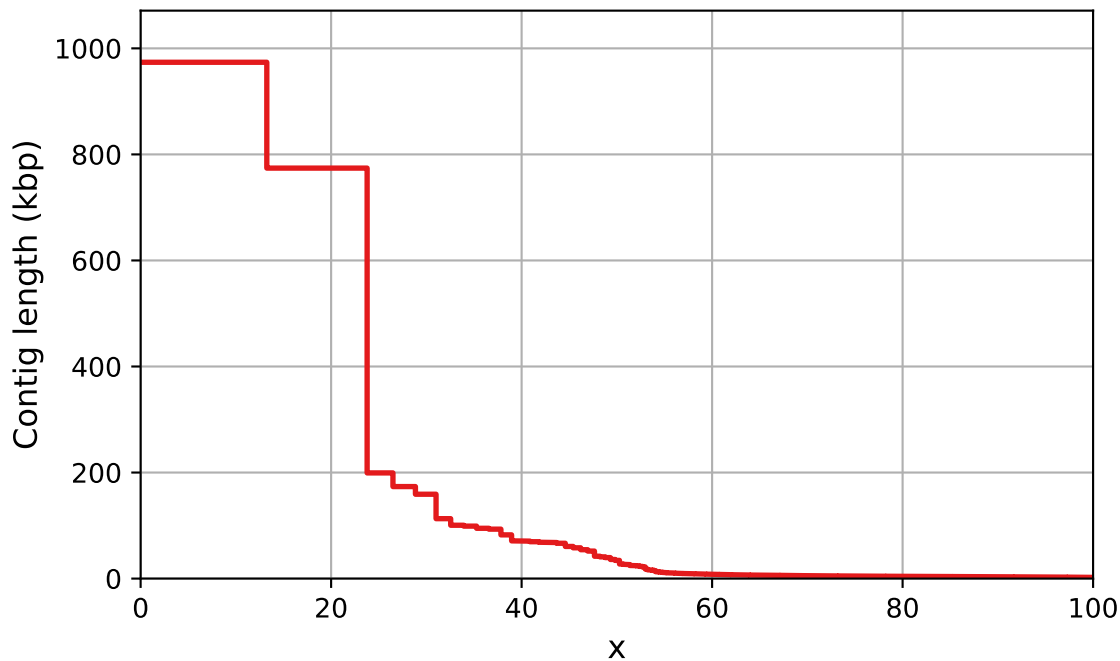


# Report

	genome.nextpolish
# contigs (>= 0 bp)	795
# contigs (>= 1000 bp)	795
# contigs (>= 5000 bp)	238
# contigs (>= 10000 bp)	54
# contigs (>= 25000 bp)	29
# contigs (>= 50000 bp)	21
Total length (>= 0 bp)	7350473
Total length (>= 1000 bp)	7350473
Total length (>= 5000 bp)	5363923
Total length (>= 10000 bp)	4144034
Total length (>= 25000 bp)	3776296
Total length (>= 50000 bp)	3502427
# contigs	795
Largest contig	973775
Total length	7350473
GC (%)	38.46
N50	34309
N90	3430
auN	241168.2
L50	26
L90	542
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	20 + 7 part

