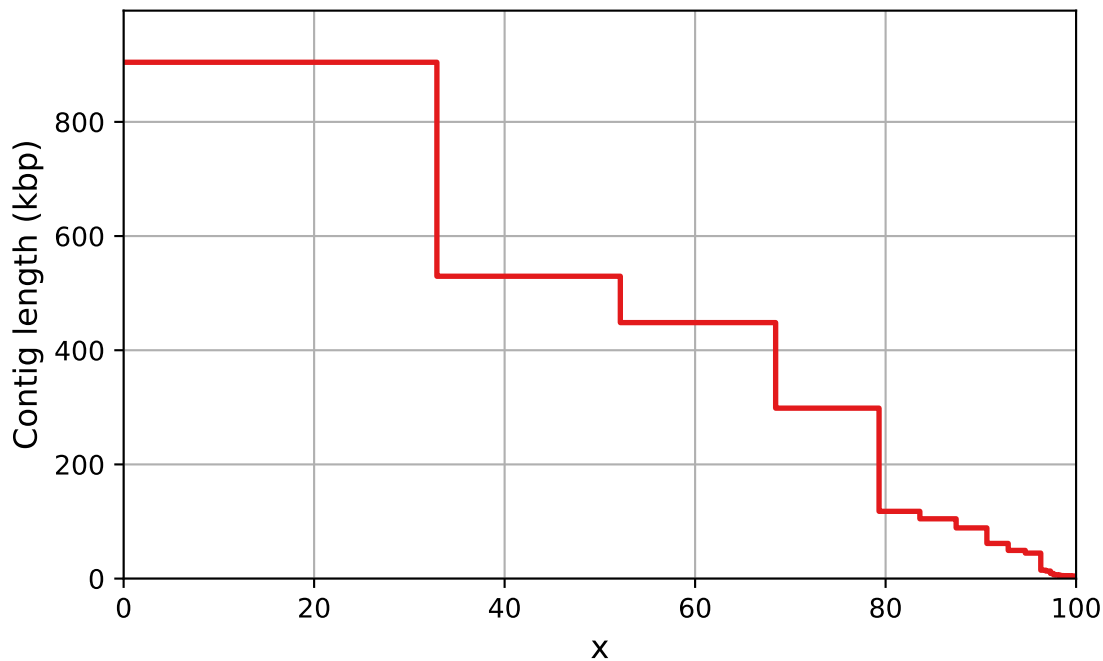


Report

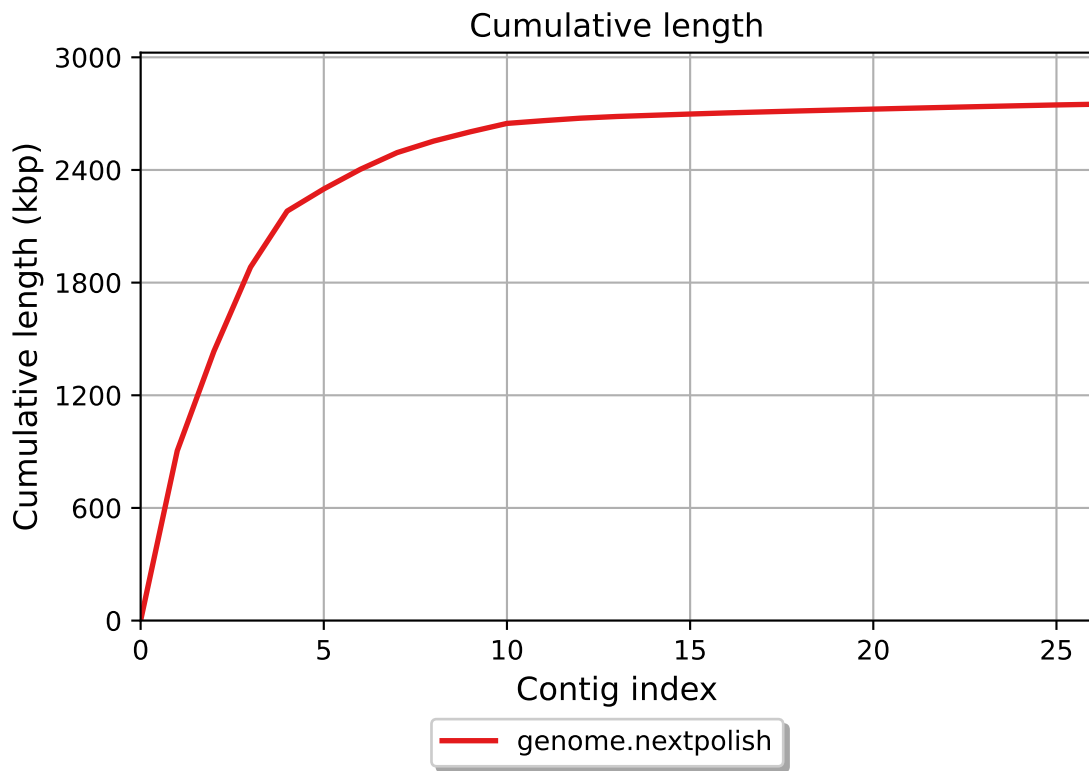
	genome.nextpolish
# contigs (>= 0 bp)	26
# contigs (>= 1000 bp)	26
# contigs (>= 5000 bp)	18
# contigs (>= 10000 bp)	12
# contigs (>= 25000 bp)	10
# contigs (>= 50000 bp)	8
Total length (>= 0 bp)	2749882
Total length (>= 1000 bp)	2749882
Total length (>= 5000 bp)	2714476
Total length (>= 10000 bp)	2675788
Total length (>= 25000 bp)	2647886
Total length (>= 50000 bp)	2553894
# contigs	26
Largest contig	904433
Total length	2749882
GC (%)	34.38
N50	529611
N90	88779
auN	520182.1
L50	2
L90	7
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	12 + 2 part

