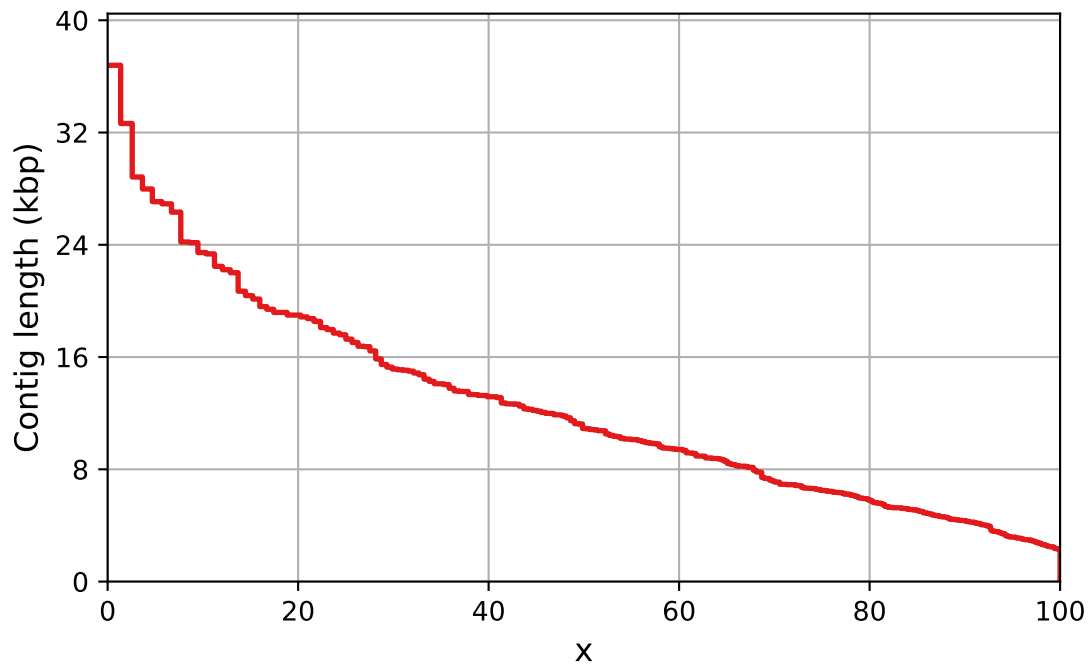


Report

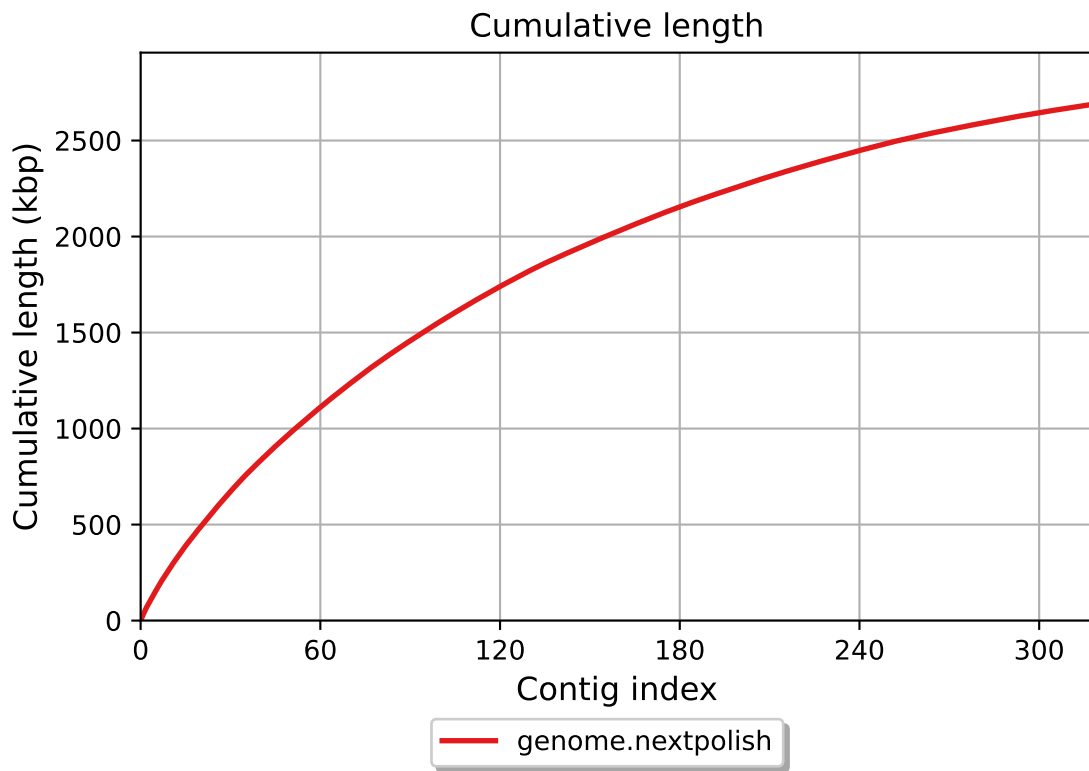
	genome.nextpolish
# contigs (>= 0 bp)	318
# contigs (>= 1000 bp)	318
# contigs (>= 5000 bp)	207
# contigs (>= 10000 bp)	95
# contigs (>= 25000 bp)	7
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2688843
Total length (>= 1000 bp)	2688843
Total length (>= 5000 bp)	2298105
Total length (>= 10000 bp)	1507977
Total length (>= 25000 bp)	206564
Total length (>= 50000 bp)	0
# contigs	318
Largest contig	36792
Total length	2688843
GC (%)	33.25
N50	10910
N90	4342
auN	12638.3
L50	80
L90	234
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 2 part

