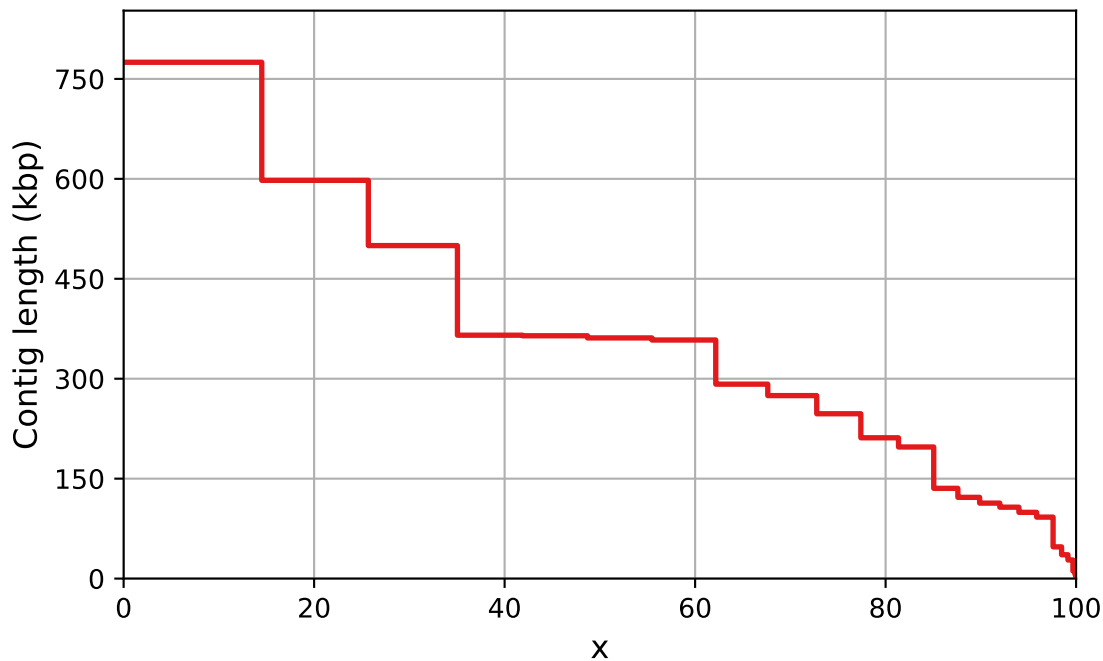


# Report

	genome.nextpolish
# contigs (>= 0 bp)	23
# contigs (>= 1000 bp)	23
# contigs (>= 5000 bp)	23
# contigs (>= 10000 bp)	22
# contigs (>= 25000 bp)	21
# contigs (>= 50000 bp)	18
Total length (>= 0 bp)	5343106
Total length (>= 1000 bp)	5343106
Total length (>= 5000 bp)	5343106
Total length (>= 10000 bp)	5336736
Total length (>= 25000 bp)	5325314
Total length (>= 50000 bp)	5213998
# contigs	23
Largest contig	774977
Total length	5343106
GC (%)	57.45
N50	361238
N90	113282
auN	396526.3
L50	6
L90	15
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	23 + 0 part

