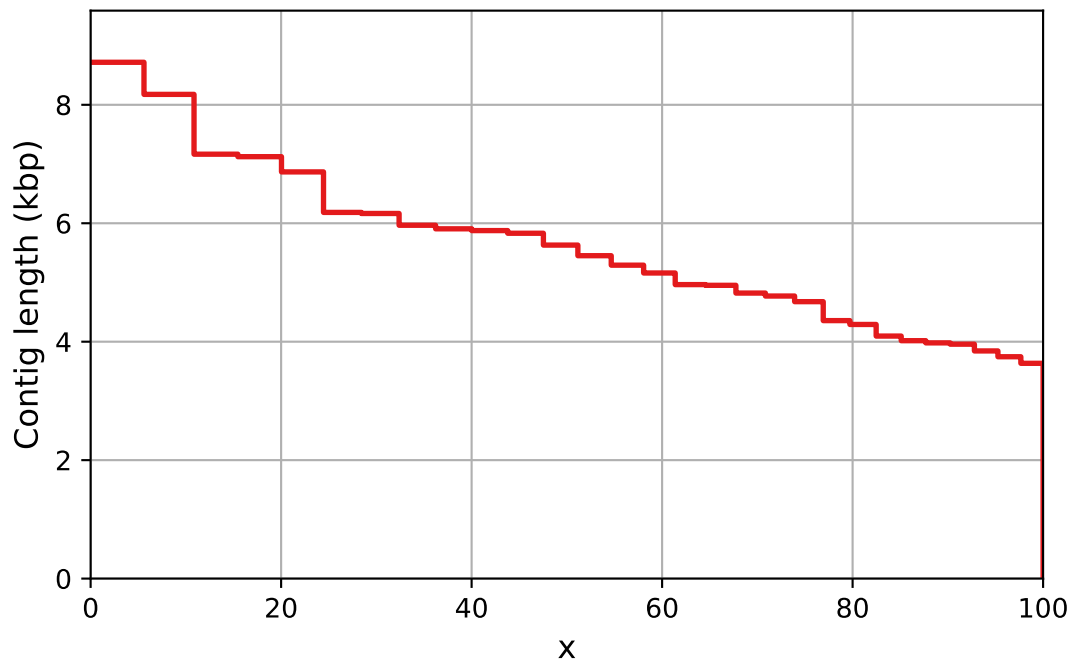


Report

	genome.nextpolish
# contigs (>= 0 bp)	29
# contigs (>= 1000 bp)	29
# contigs (>= 5000 bp)	15
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	155619
Total length (>= 1000 bp)	155619
Total length (>= 5000 bp)	95513
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	29
Largest contig	8718
Total length	155619
GC (%)	38.33
N50	5631
N90	3980
auN	5682.2
L50	12
L90	25
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 1 part

