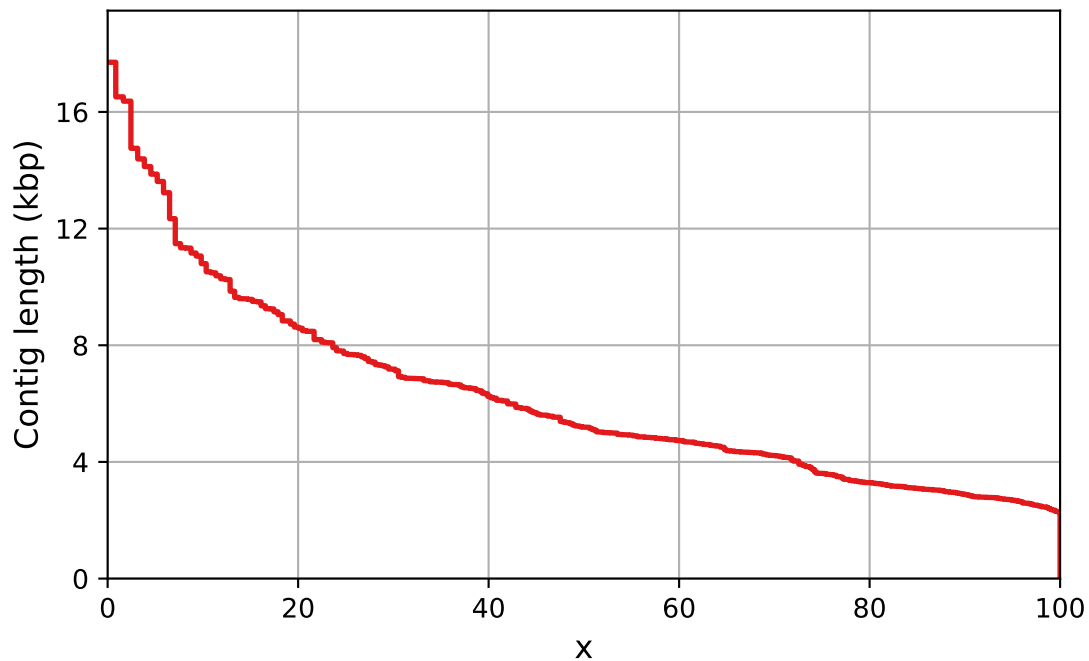


Report

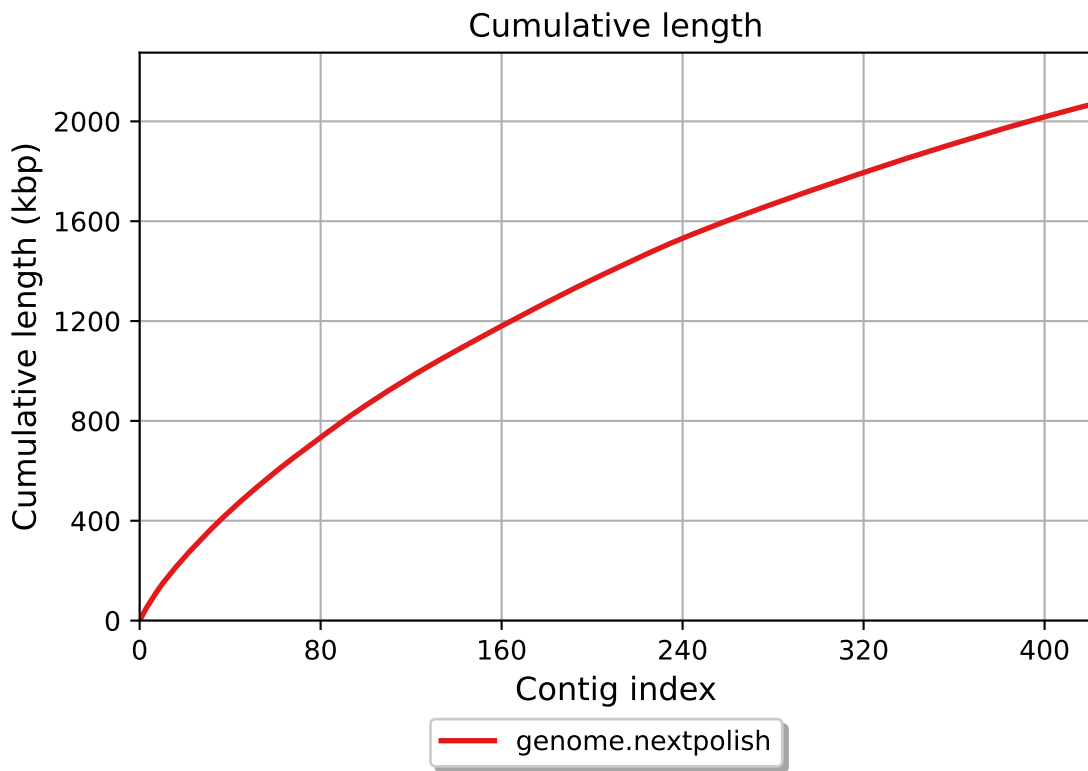
	genome.nextpolish
# contigs (>= 0 bp)	421
# contigs (>= 1000 bp)	421
# contigs (>= 5000 bp)	141
# contigs (>= 10000 bp)	21
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2068546
Total length (>= 1000 bp)	2068546
Total length (>= 5000 bp)	1086781
Total length (>= 10000 bp)	266002
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	421
Largest contig	17702
Total length	2068546
GC (%)	33.75
N50	5195
N90	2889
auN	6248.4
L50	131
L90	343
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 1 part