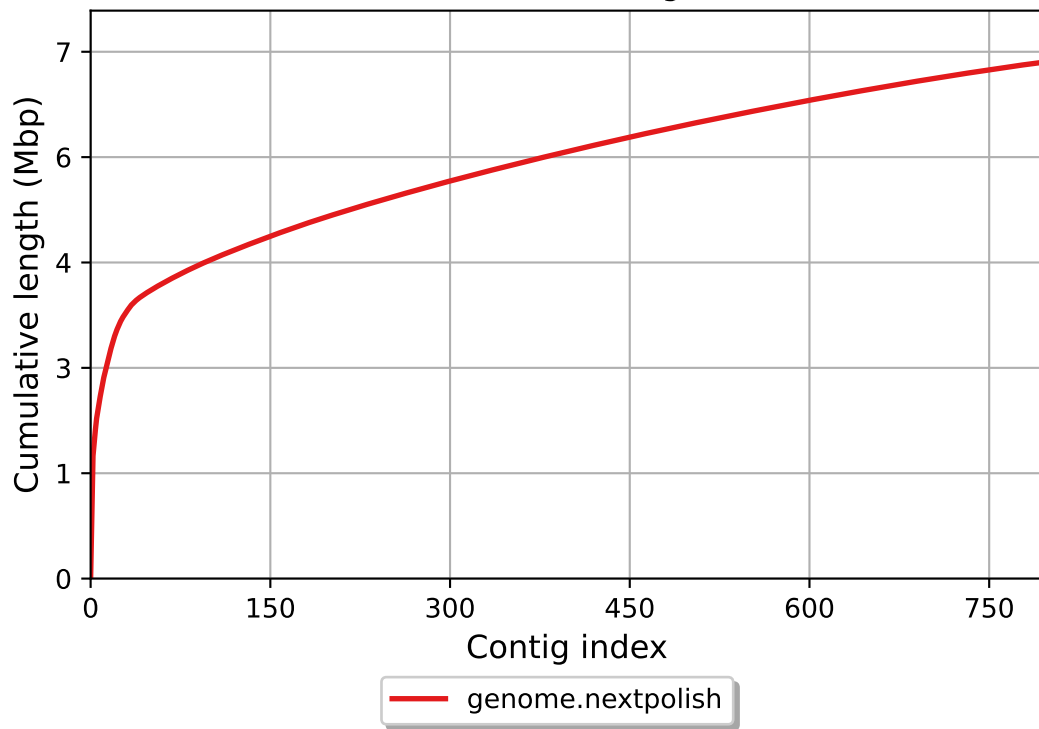
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

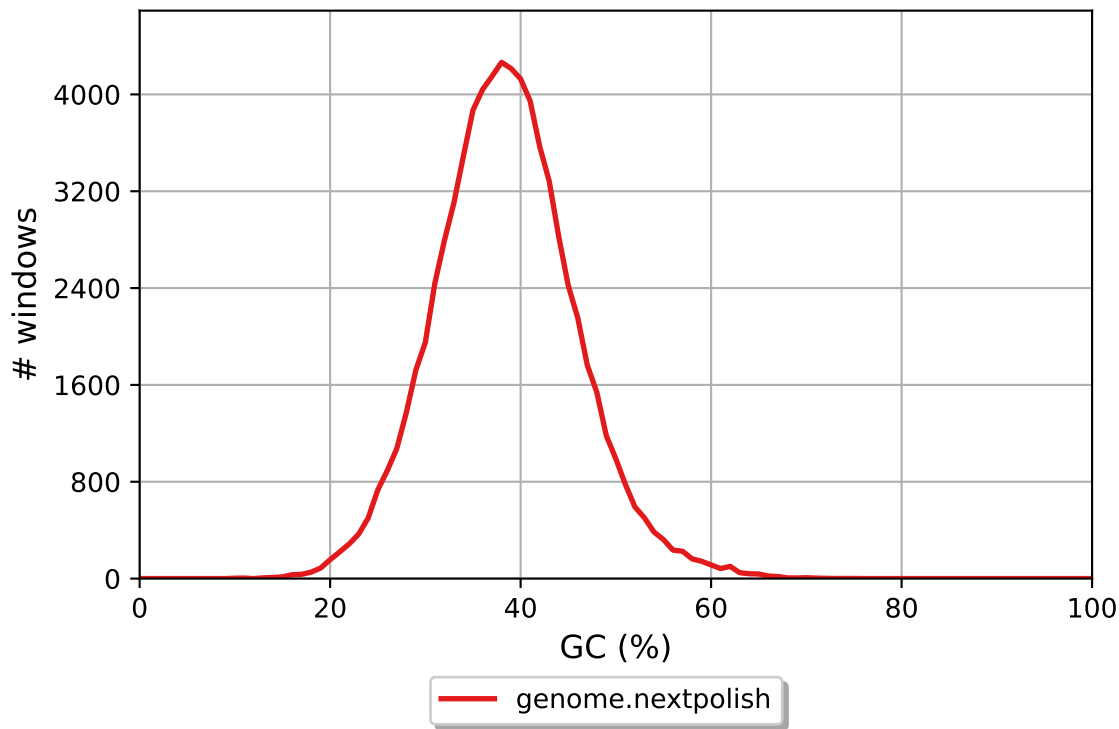


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

