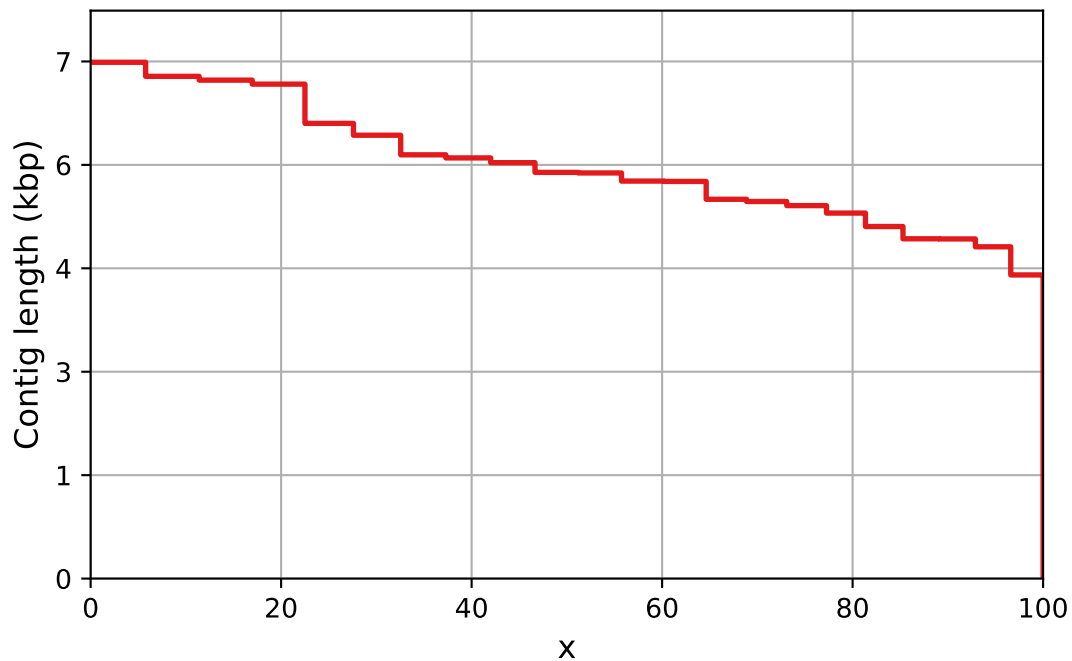


Report

	genome.nextpolish
# contigs (>= 0 bp)	22
# contigs (>= 1000 bp)	22
# contigs (>= 5000 bp)	18
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	129638
Total length (>= 1000 bp)	129638
Total length (>= 5000 bp)	110568
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	22
Largest contig	7488
Total length	129638
GC (%)	55.58
N50	5891
N90	4926
auN	6013.3
L50	10
L90	20
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 0 part

