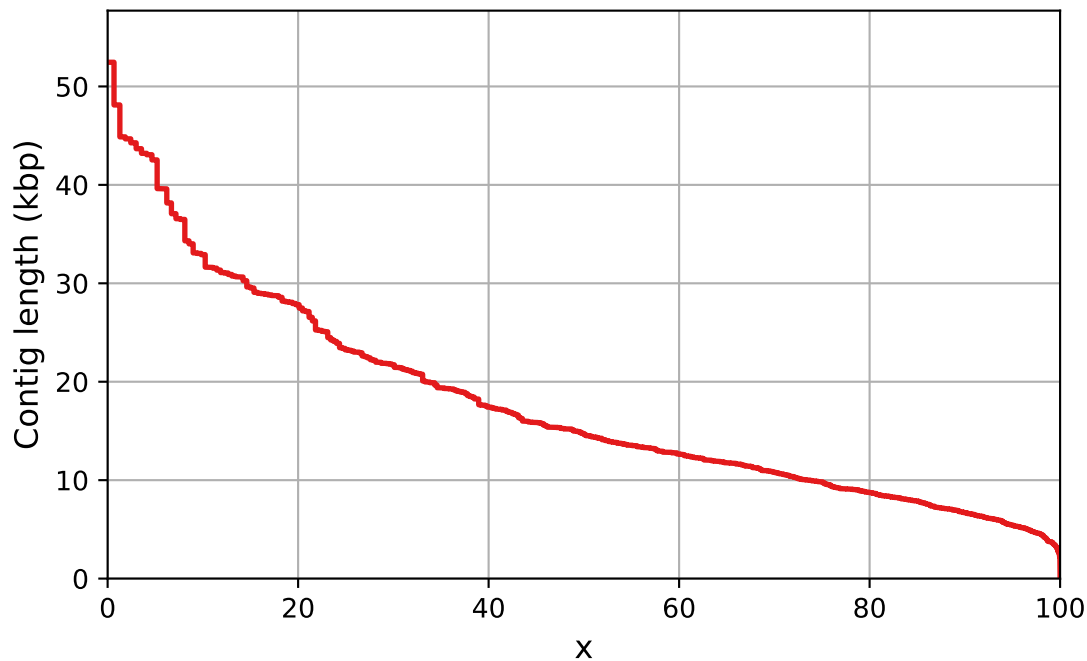


# Report

	genome.nextpolish
# contigs (>= 0 bp)	631
# contigs (>= 1000 bp)	630
# contigs (>= 5000 bp)	562
# contigs (>= 10000 bp)	320
# contigs (>= 25000 bp)	55
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	7825609
Total length (>= 1000 bp)	7824776
Total length (>= 5000 bp)	7555911
Total length (>= 10000 bp)	5766423
Total length (>= 25000 bp)	1808539
Total length (>= 50000 bp)	52456
# contigs	631
Largest contig	52456
Total length	7825609
GC (%)	46.32
N50	14733
N90	6703
auN	17842.4
L50	168
L90	475
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	17 + 2 part