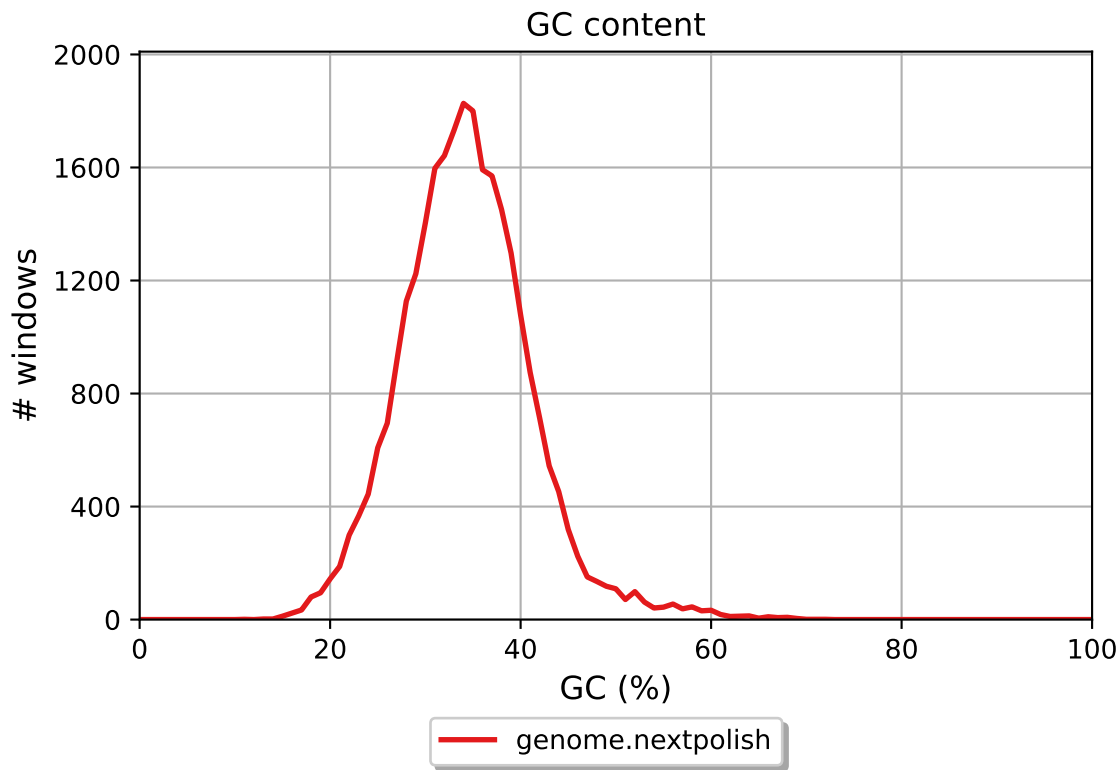
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

