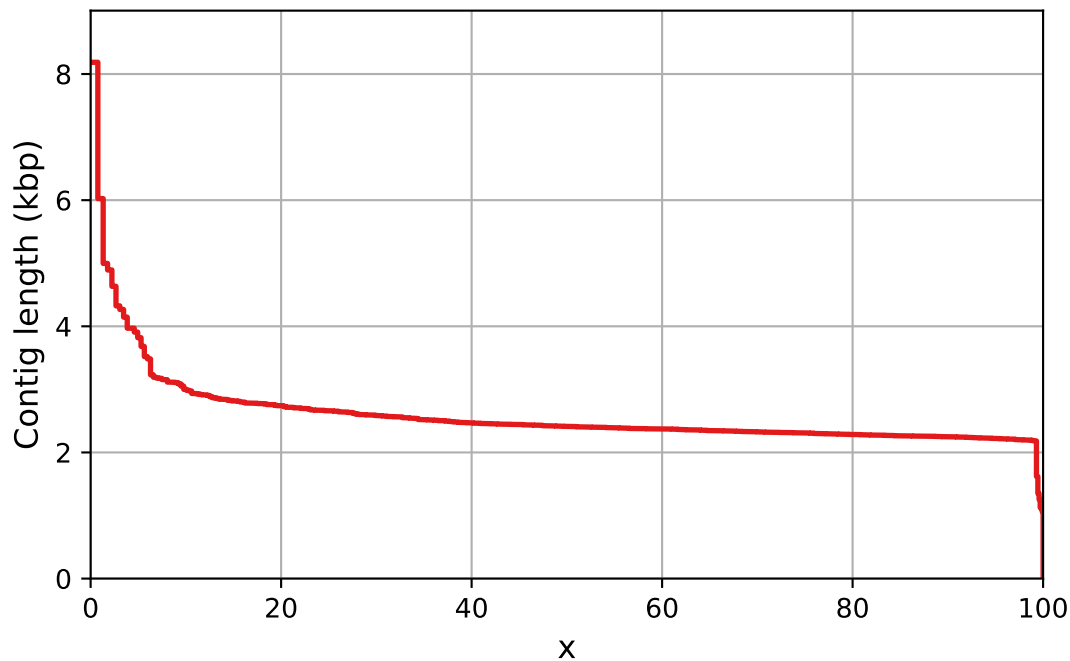


Report

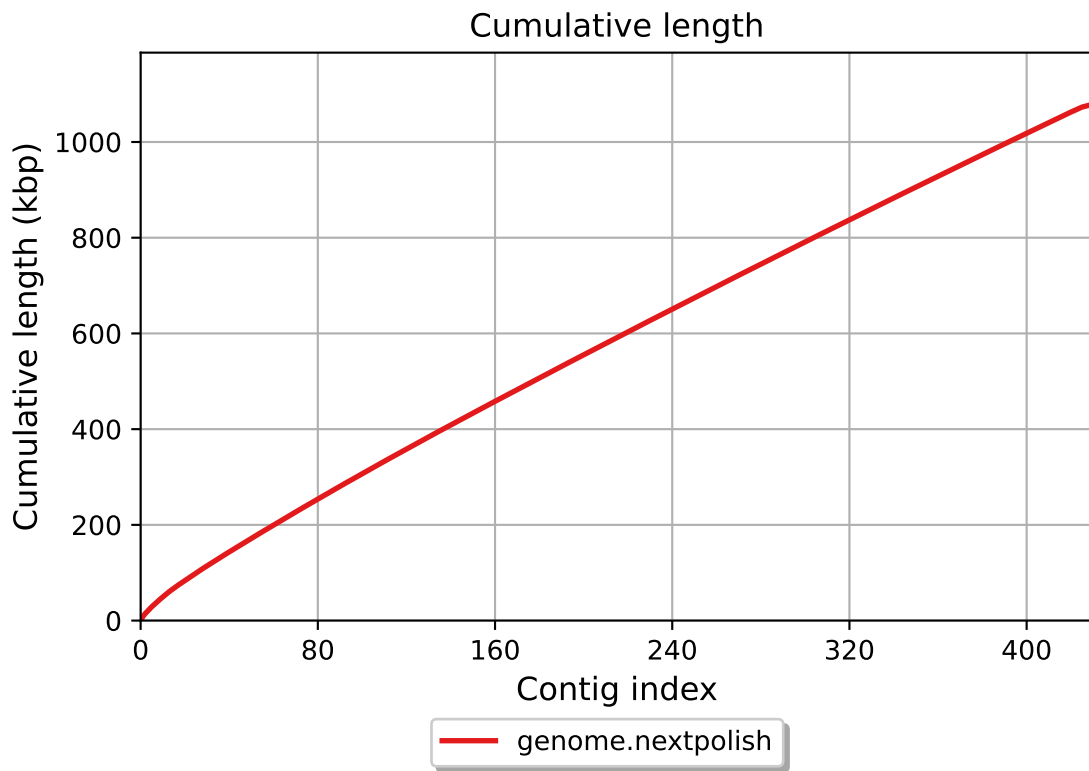
	genome.nextpolish
# contigs (>= 0 bp)	430
# contigs (>= 1000 bp)	430
# contigs (>= 5000 bp)	2
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1078833
Total length (>= 1000 bp)	1078833
Total length (>= 5000 bp)	14211
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	430
Largest contig	8186
Total length	1078833
GC (%)	37.97
N50	2414
N90	2249
auN	2612.5
L50	194
L90	379
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 2 part