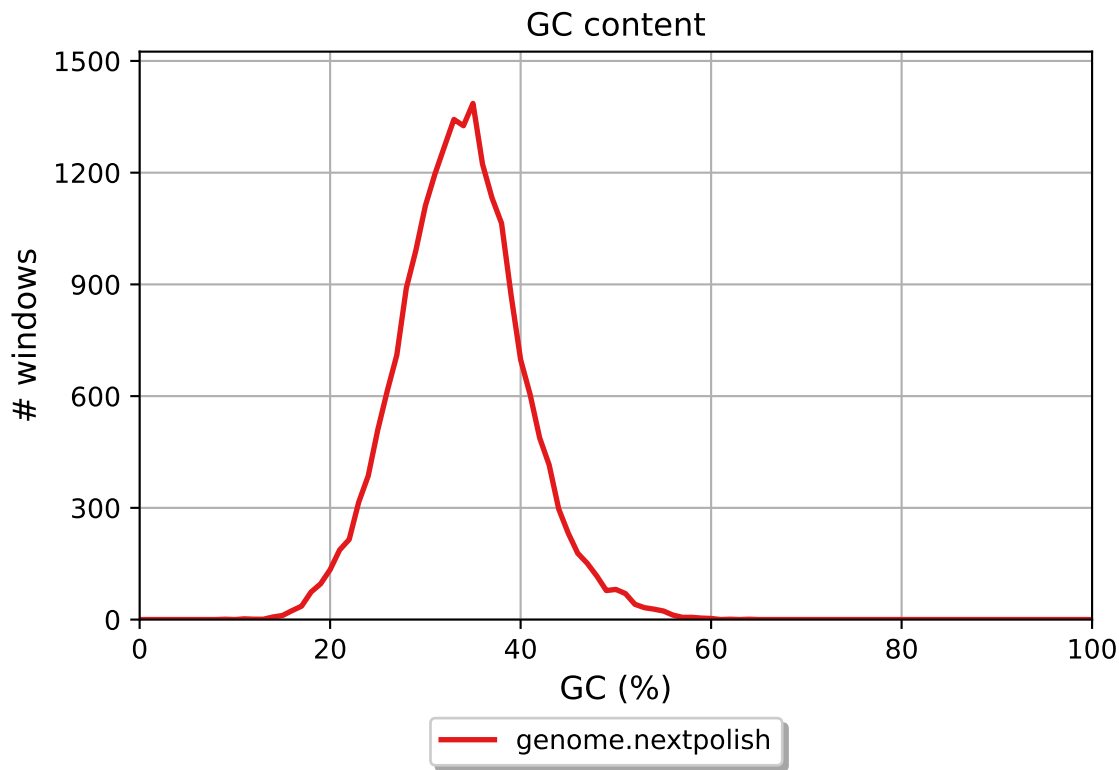
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

