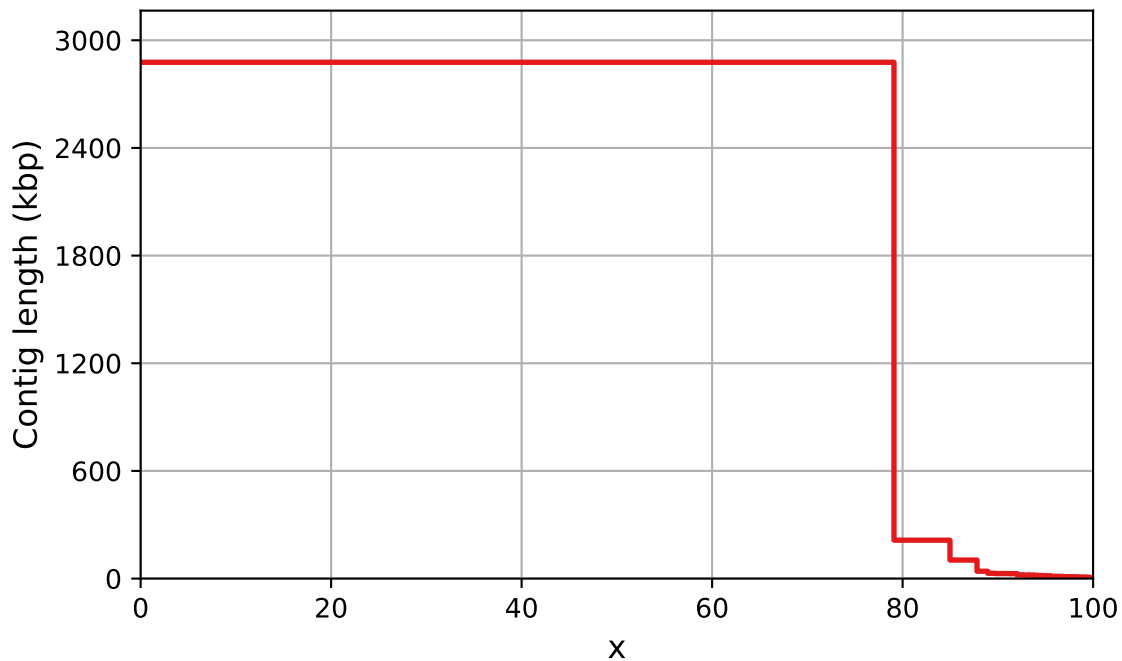


Report

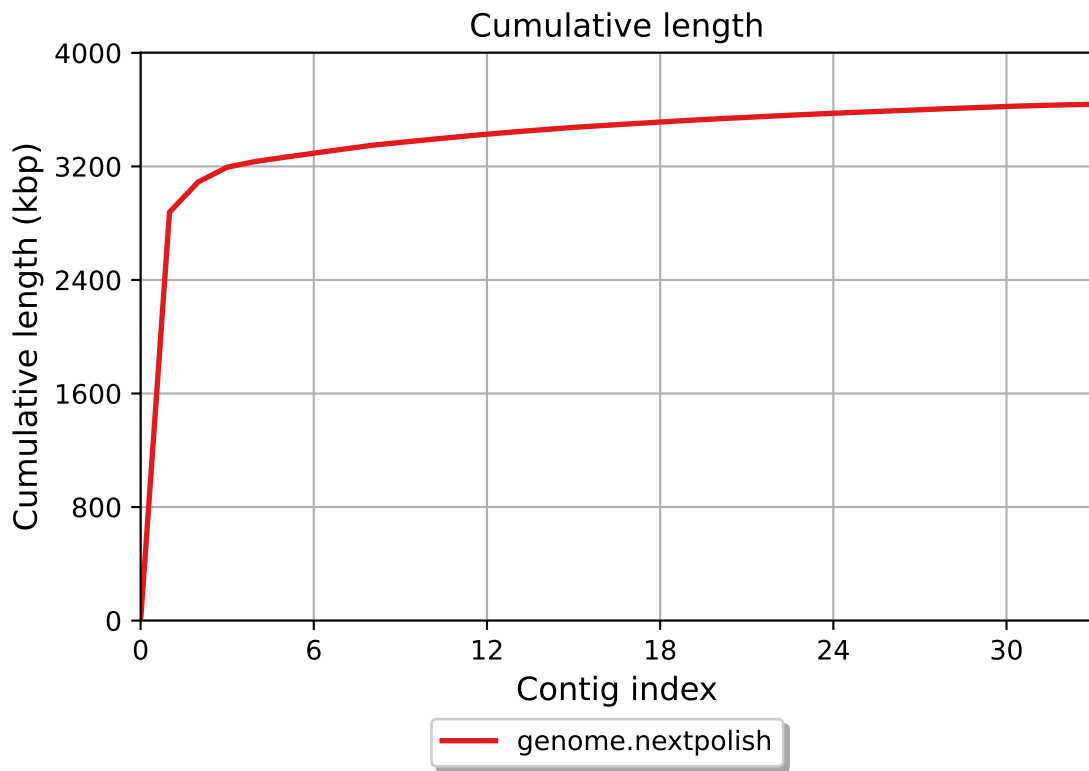
	genome.nextpolish
# contigs (>= 0 bp)	33
# contigs (>= 1000 bp)	33
# contigs (>= 5000 bp)	32
# contigs (>= 10000 bp)	22
# contigs (>= 25000 bp)	8
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	3637856
Total length (>= 1000 bp)	3637856
Total length (>= 5000 bp)	3633718
Total length (>= 10000 bp)	3555627
Total length (>= 25000 bp)	3348513
Total length (>= 50000 bp)	3194476
# contigs	33
Largest contig	2877310
Total length	3637856
GC (%)	37.73
N50	2877310
N90	28027
auN	2293688.3
L50	1
L90	6
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	18 + 0 part