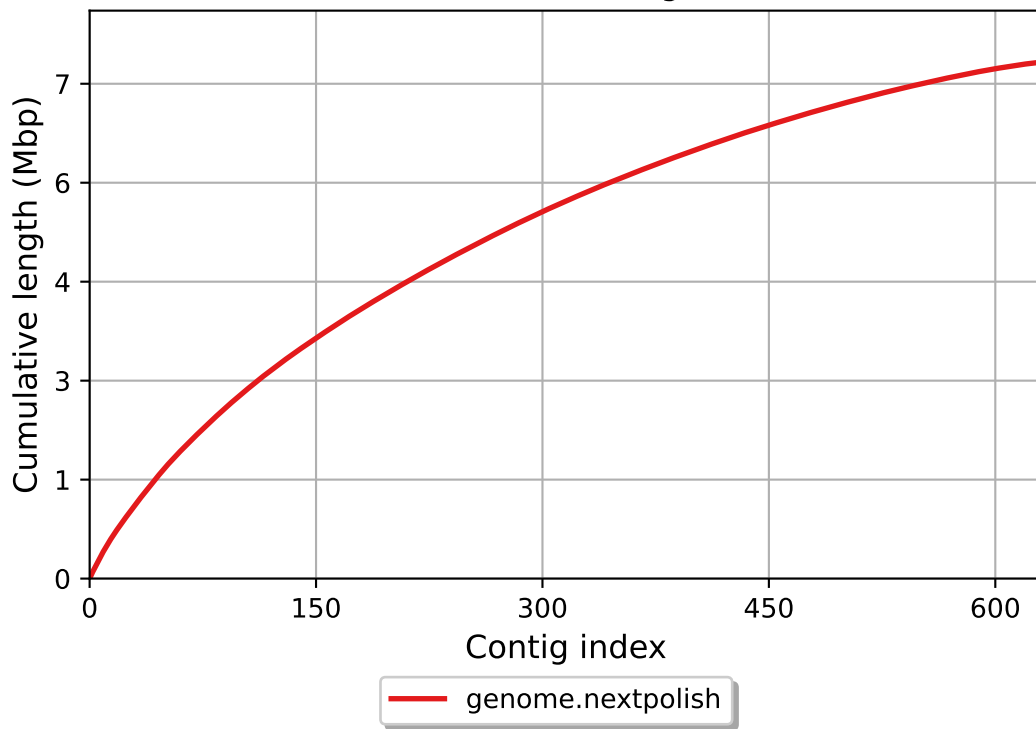
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

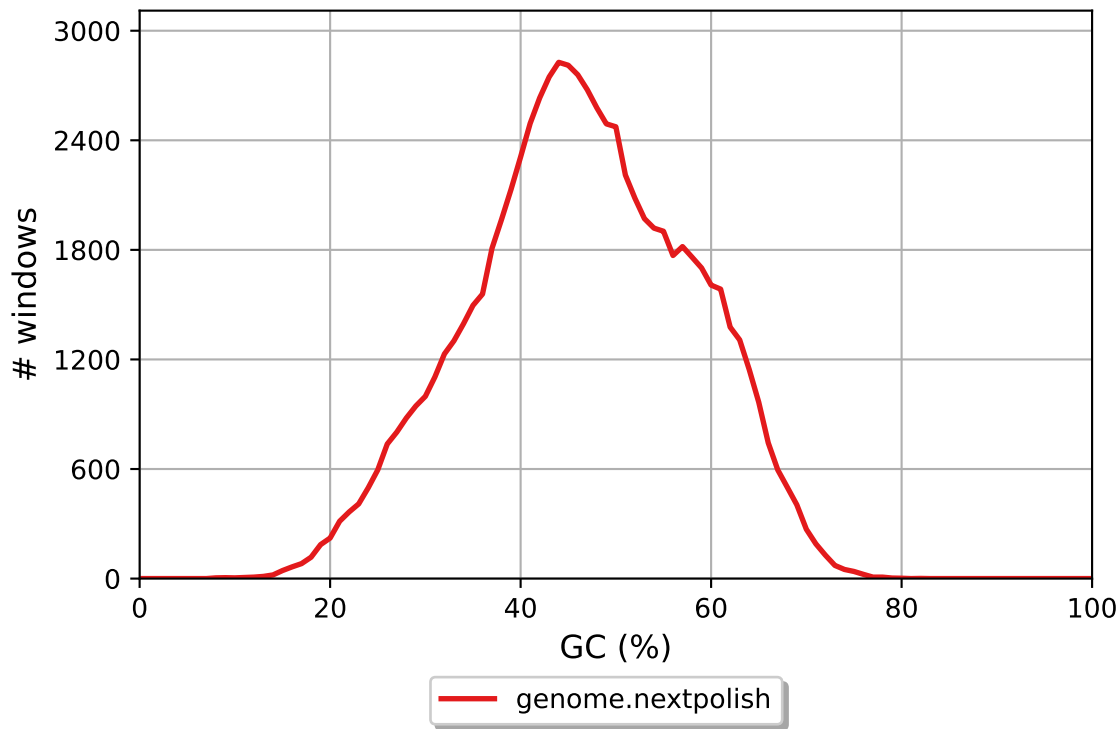


— genome.nextpolish

Cumulative length



# GC content



genome.nextpolish GC content