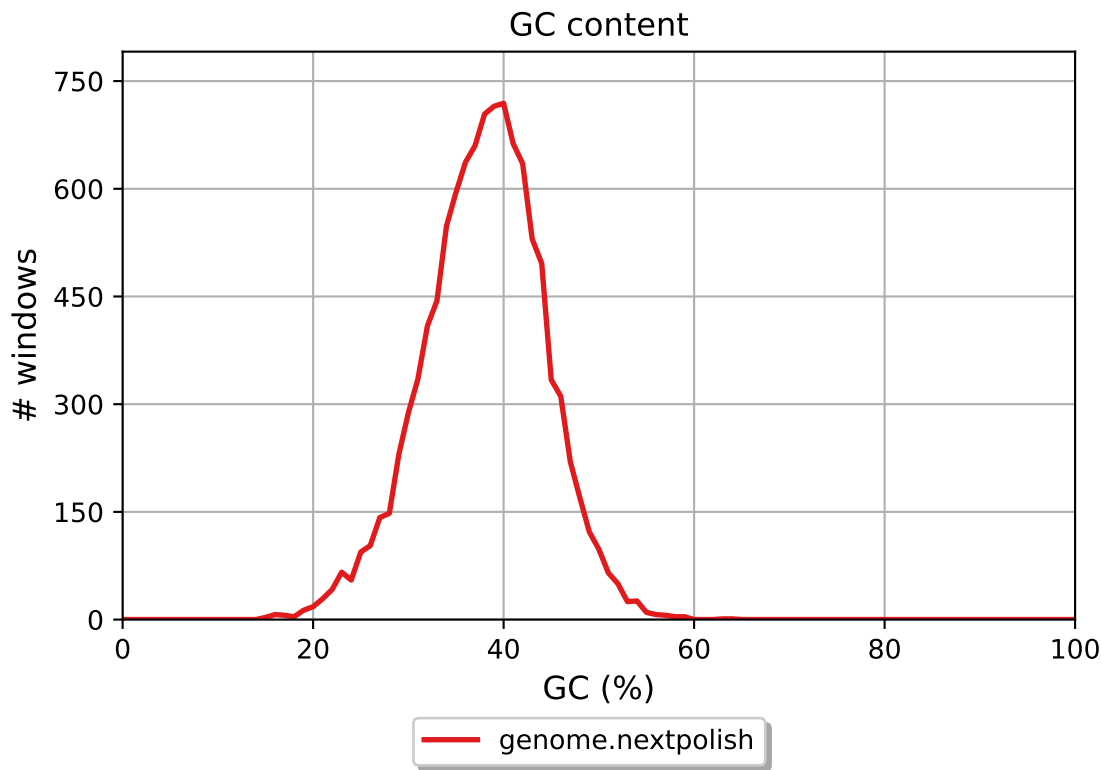
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

