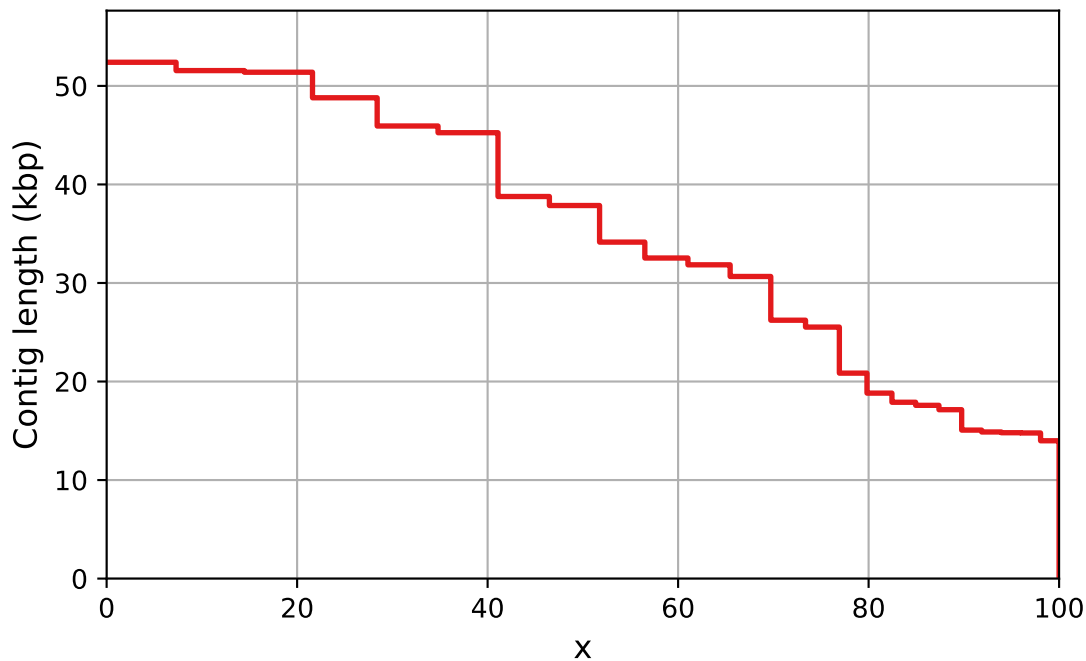


Report

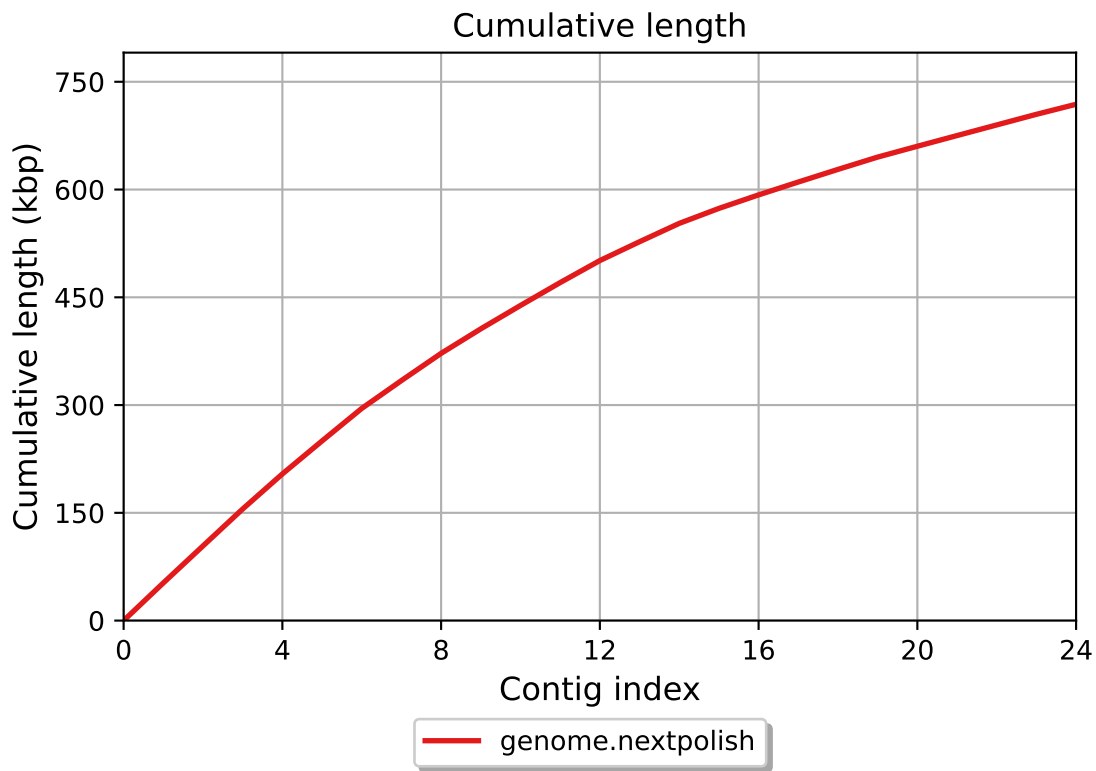
	genome.nextpolish
# contigs (>= 0 bp)	24
# contigs (>= 1000 bp)	24
# contigs (>= 5000 bp)	24
# contigs (>= 10000 bp)	24
# contigs (>= 25000 bp)	14
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	718690
Total length (>= 1000 bp)	718690
Total length (>= 5000 bp)	718690
Total length (>= 10000 bp)	718690
Total length (>= 25000 bp)	552884
Total length (>= 50000 bp)	155341
# contigs	24
Largest contig	52399
Total length	718690
GC (%)	38.18
N50	37861
N90	15069
auN	35941.5
L50	8
L90	20
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 0 part

