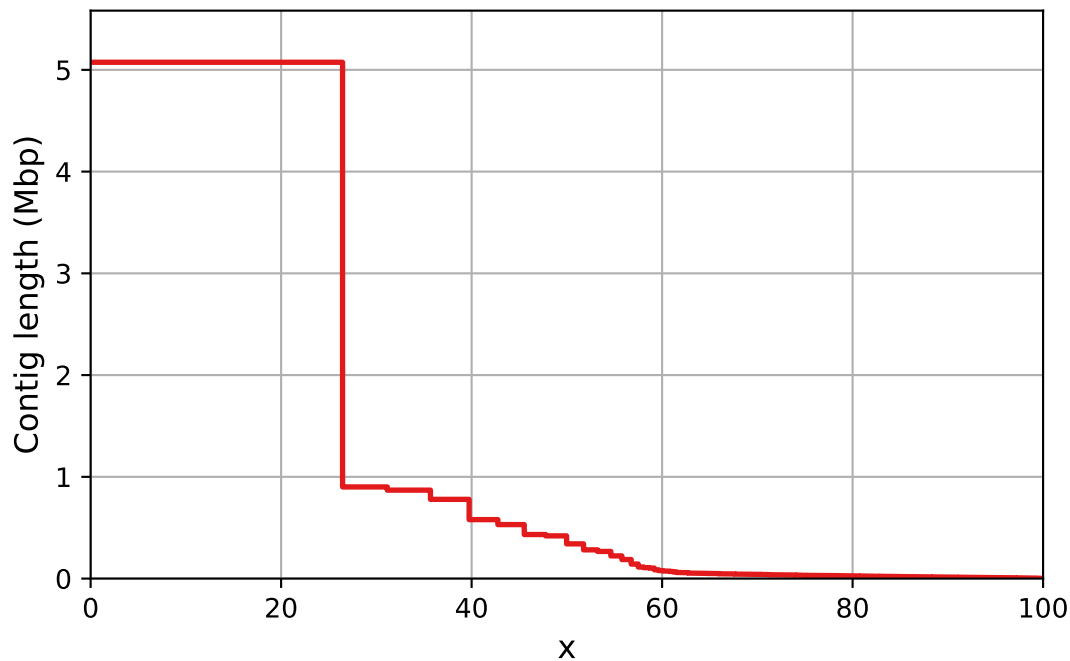


Report

	genome.nextpolish
# contigs (>= 0 bp)	451
# contigs (>= 1000 bp)	451
# contigs (>= 5000 bp)	404
# contigs (>= 10000 bp)	282
# contigs (>= 25000 bp)	125
# contigs (>= 50000 bp)	37
Total length (>= 0 bp)	19183468
Total length (>= 1000 bp)	19183468
Total length (>= 5000 bp)	19016524
Total length (>= 10000 bp)	18112085
Total length (>= 25000 bp)	15599265
Total length (>= 50000 bp)	12555014
# contigs	451
Largest contig	5074511
Total length	19183468
GC (%)	57.88
N50	341403
N90	14749
auN	1540245.2
L50	9
L90	213
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	45 + 0 part

