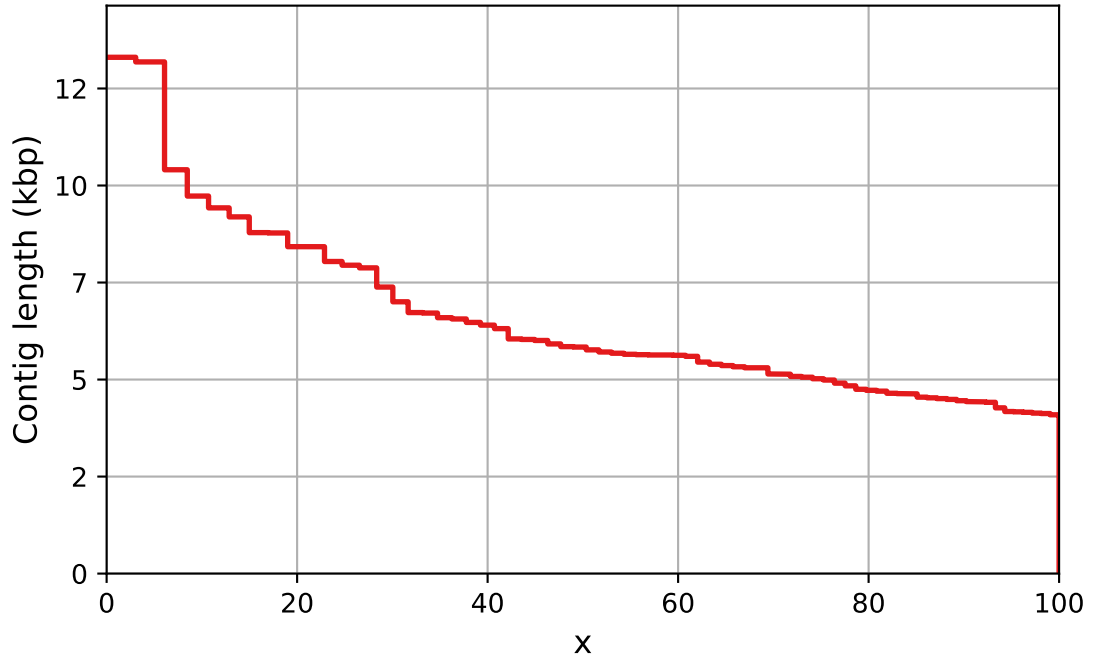


# Report

	genome.nextpolish
# contigs (>= 0 bp)	72
# contigs (>= 1000 bp)	72
# contigs (>= 5000 bp)	48
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	435626
Total length (>= 1000 bp)	435626
Total length (>= 5000 bp)	327888
Total length (>= 10000 bp)	36895
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	72
Largest contig	13304
Total length	435626
GC (%)	50.72
N50	5837
N90	4455
auN	6665.5
L50	28
L90	62
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 1 part

