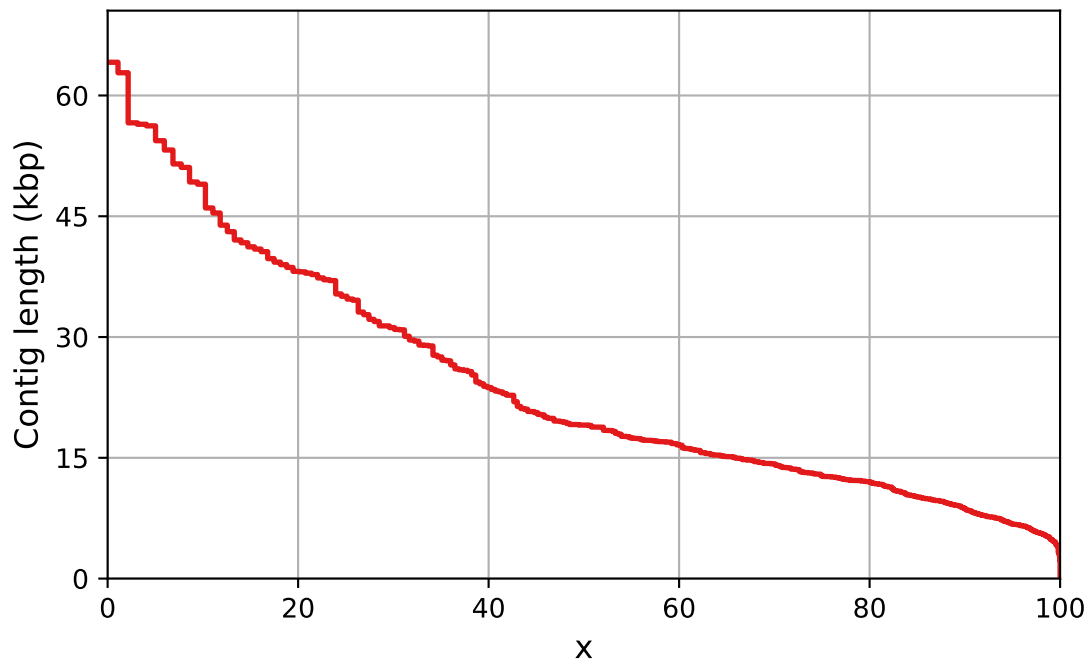


# Report

	genome.nextpolish
# contigs (>= 0 bp)	359
# contigs (>= 1000 bp)	359
# contigs (>= 5000 bp)	344
# contigs (>= 10000 bp)	239
# contigs (>= 25000 bp)	60
# contigs (>= 50000 bp)	9
Total length (>= 0 bp)	5888042
Total length (>= 1000 bp)	5888042
Total length (>= 5000 bp)	5827086
Total length (>= 10000 bp)	5035430
Total length (>= 25000 bp)	2275967
Total length (>= 50000 bp)	506381
# contigs	359
Largest contig	64124
Total length	5888042
GC (%)	53.40
N50	19067
N90	8758
auN	24356.0
L50	92
L90	267
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	18 + 2 part