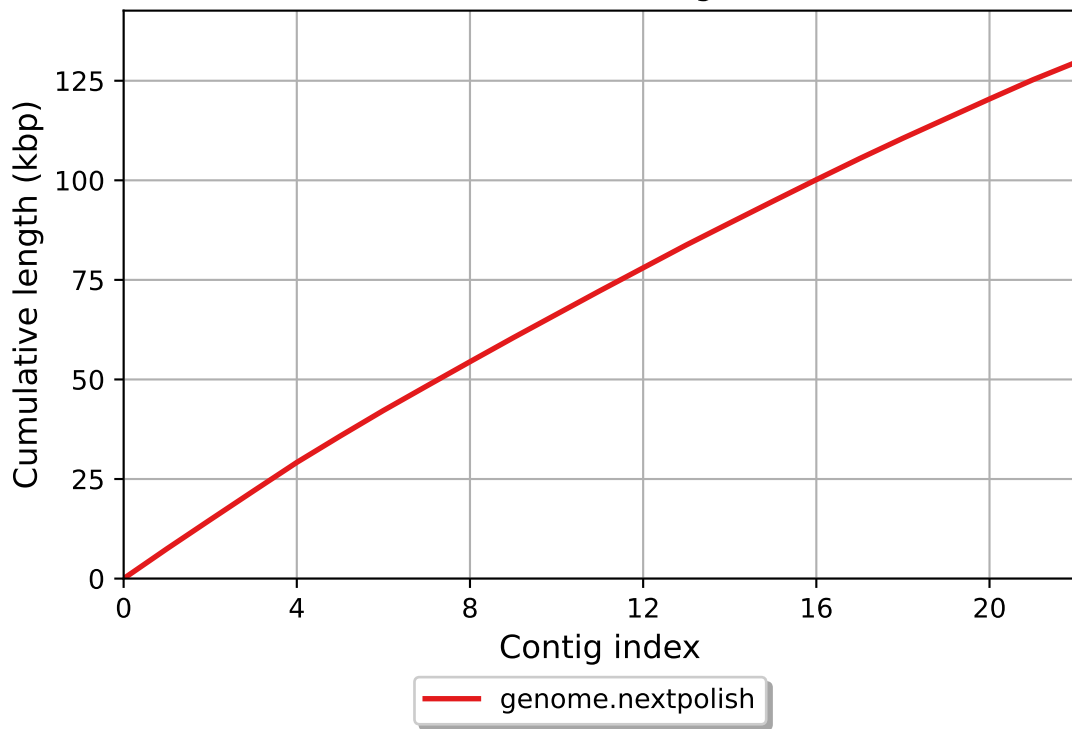
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

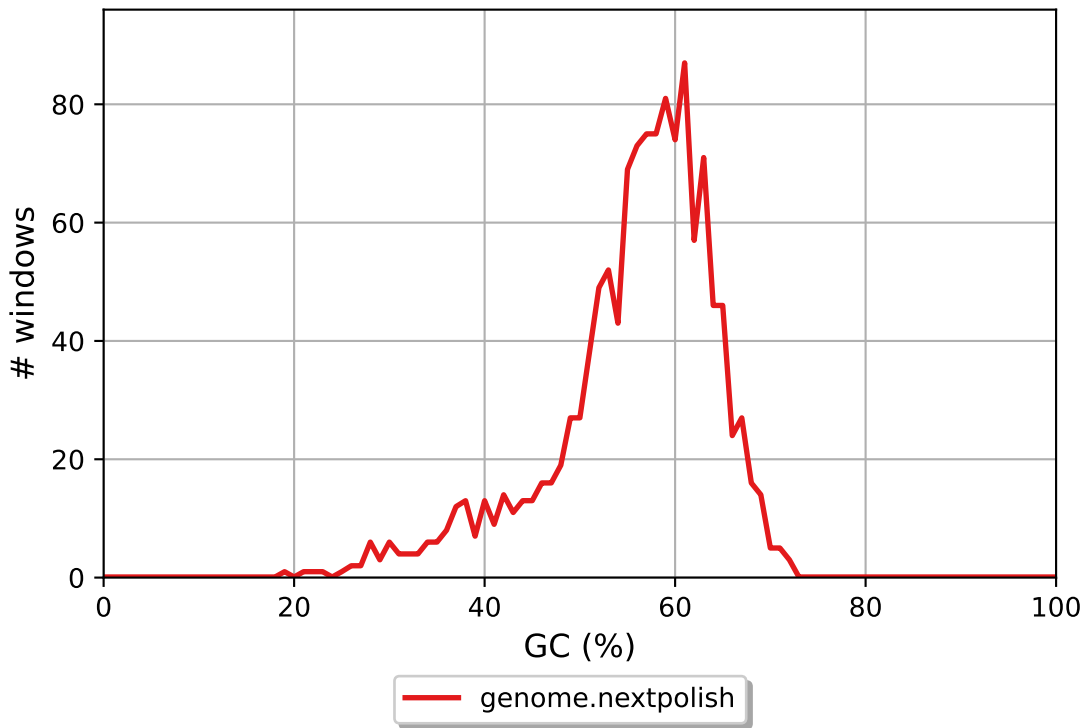


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