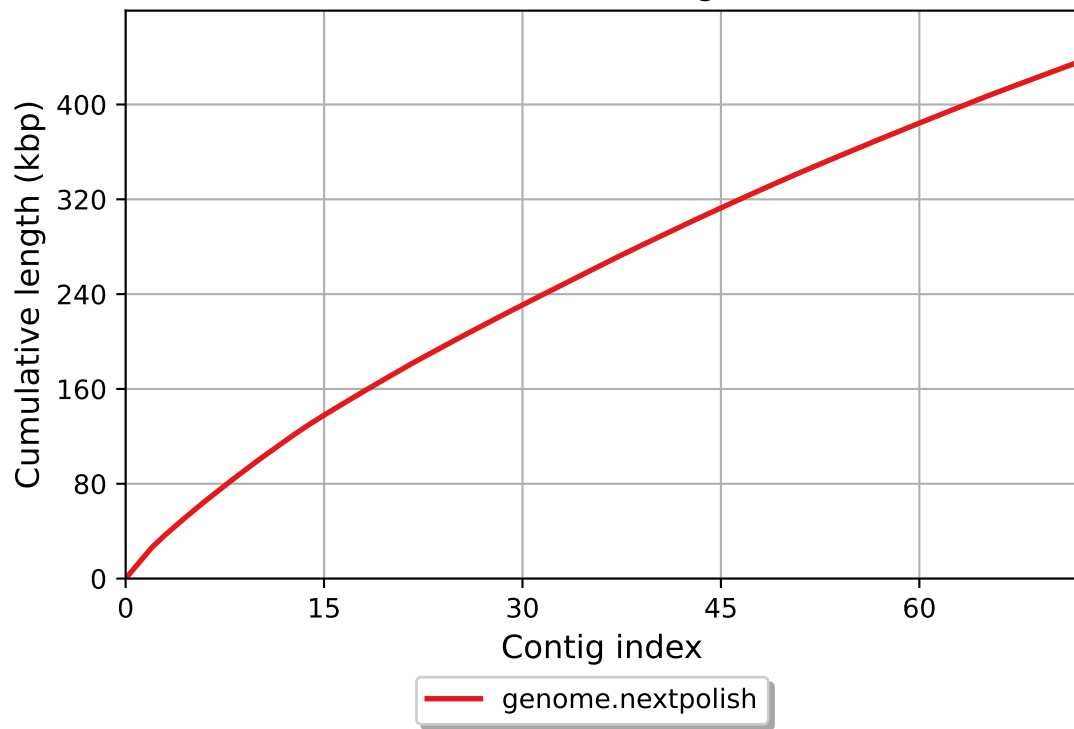
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

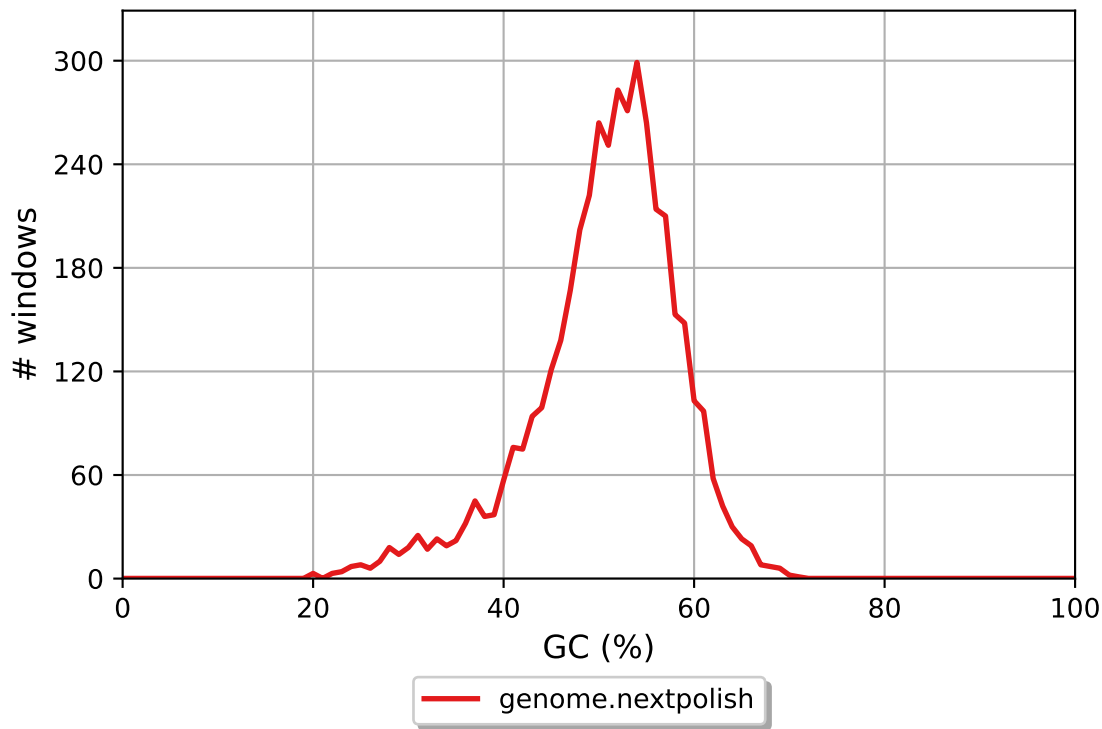


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

