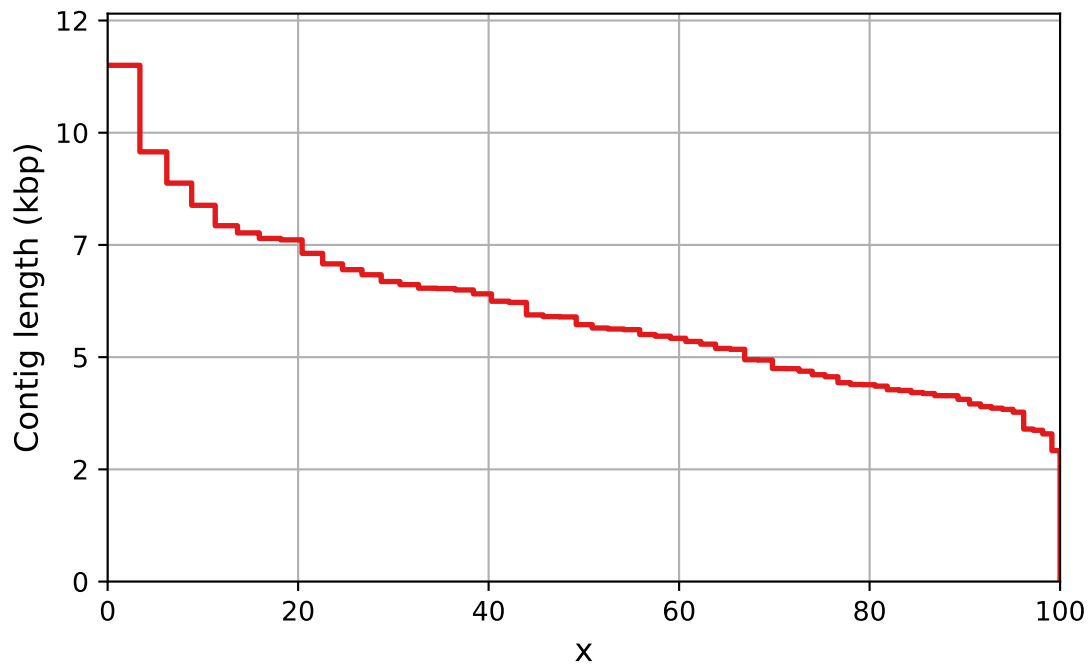


# Report

	genome.nextpolish
# contigs (>= 0 bp)	61
# contigs (>= 1000 bp)	61
# contigs (>= 5000 bp)	34
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	339324
Total length (>= 1000 bp)	339324
Total length (>= 5000 bp)	226955
Total length (>= 10000 bp)	11502
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	61
Largest contig	11502
Total length	339324
GC (%)	58.07
N50	5724
N90	4059
auN	6050.5
L50	24
L90	52
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 0 part