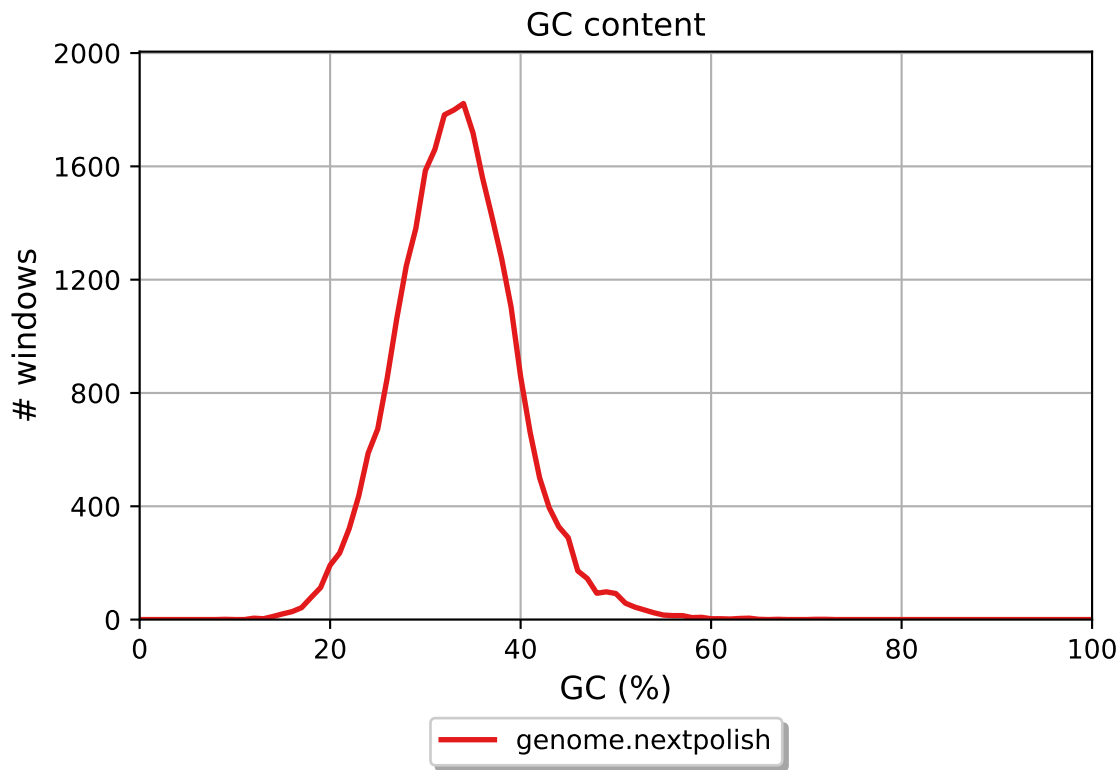
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

