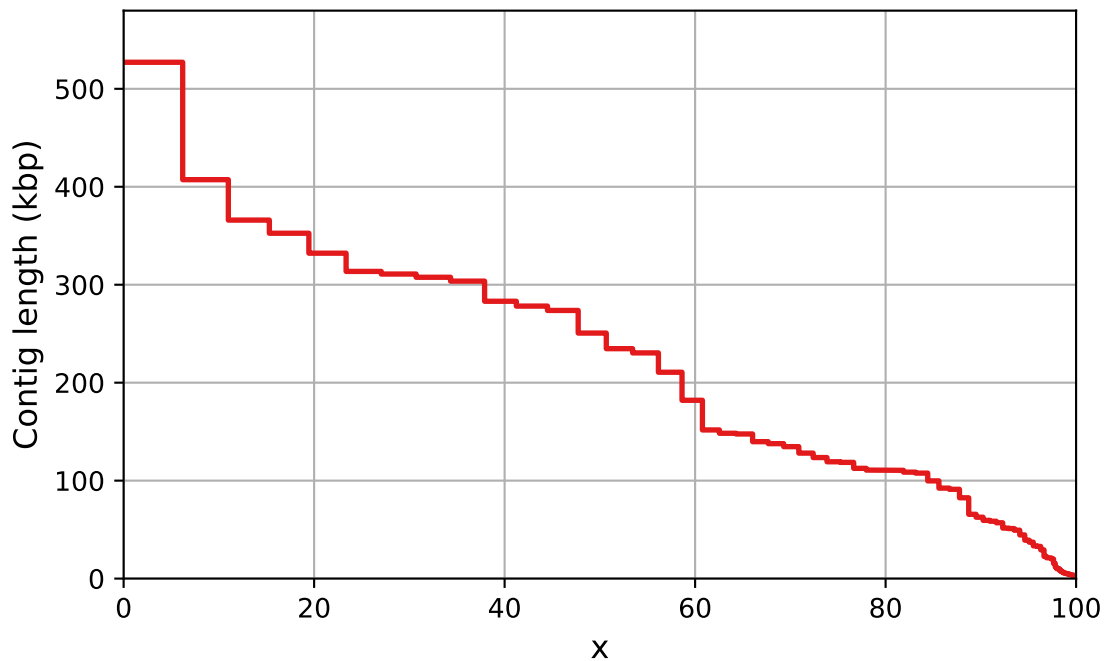


# Report

	genome.nextpolish
# contigs (>= 0 bp)	100
# contigs (>= 1000 bp)	91
# contigs (>= 5000 bp)	70
# contigs (>= 10000 bp)	59
# contigs (>= 25000 bp)	51
# contigs (>= 50000 bp)	44
Total length (>= 0 bp)	8498951
Total length (>= 1000 bp)	8493100
Total length (>= 5000 bp)	8416790
Total length (>= 10000 bp)	8344876
Total length (>= 25000 bp)	8212238
Total length (>= 50000 bp)	7946313
# contigs	99
Largest contig	527101
Total length	8498509
GC (%)	57.81
N50	250639
N90	62661
auN	235599.5
L50	13
L90	39
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	28 + 7 part

