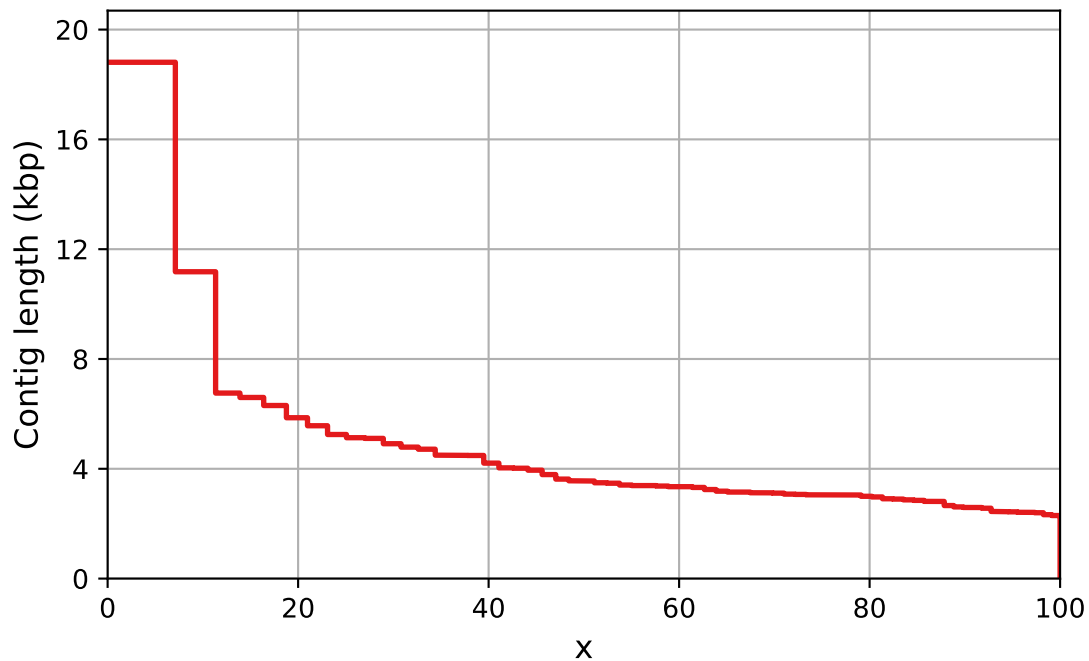


# Report

	genome.nextpolish
# contigs (>= 0 bp)	68
# contigs (>= 1000 bp)	68
# contigs (>= 5000 bp)	10
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	264502
Total length (>= 1000 bp)	264502
Total length (>= 5000 bp)	76564
Total length (>= 10000 bp)	29987
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	68
Largest contig	18809
Total length	264502
GC (%)	33.93
N50	3554
N90	2591
auN	5242.6
L50	24
L90	58
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 0 part