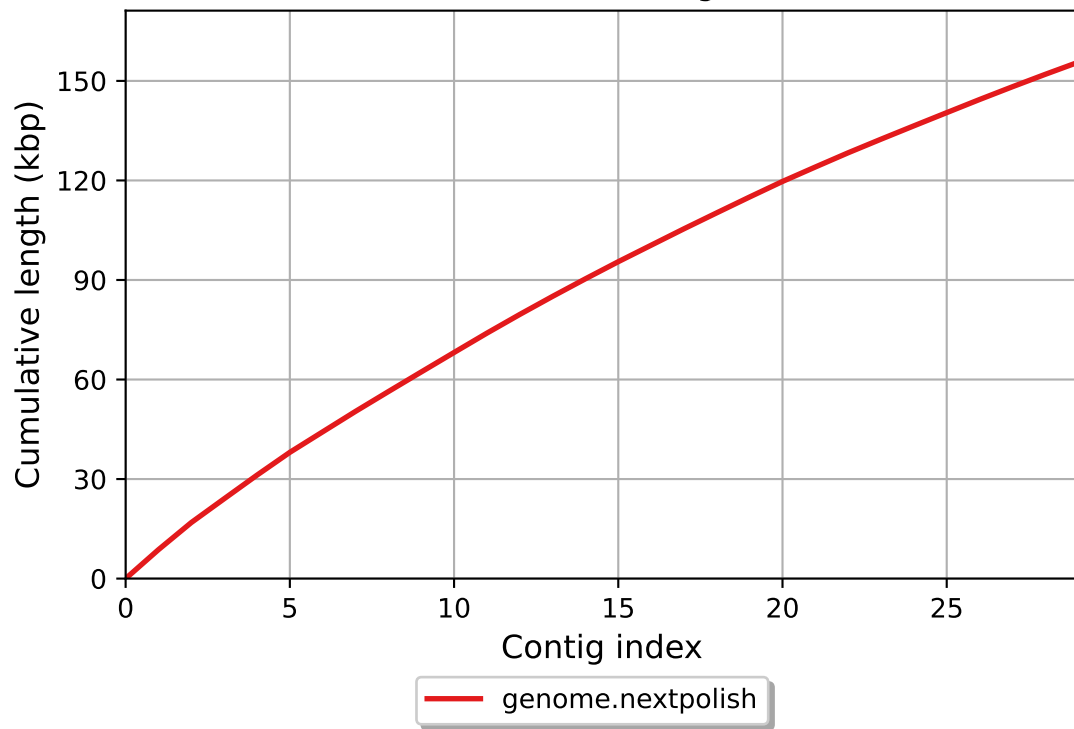
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

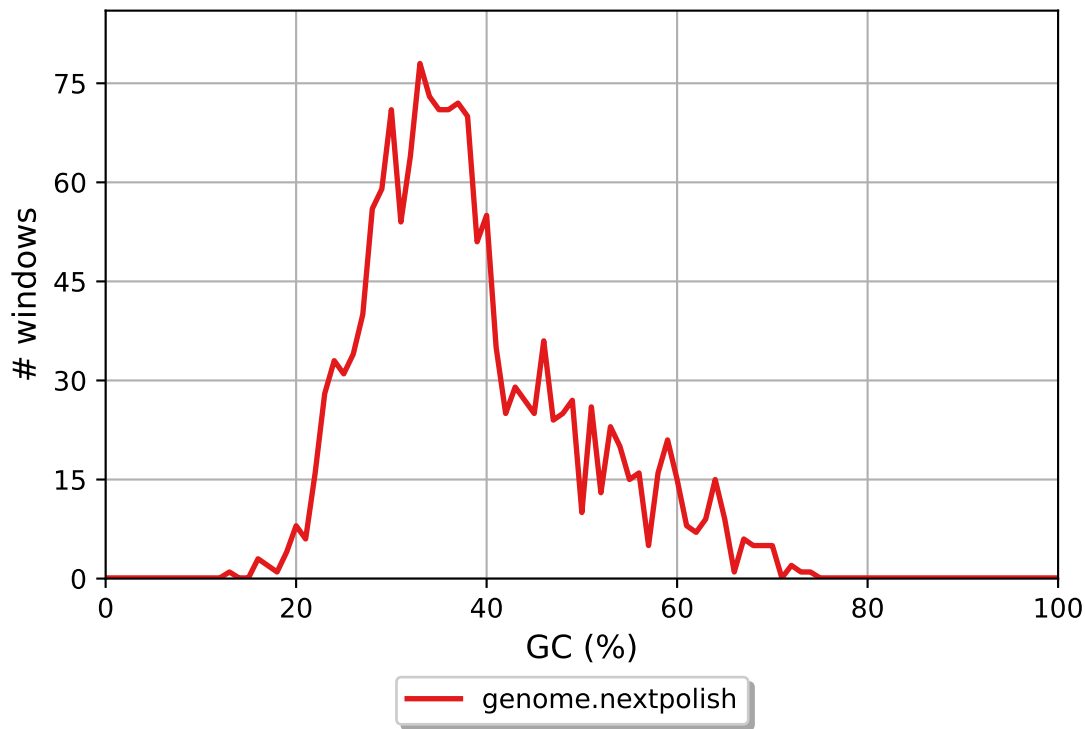


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