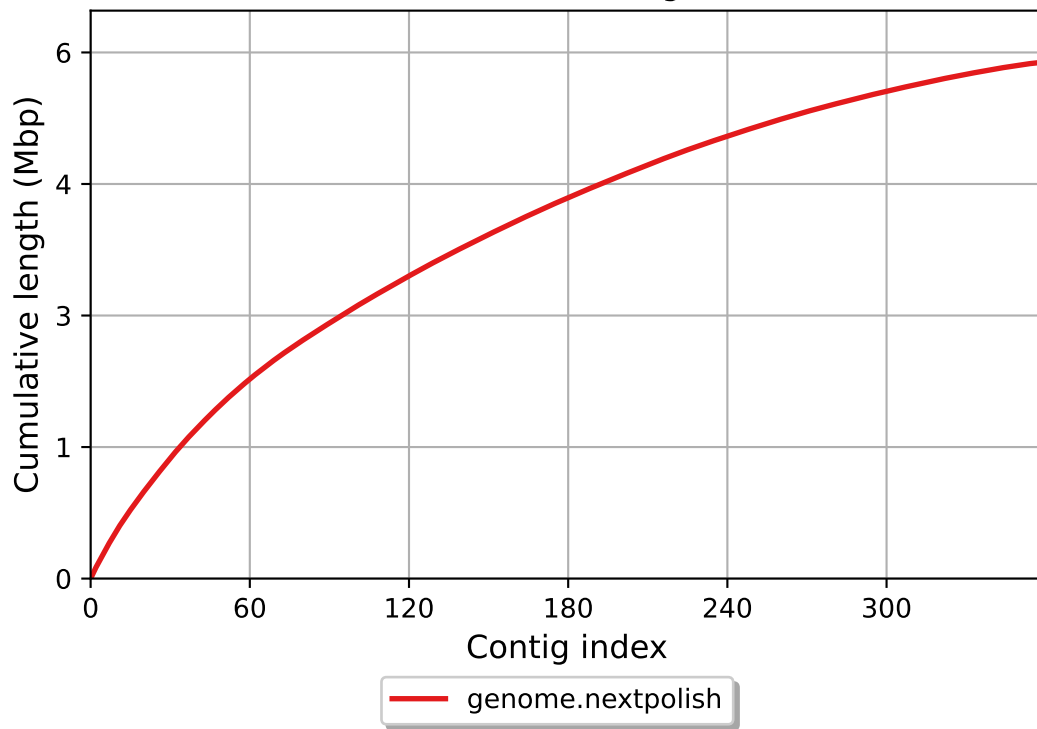
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

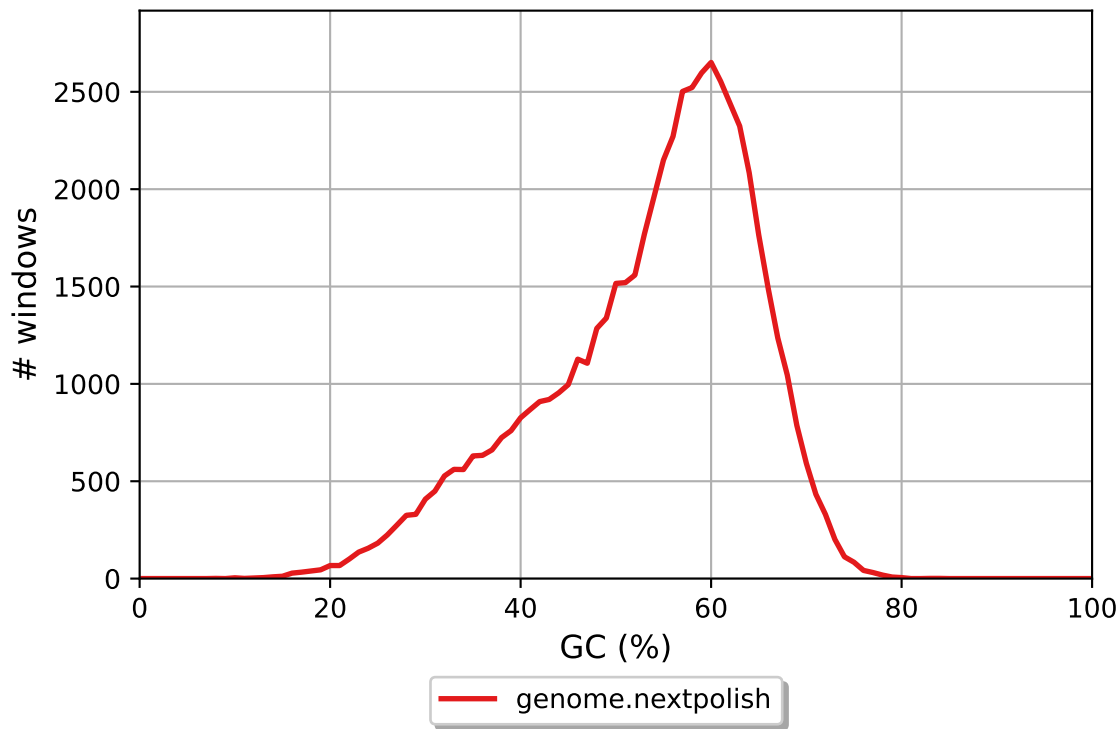


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