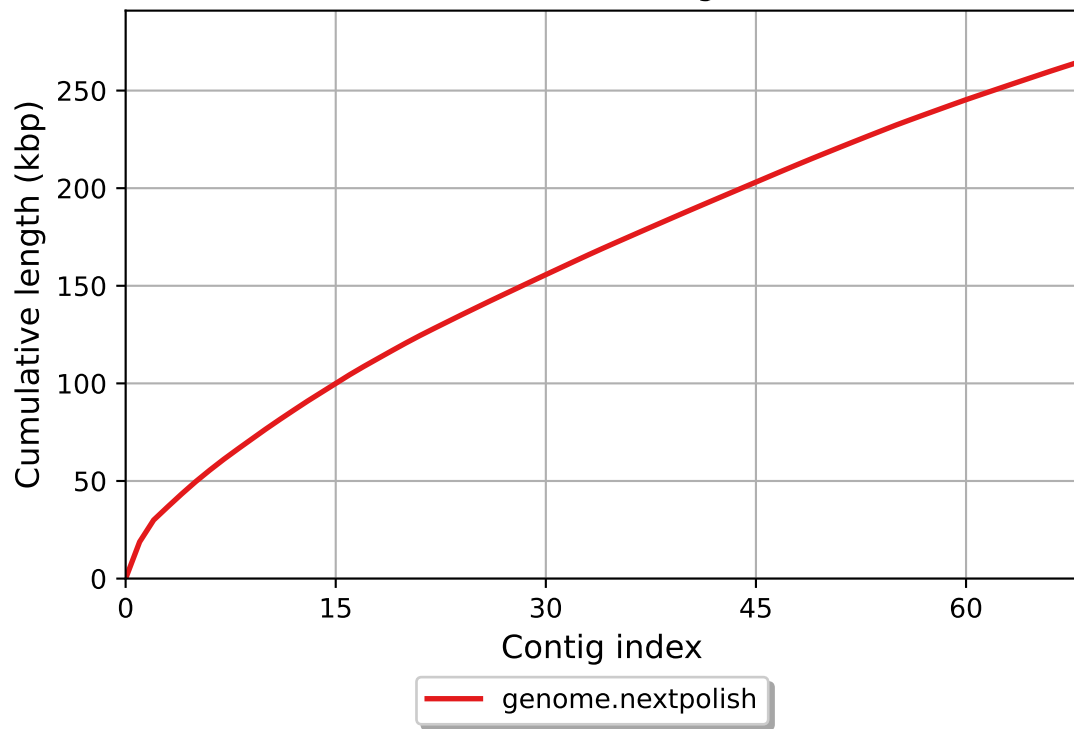
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

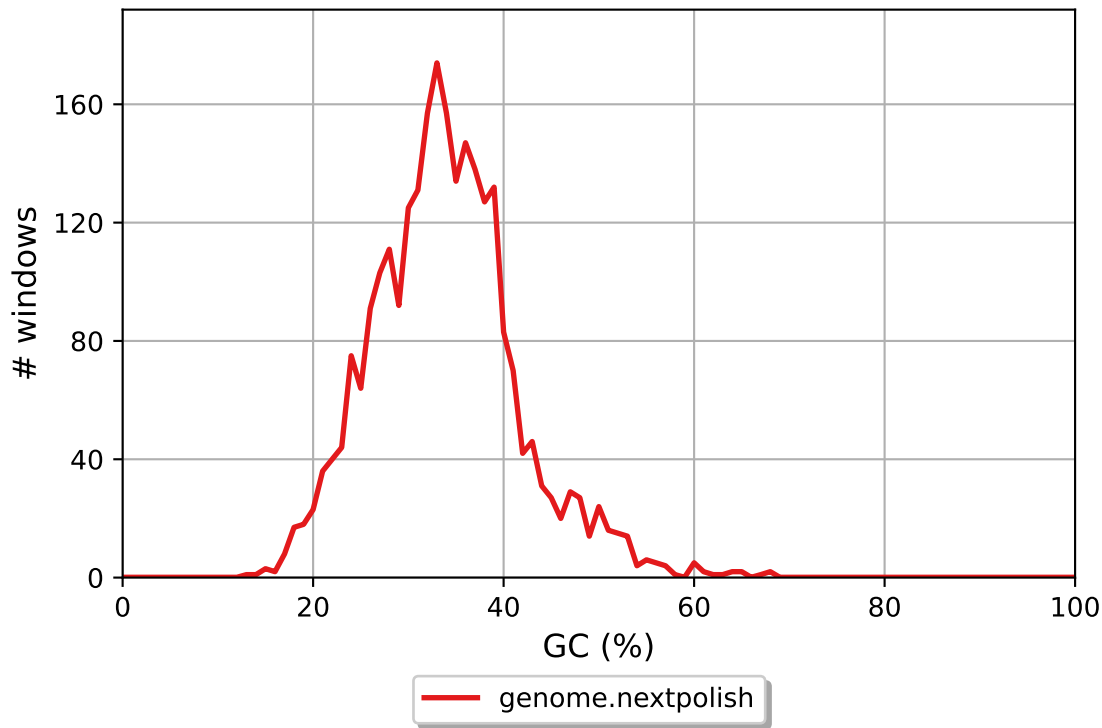


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

