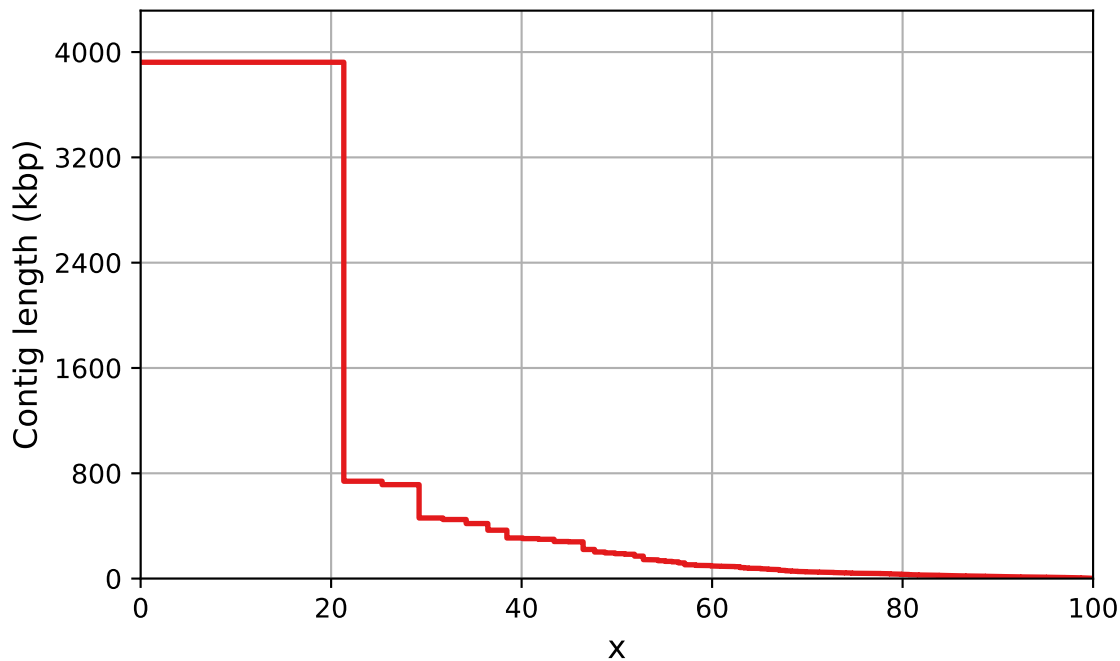


Report

	genome.nextpolish
# contigs (>= 0 bp)	402
# contigs (>= 1000 bp)	402
# contigs (>= 5000 bp)	355
# contigs (>= 10000 bp)	260
# contigs (>= 25000 bp)	124
# contigs (>= 50000 bp)	58
Total length (>= 0 bp)	18387671
Total length (>= 1000 bp)	18387671
Total length (>= 5000 bp)	18229740
Total length (>= 10000 bp)	17524263
Total length (>= 25000 bp)	15404907
Total length (>= 50000 bp)	13043113
# contigs	402
Largest contig	3922506
Total length	18387671
GC (%)	59.75
N50	189263
N90	15554
auN	993233.5
L50	16
L90	183
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	42 + 1 part