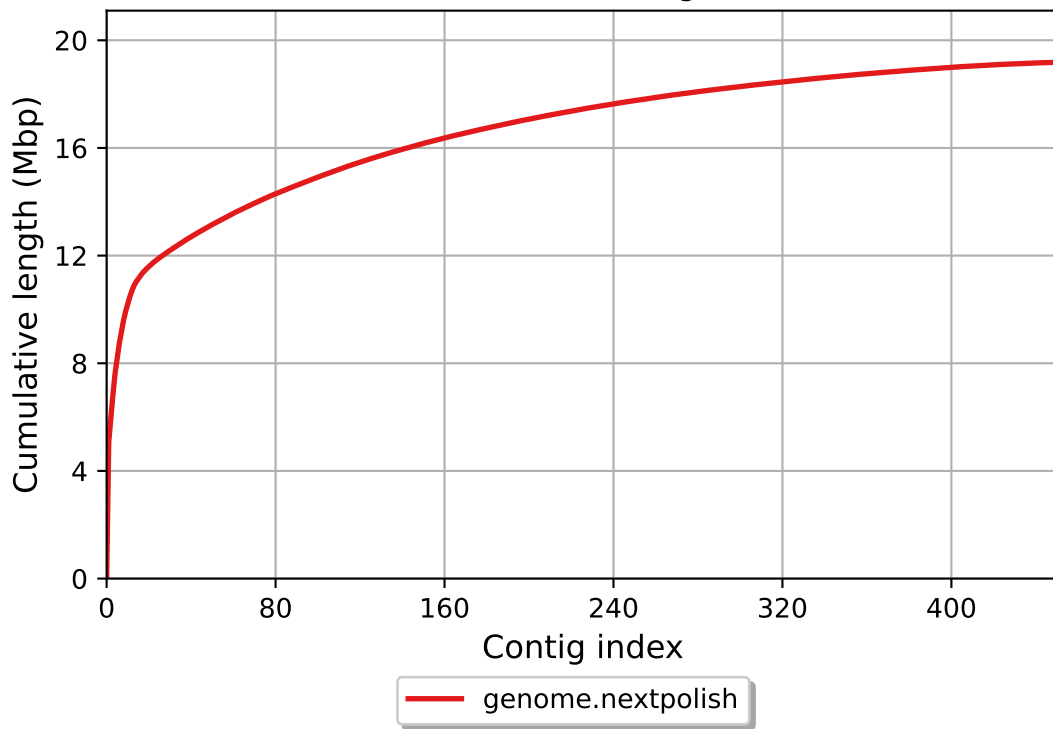
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

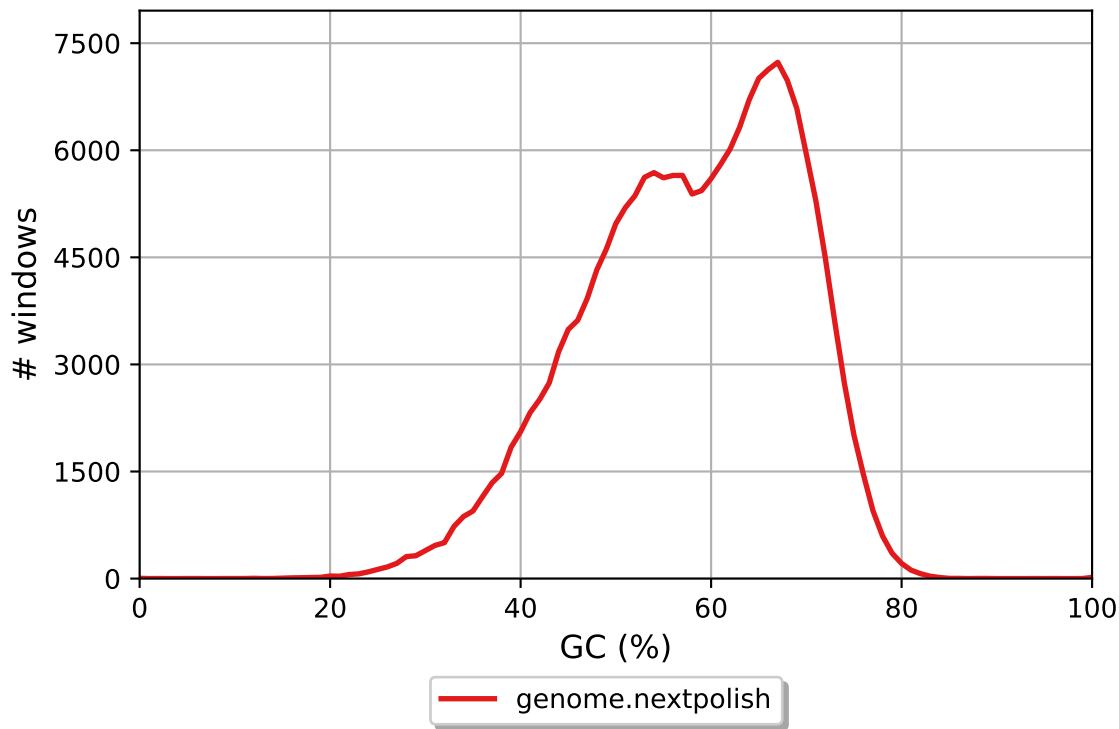


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

