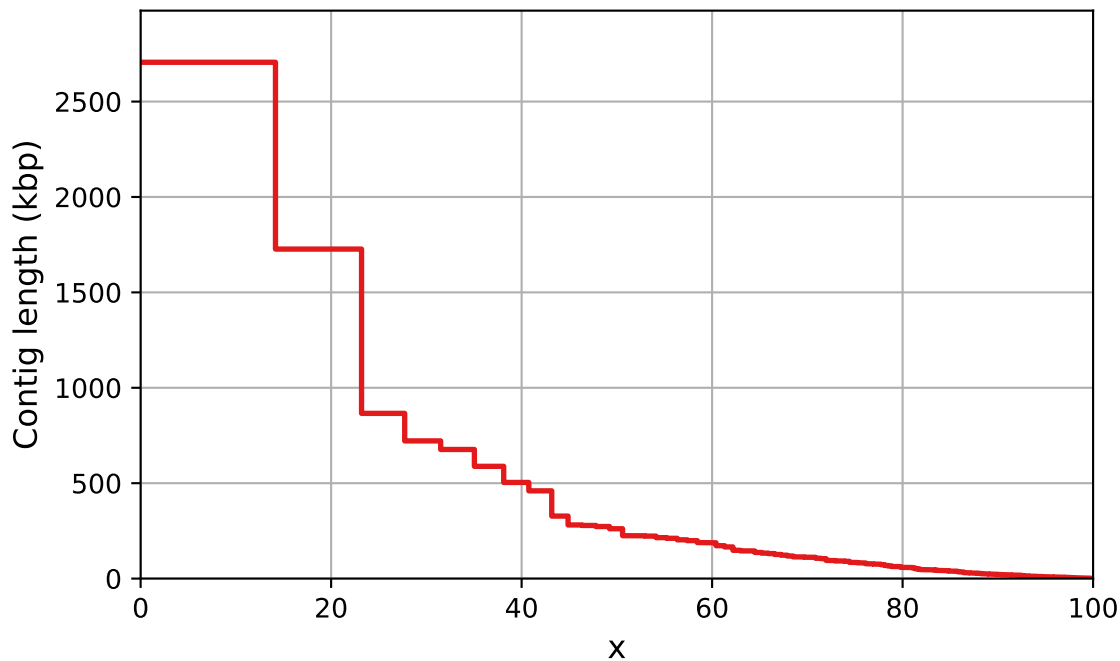


Report

	genome.nextpolish
# contigs (>= 0 bp)	409
# contigs (>= 1000 bp)	404
# contigs (>= 5000 bp)	253
# contigs (>= 10000 bp)	171
# contigs (>= 25000 bp)	102
# contigs (>= 50000 bp)	64
Total length (>= 0 bp)	19112074
Total length (>= 1000 bp)	19108082
Total length (>= 5000 bp)	18717596
Total length (>= 10000 bp)	18102923
Total length (>= 25000 bp)	16967318
Total length (>= 50000 bp)	15595365
# contigs	409
Largest contig	2705999
Total length	19112074
GC (%)	38.52
N50	261323
N90	22013
auN	740634.8
L50	13
L90	112
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	69 + 6 part