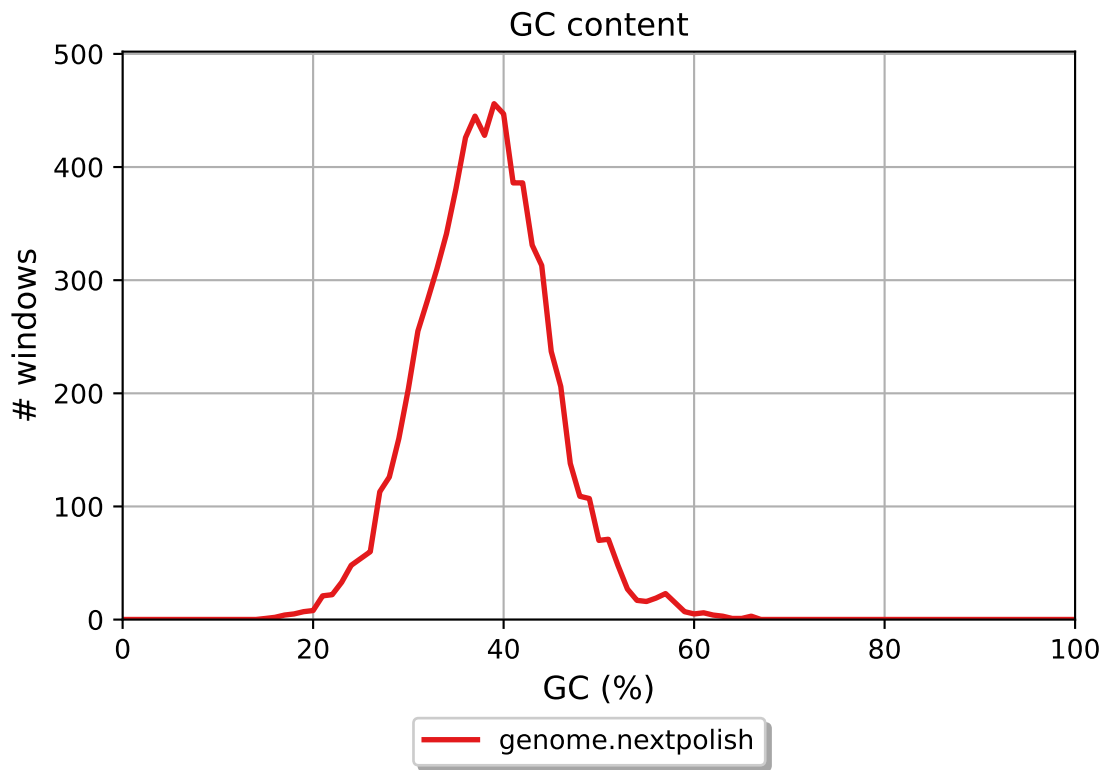
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

