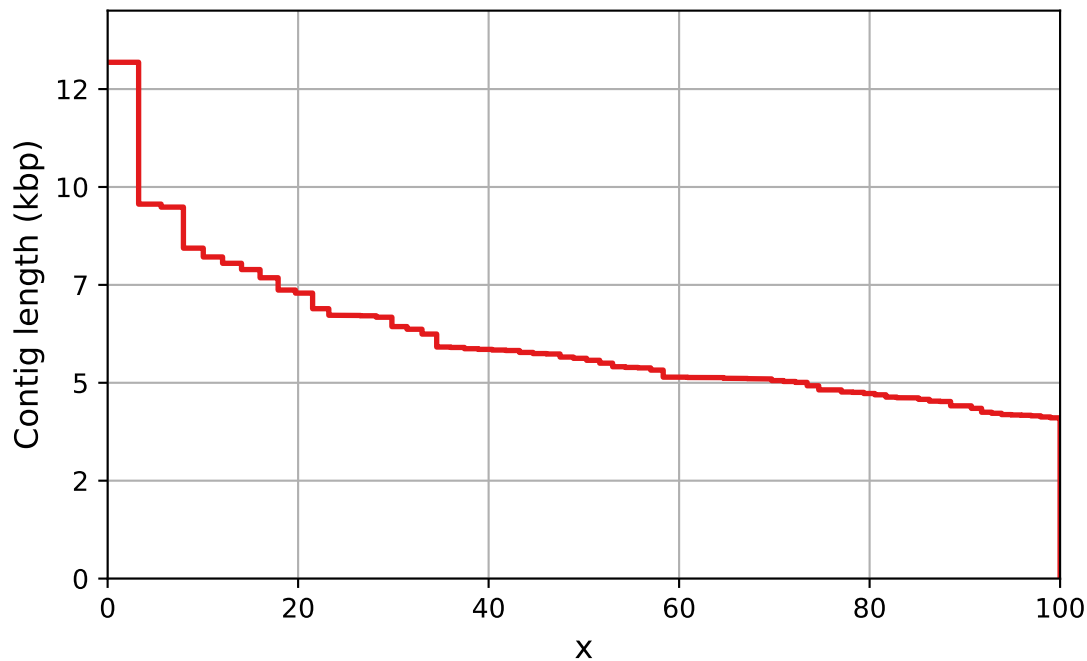


# Report

	genome.nextpolish
# contigs (>= 0 bp)	71
# contigs (>= 1000 bp)	71
# contigs (>= 5000 bp)	47
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	405023
Total length (>= 1000 bp)	405023
Total length (>= 5000 bp)	297463
Total length (>= 10000 bp)	13184
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	71
Largest contig	13184
Total length	405023
GC (%)	50.93
N50	5625
N90	4411
auN	6118.3
L50	29
L90	62
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 1 part

