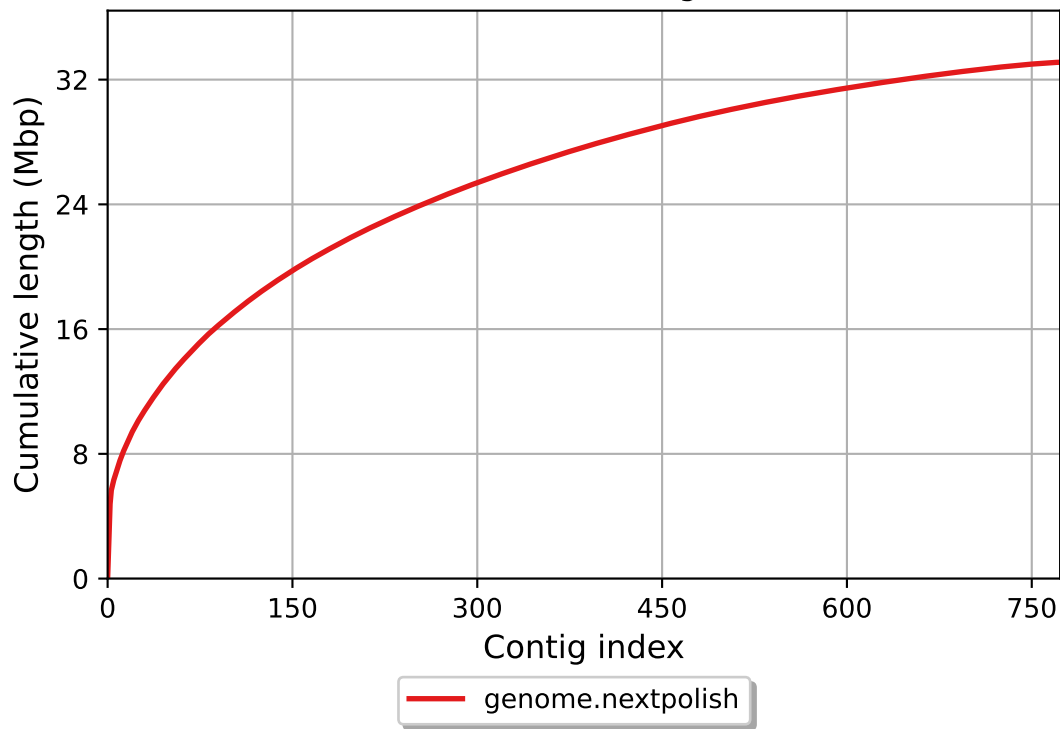
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

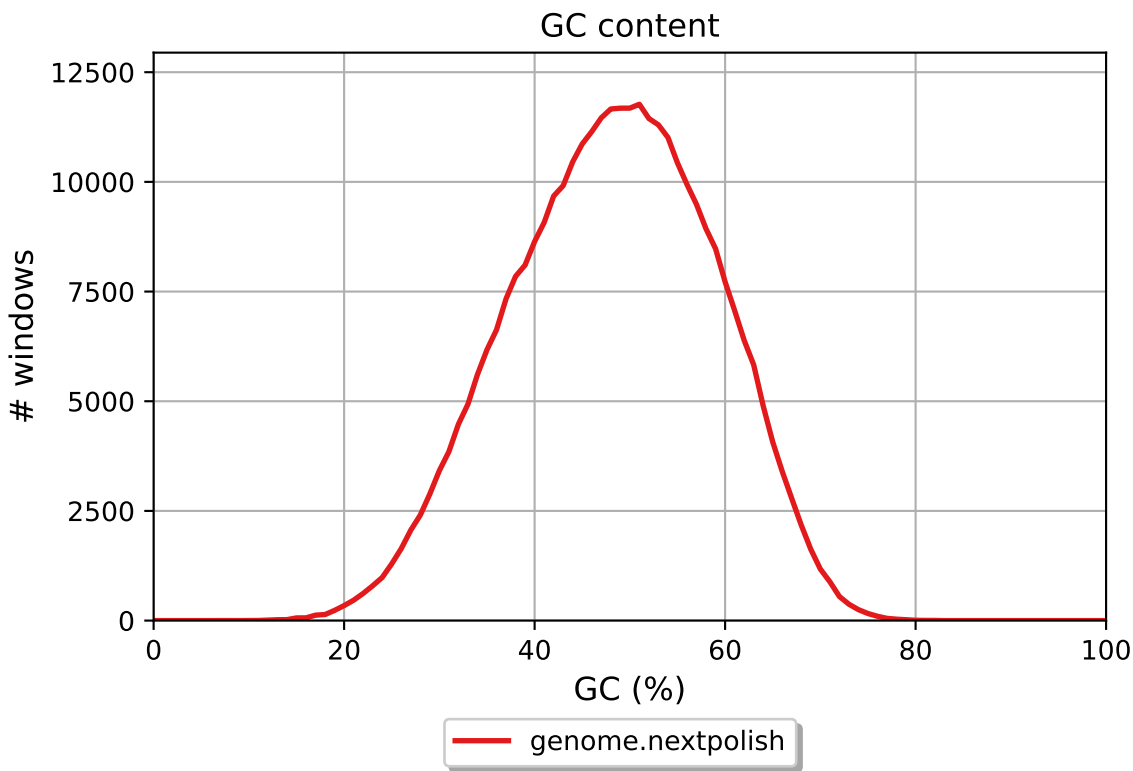
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

