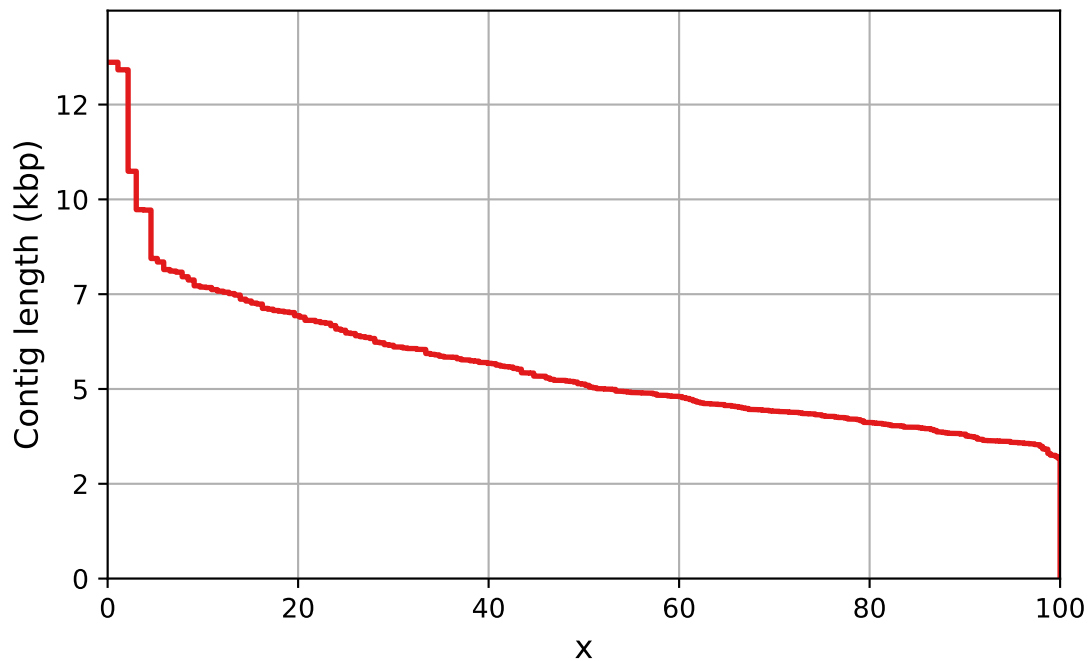


Report

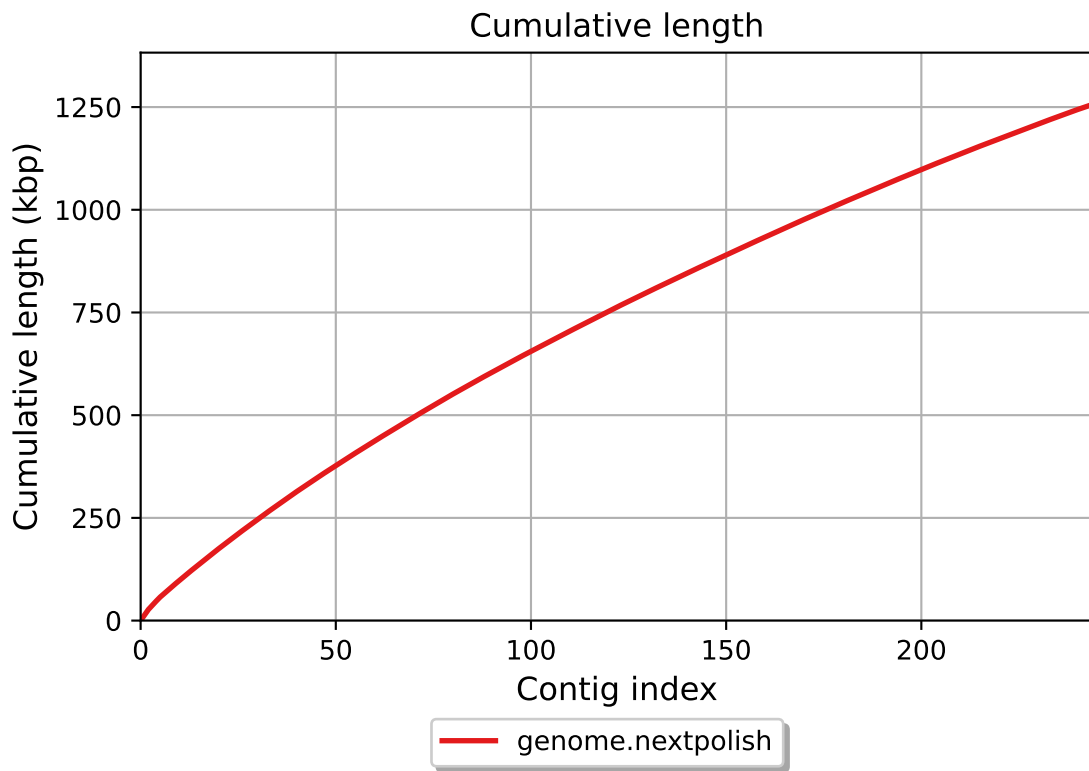
	genome.nextpolish
# contigs (>= 0 bp)	244
# contigs (>= 1000 bp)	244
# contigs (>= 5000 bp)	100
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1256822
Total length (>= 1000 bp)	1256822
Total length (>= 5000 bp)	655484
Total length (>= 10000 bp)	37773
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	244
Largest contig	13615
Total length	1256822
GC (%)	41.54
N50	5129
N90	3808
auN	5617.7
L50	95
L90	209
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 1 part

