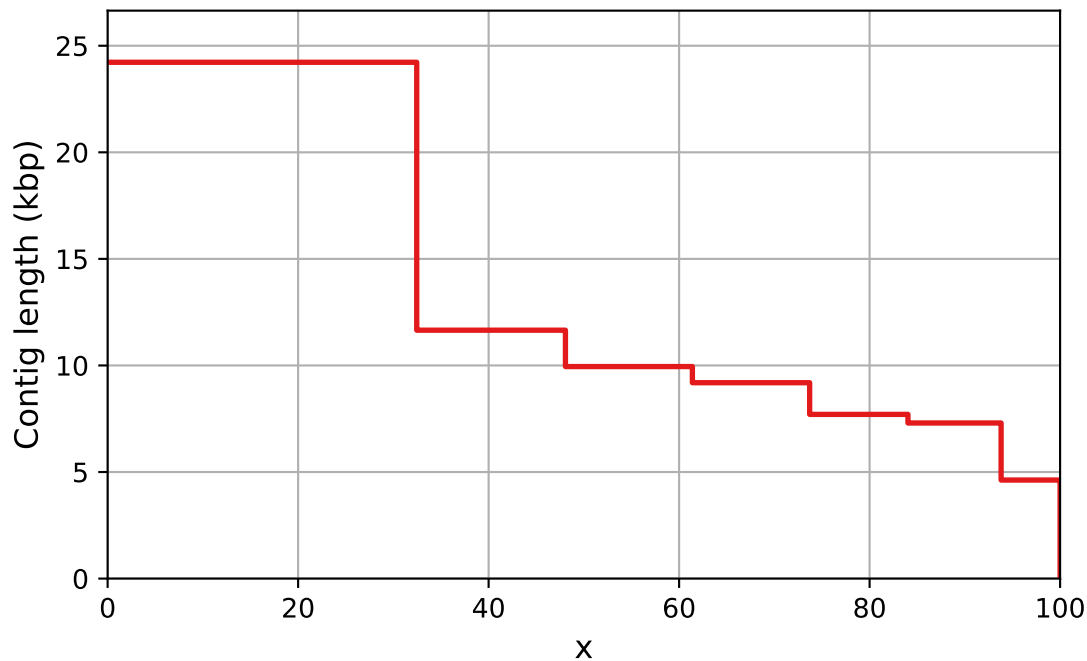


# Report

	genome.nextpolish
# contigs ( $\geq 0$ bp)	7
# contigs ( $\geq 1000$ bp)	7
# contigs ( $\geq 5000$ bp)	6
# contigs ( $\geq 10000$ bp)	2
# contigs ( $\geq 25000$ bp)	0
# contigs ( $\geq 50000$ bp)	0
Total length ( $\geq 0$ bp)	74643
Total length ( $\geq 1000$ bp)	74643
Total length ( $\geq 5000$ bp)	70020
Total length ( $\geq 10000$ bp)	35878
Total length ( $\geq 25000$ bp)	0
Total length ( $\geq 50000$ bp)	0
# contigs	7
Largest contig	24224
Total length	74643
GC (%)	49.03
N50	9948
N90	7300
auN	13933.7
L50	3
L90	6
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