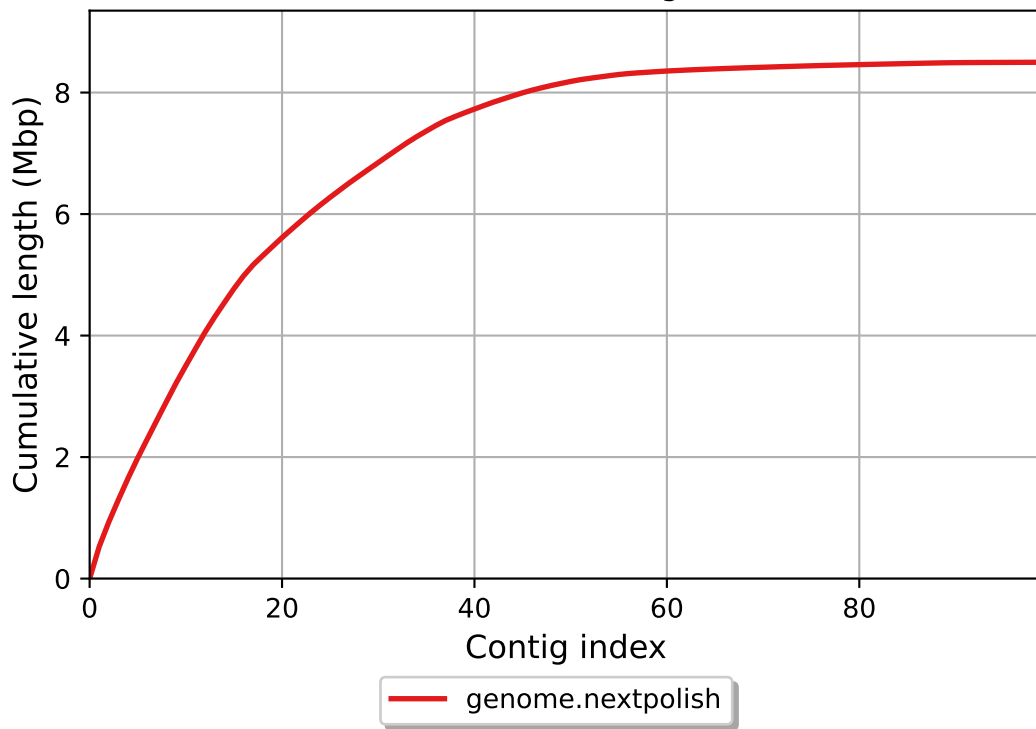
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

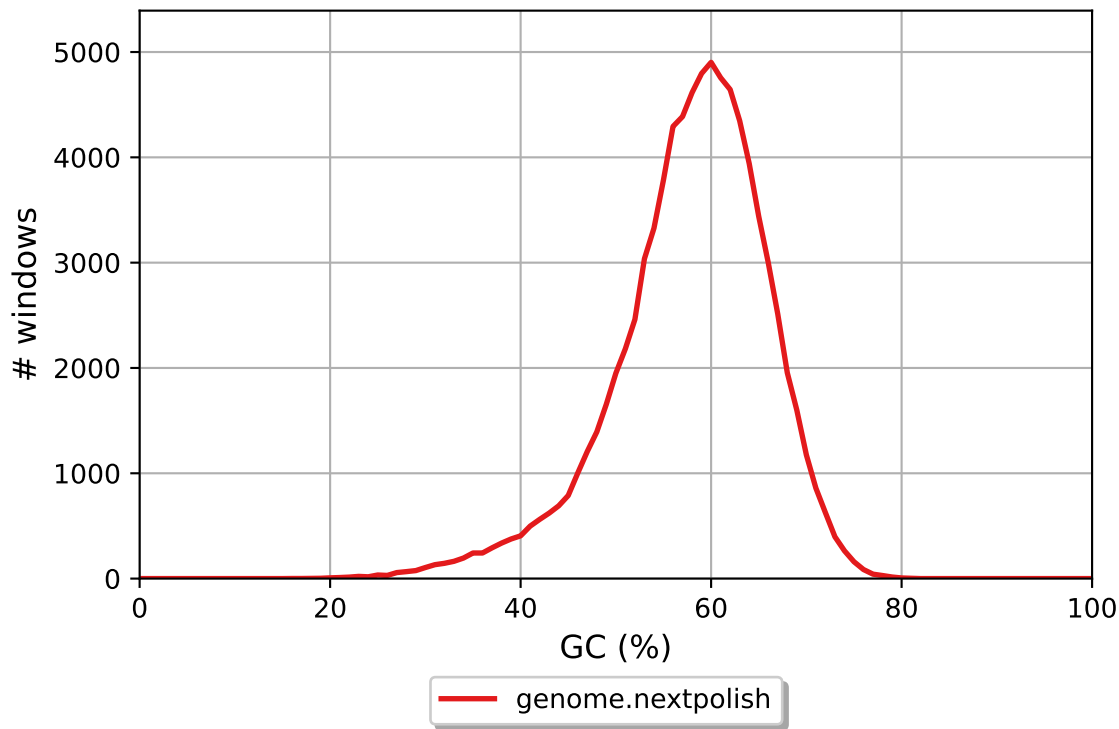


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