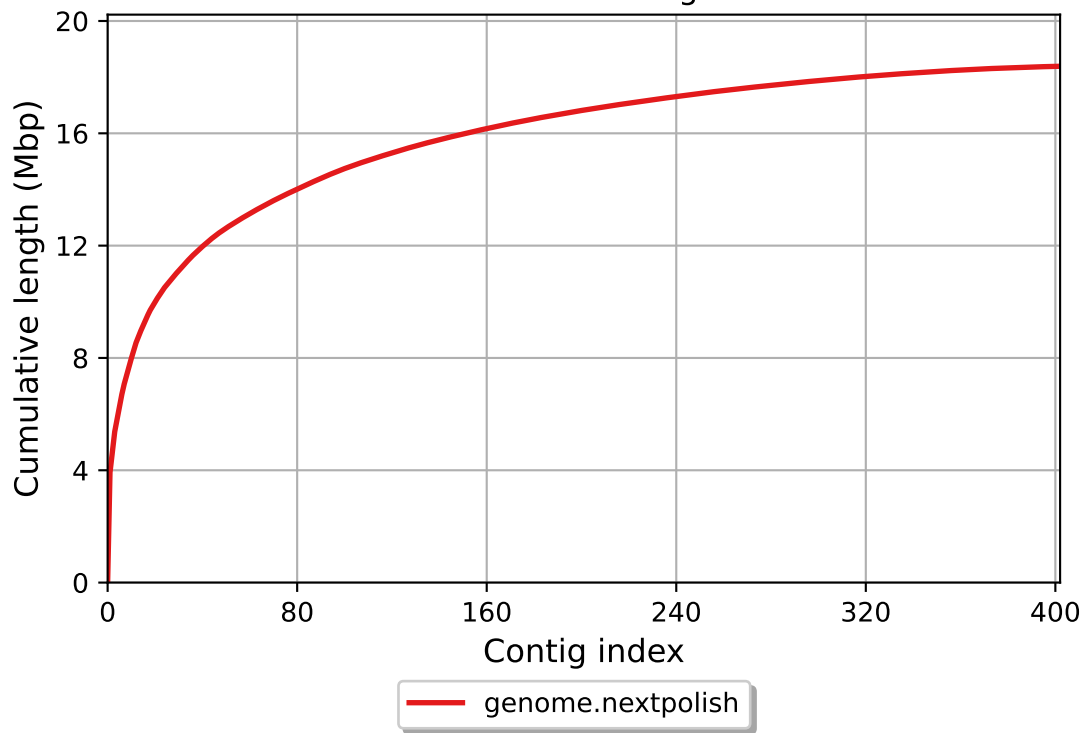
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

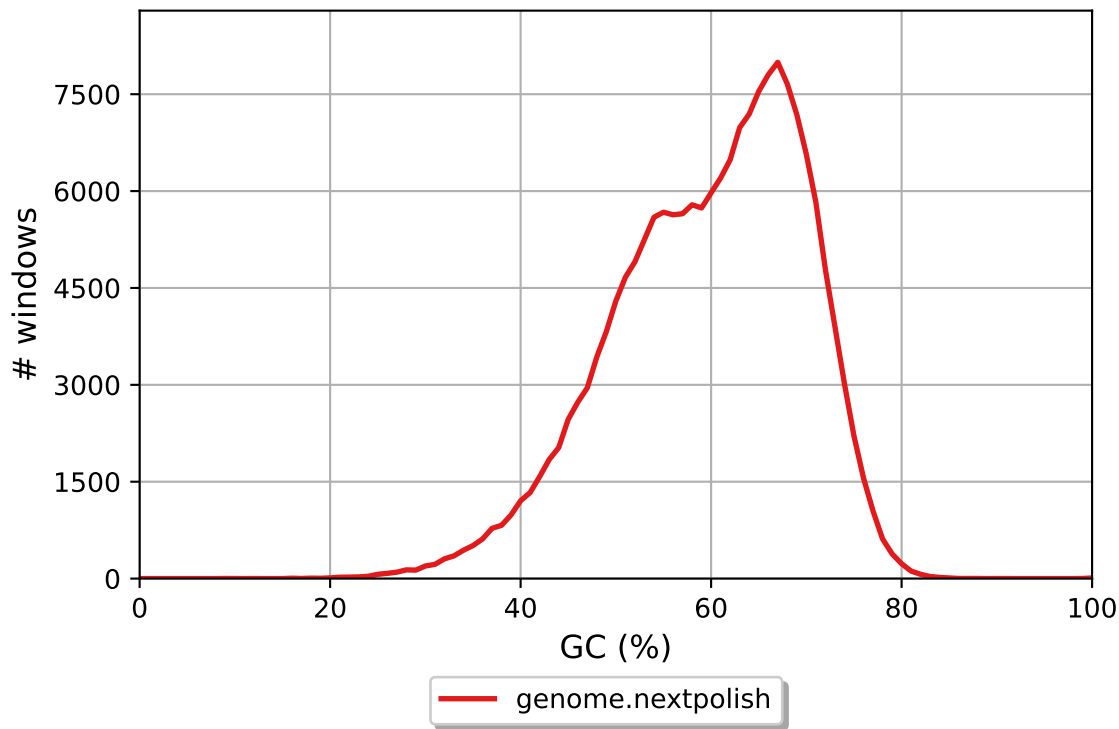


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