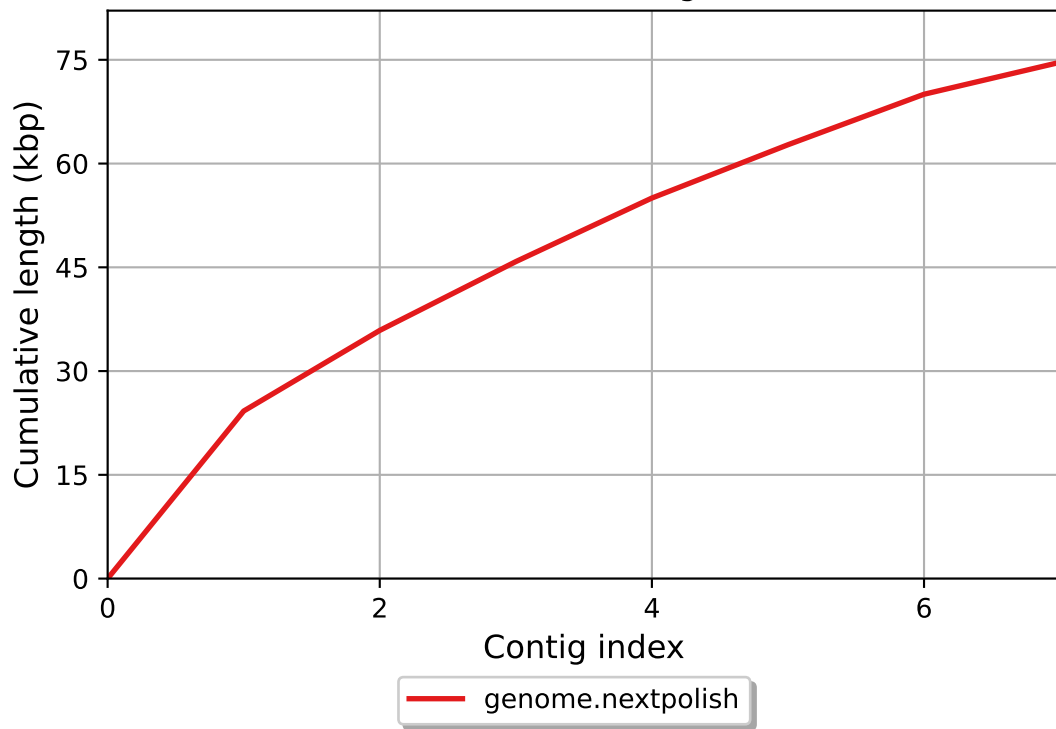
All statistics are based on contigs of size  $\geq 500$  bp, unless otherwise noted (e.g., "# contigs ( $\geq 0$  bp)" and "Total length ( $\geq 0$  bp)" include all contigs).

Nx

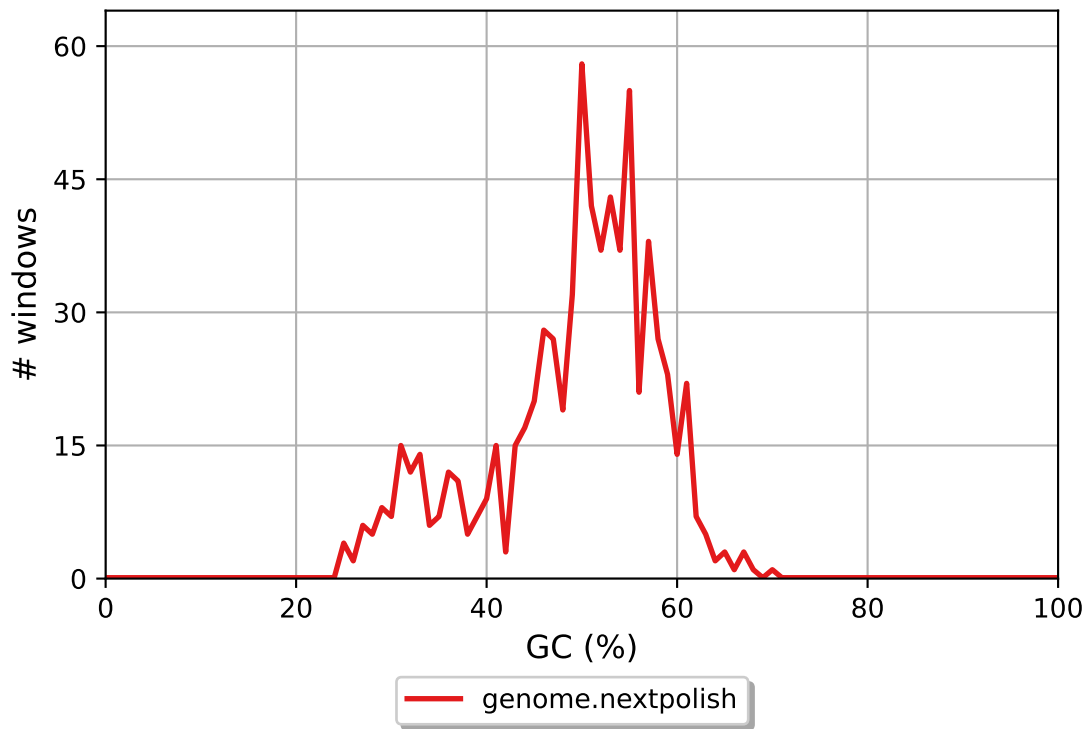


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

