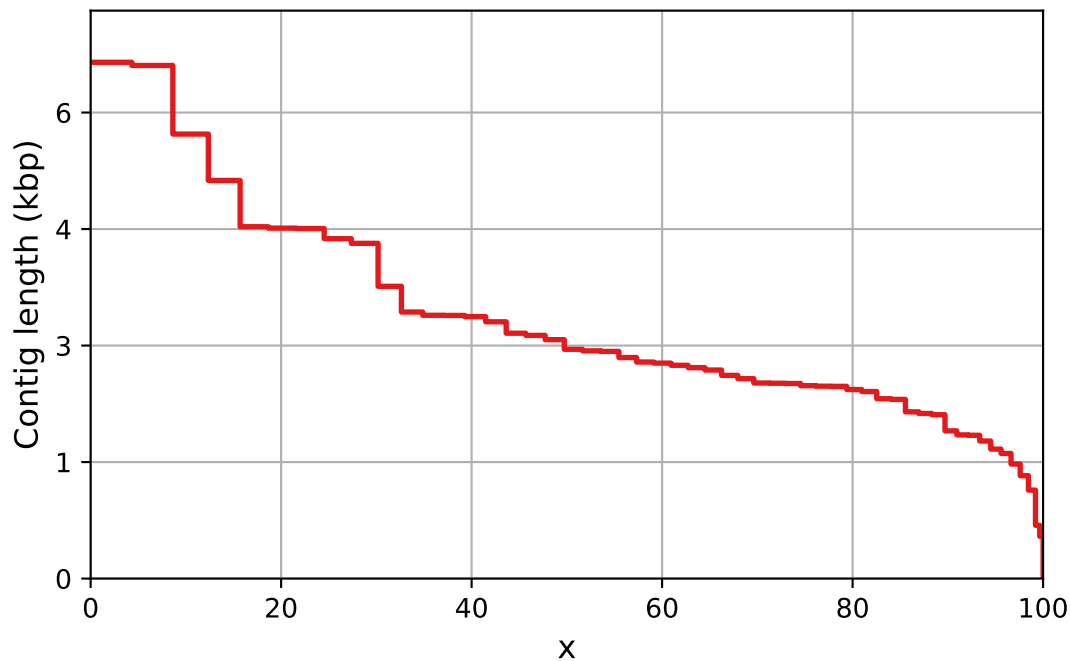


# Report

	genome.nextpolish
# contigs (>= 0 bp)	54
# contigs (>= 1000 bp)	51
# contigs (>= 5000 bp)	4
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	153797
Total length (>= 1000 bp)	152329
Total length (>= 5000 bp)	24102
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	53
Largest contig	6647
Total length	153559
GC (%)	40.40
N50	2952
N90	1902
auN	3449.2
L50	19
L90	43
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