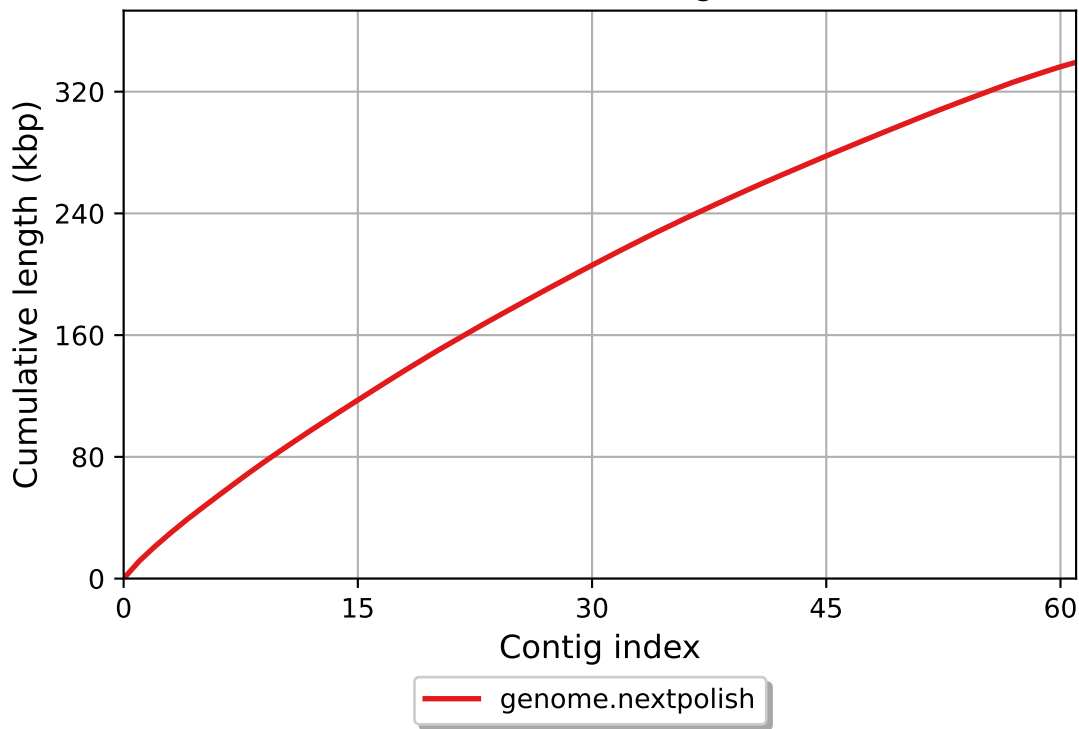
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

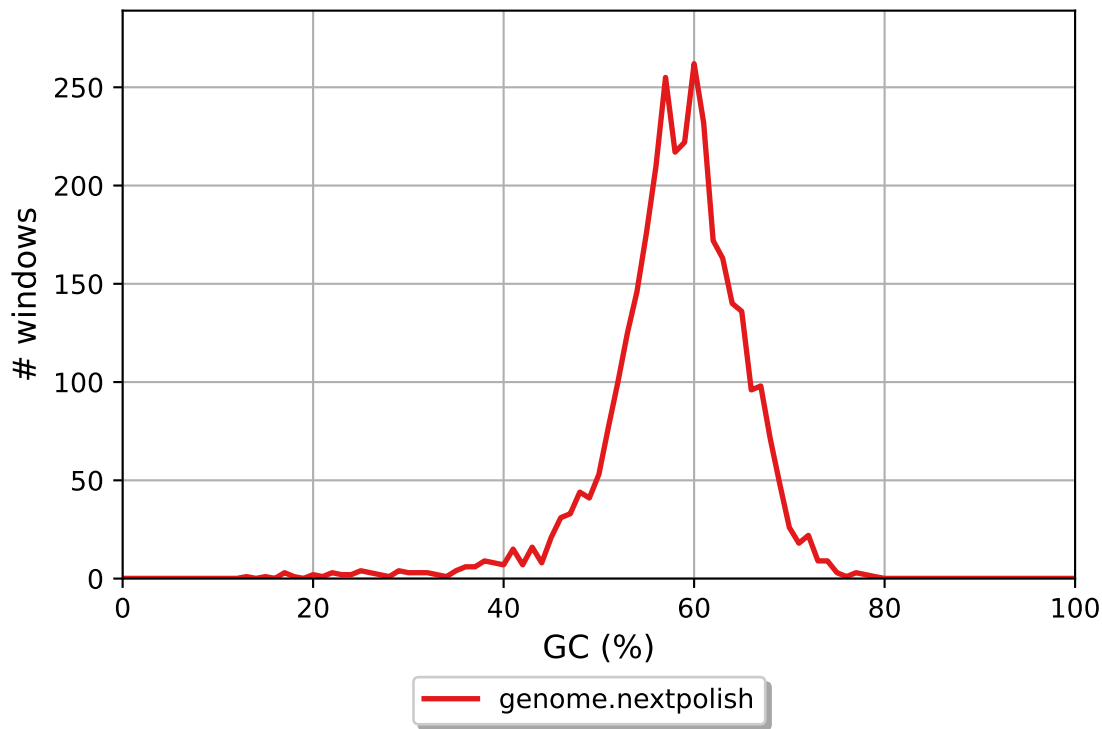


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