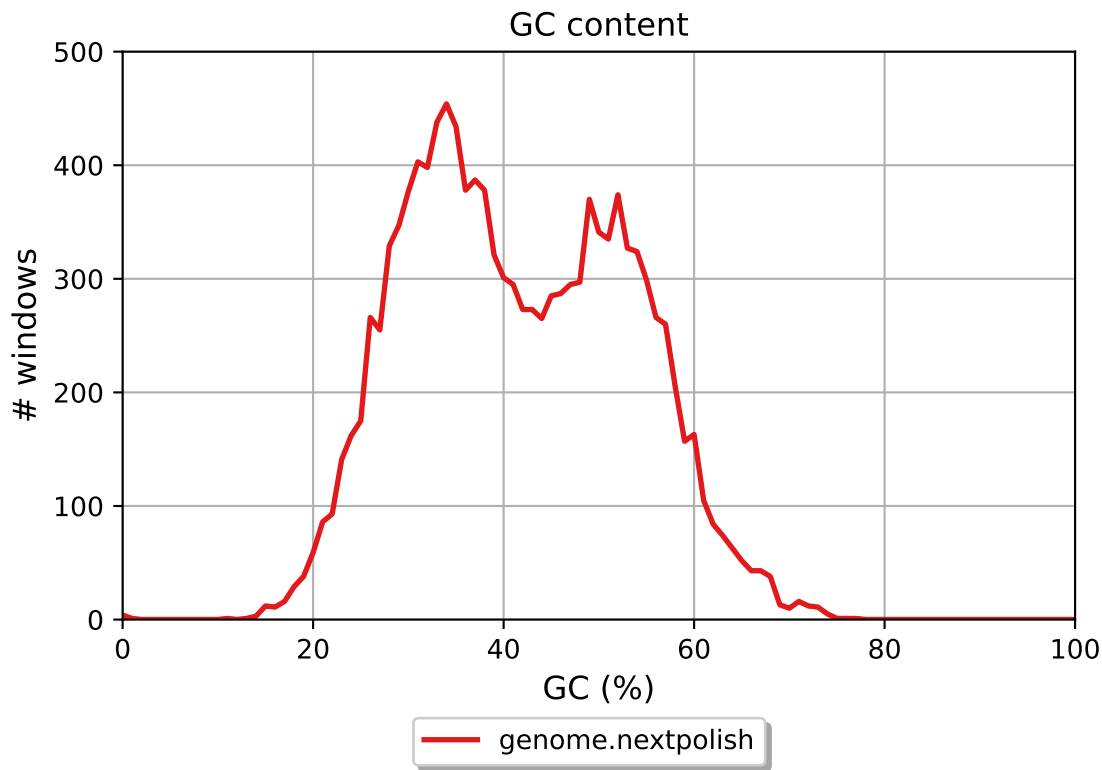
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