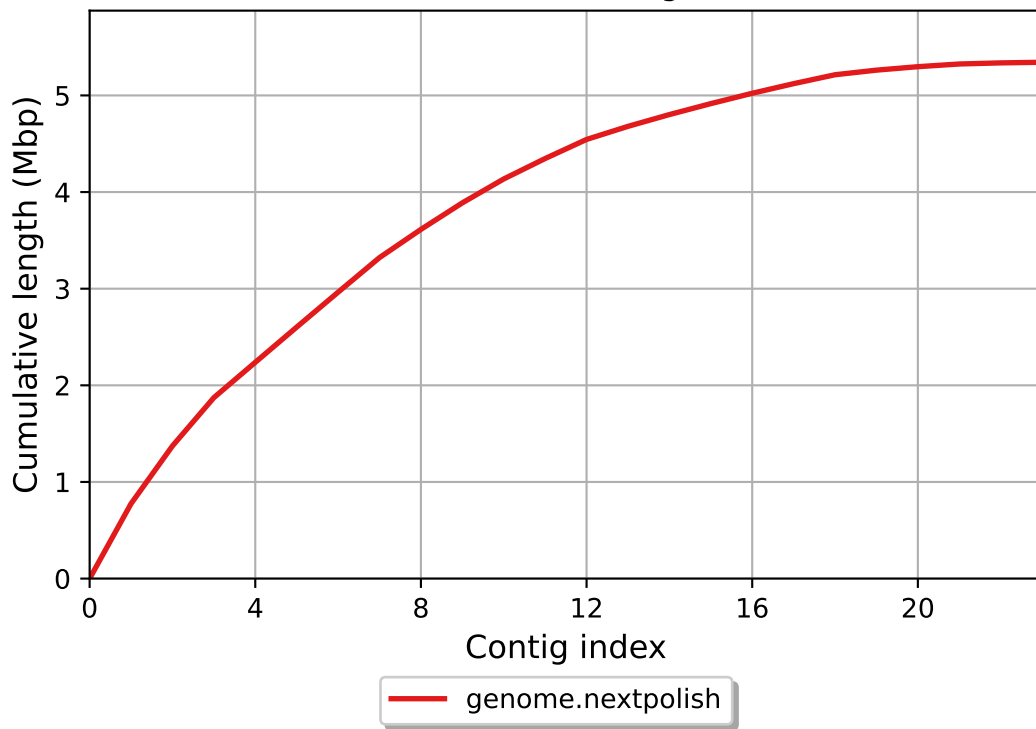
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

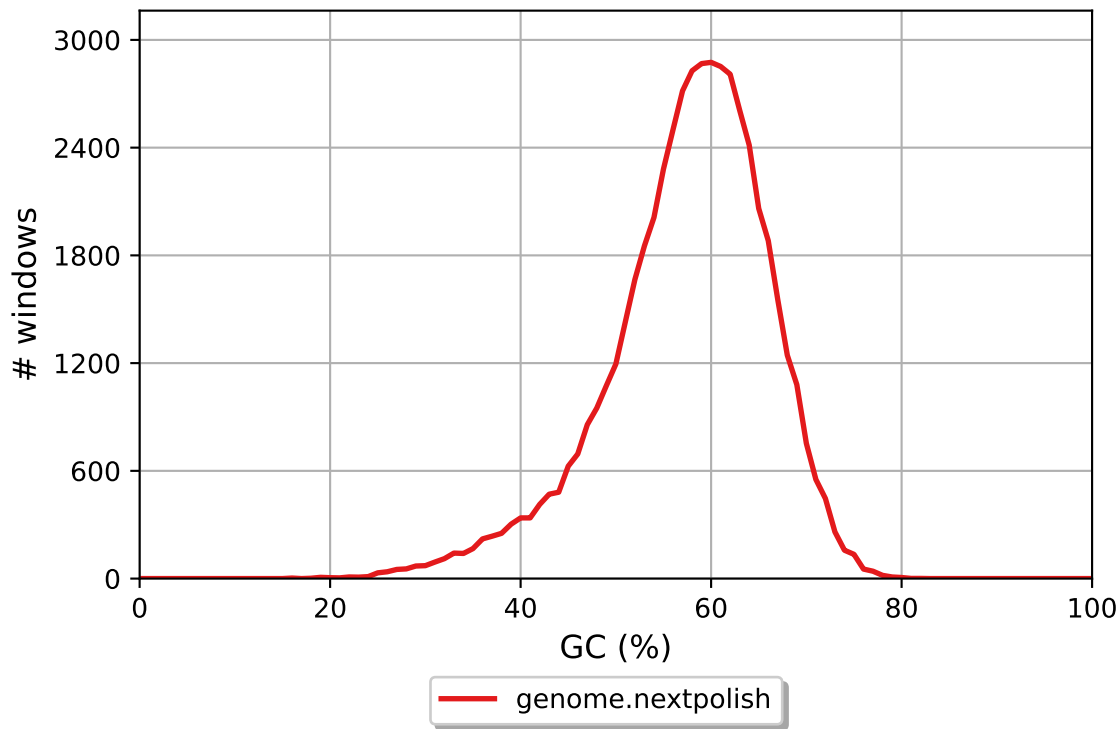


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