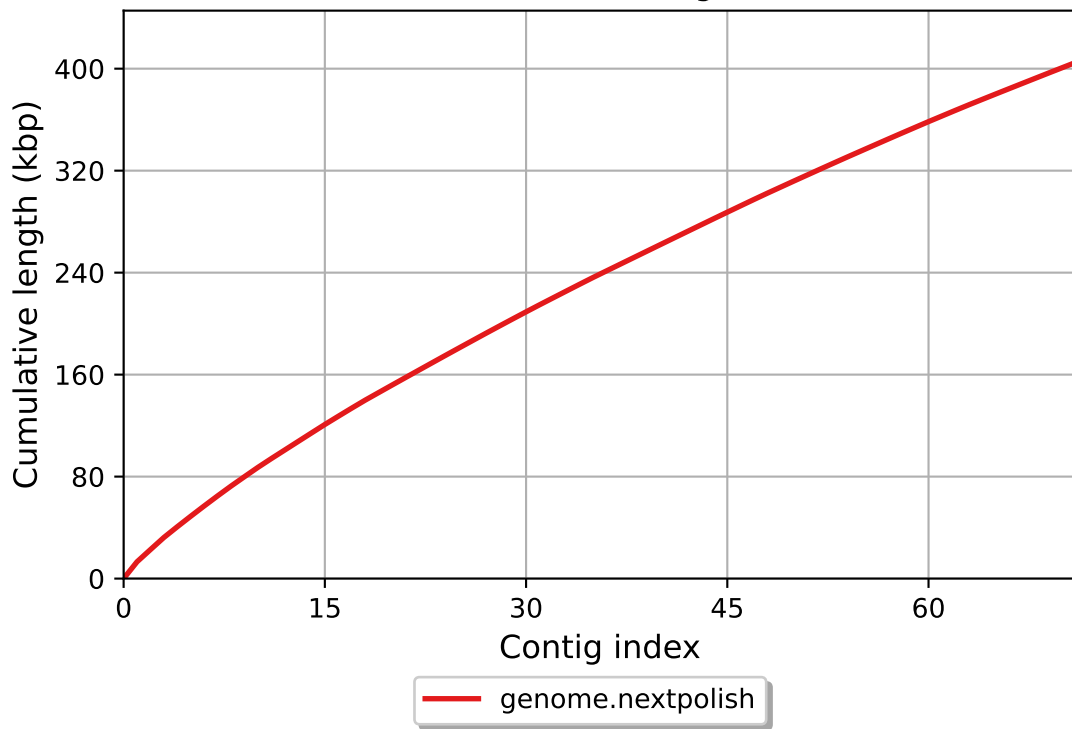
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

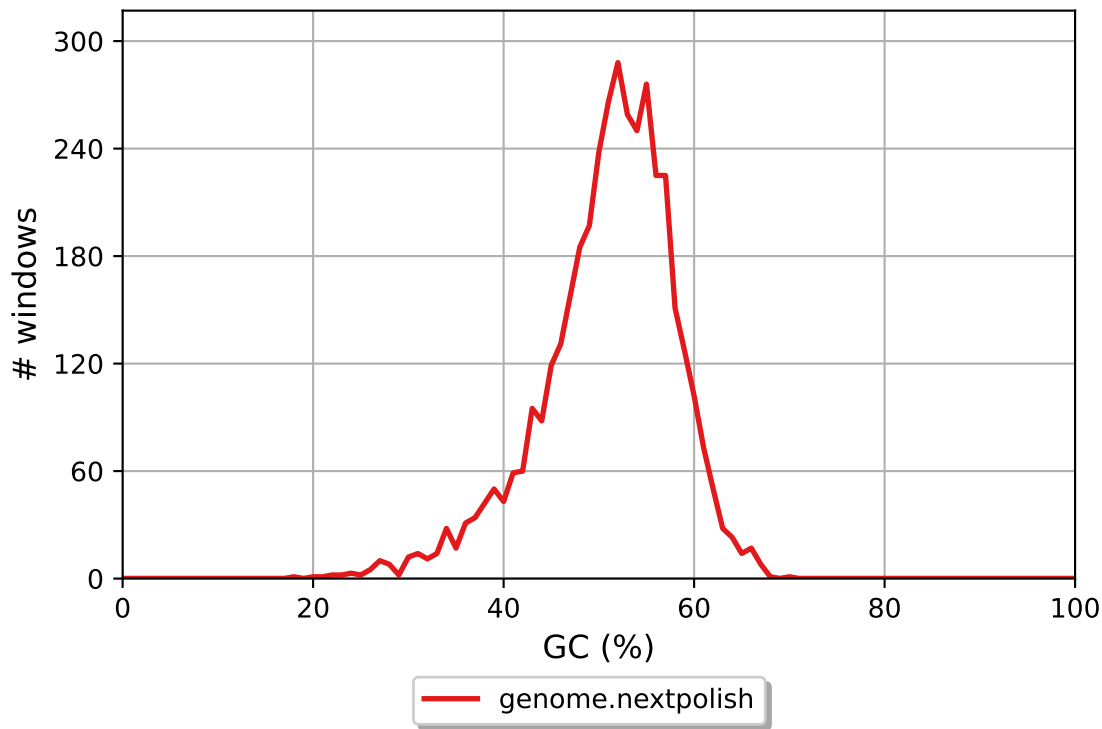


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