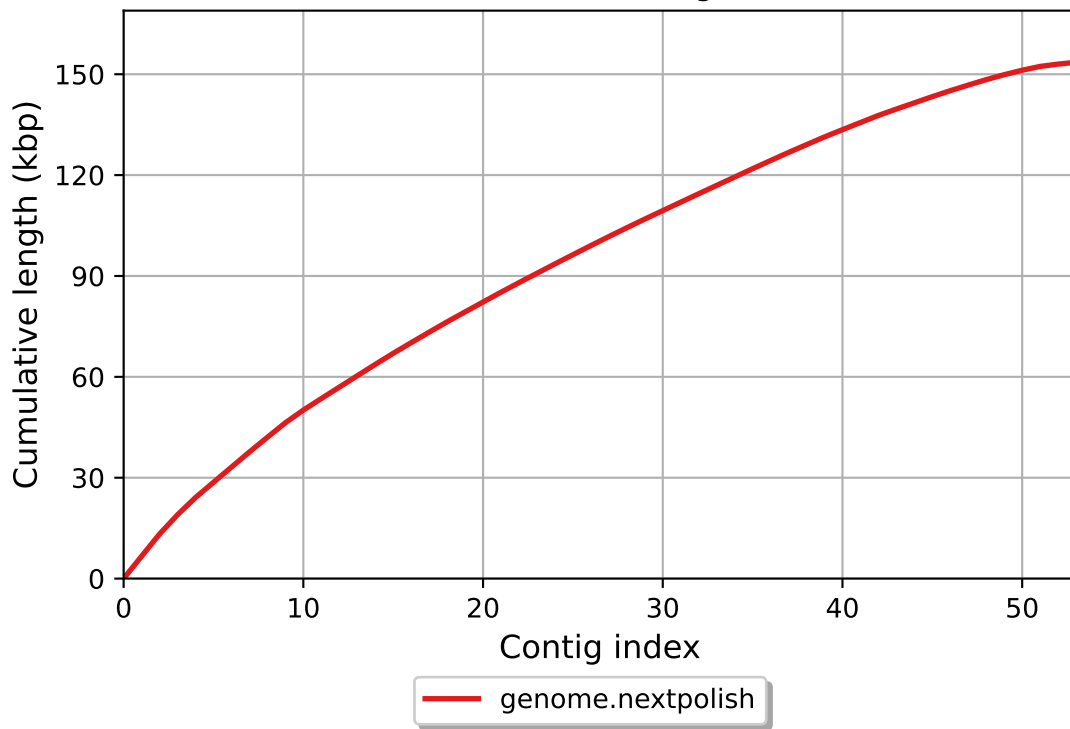
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

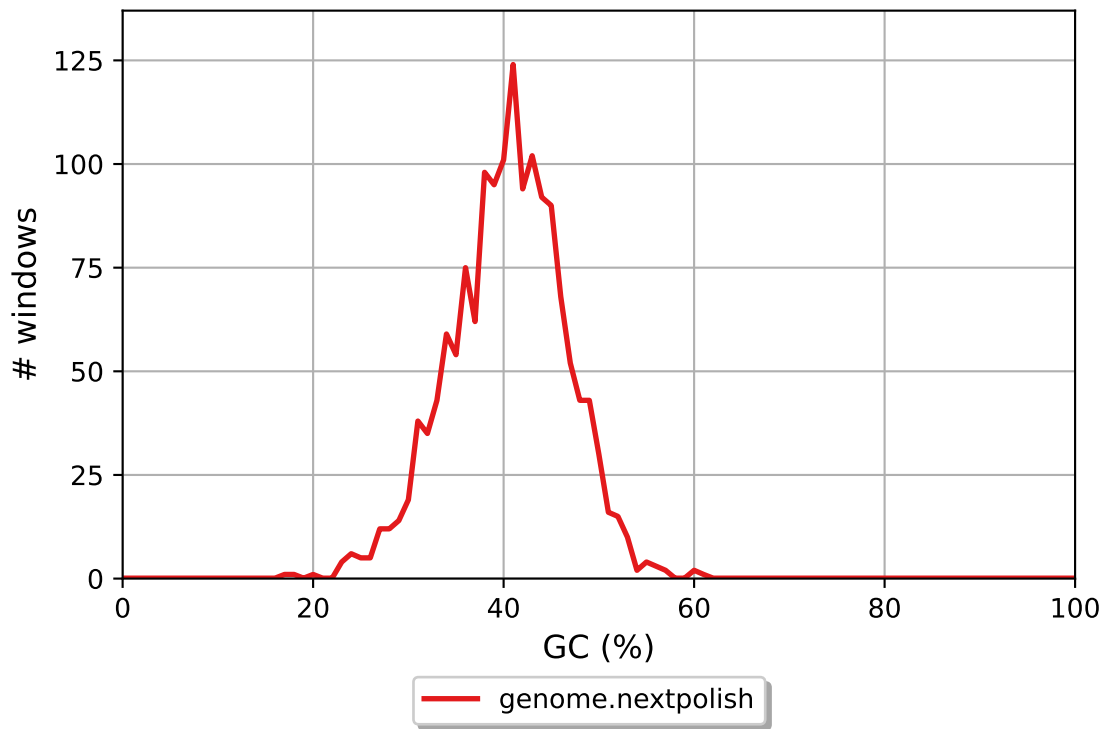


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