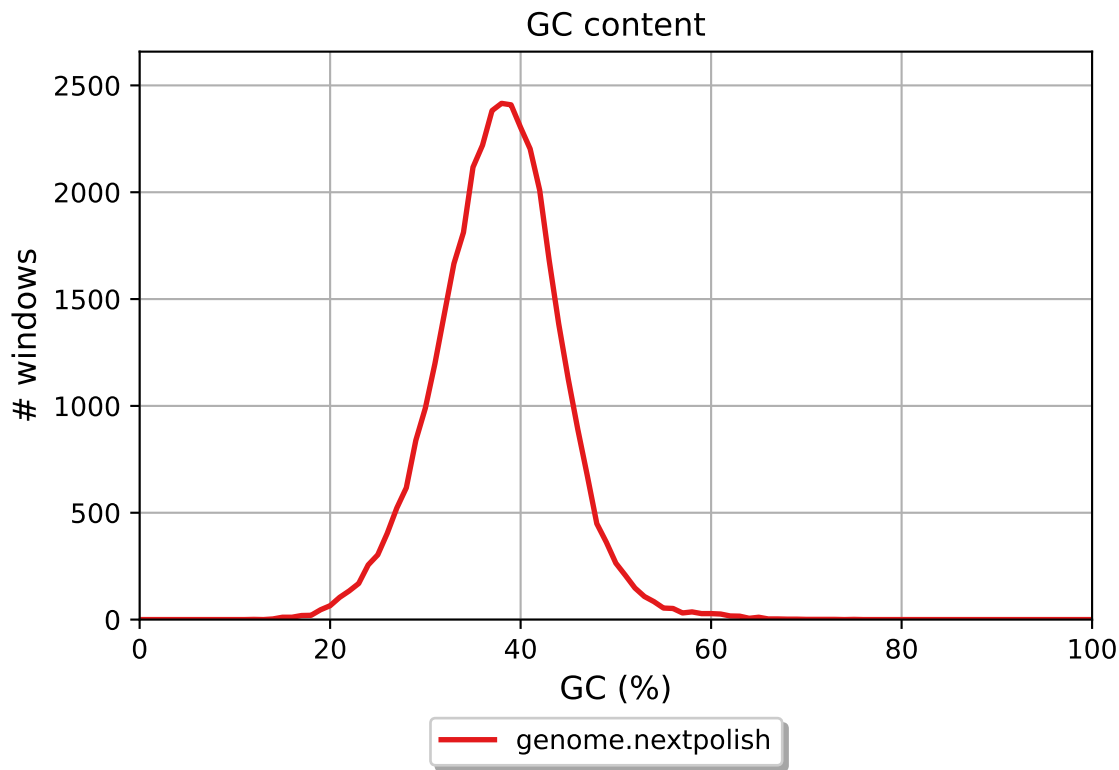
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