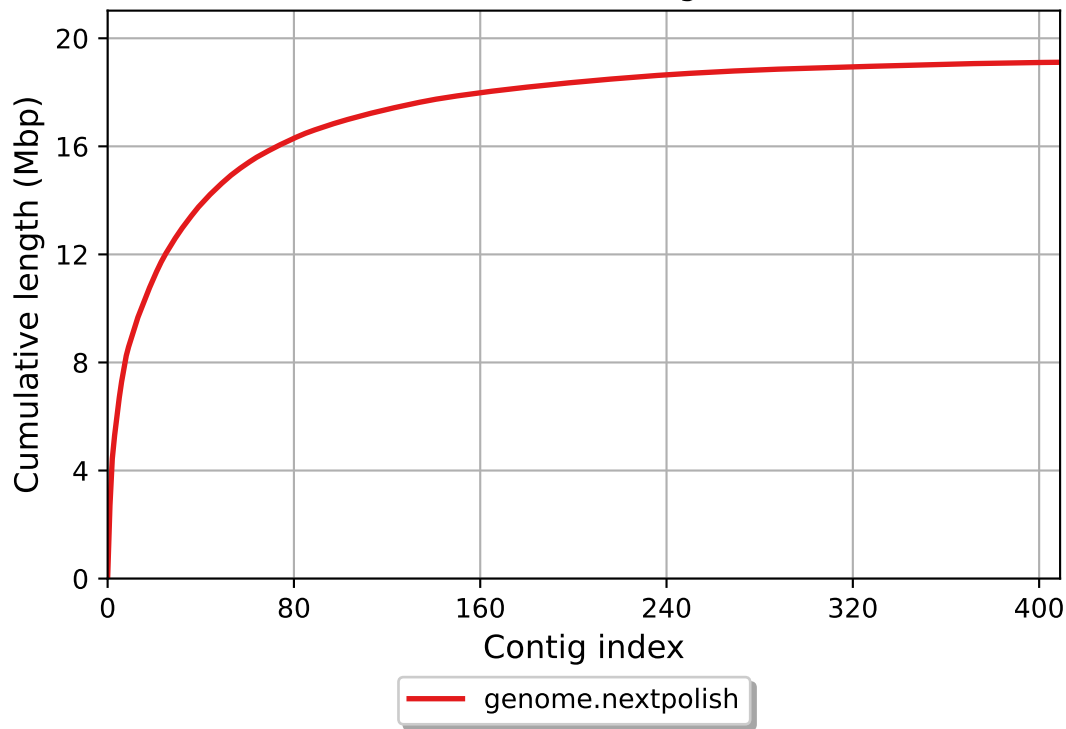
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

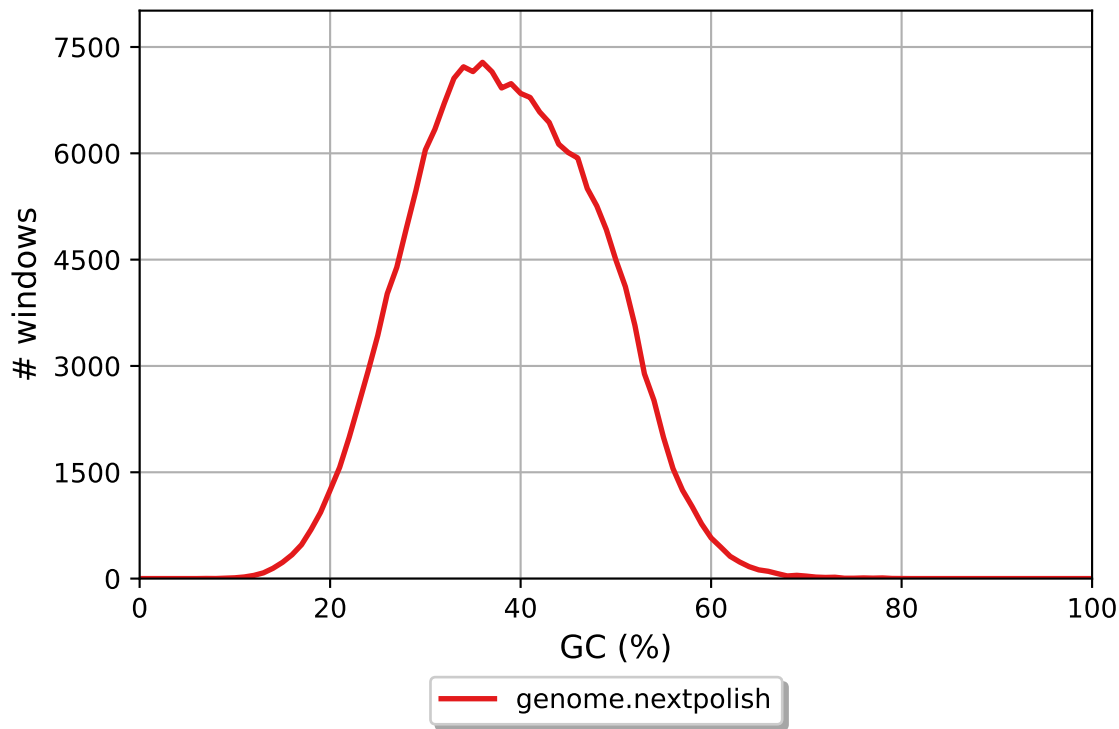


genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

