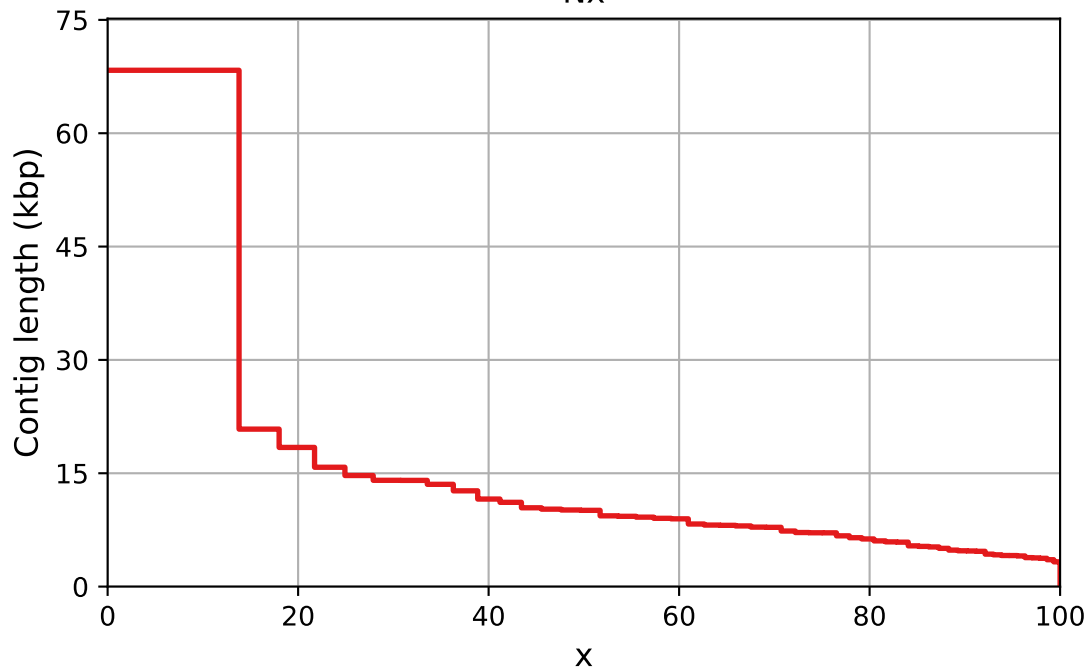


# Report

	genome.nextpolish
# contigs (>= 0 bp)	54
# contigs (>= 1000 bp)	54
# contigs (>= 5000 bp)	40
# contigs (>= 10000 bp)	15
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	495211
Total length (>= 1000 bp)	495211
Total length (>= 5000 bp)	437283
Total length (>= 10000 bp)	256051
Total length (>= 25000 bp)	68331
Total length (>= 50000 bp)	68331
# contigs	54
Largest contig	68331
Total length	495211
GC (%)	40.24
N50	10092
N90	4738
auN	18024.9
L50	15
L90	42
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 0 part

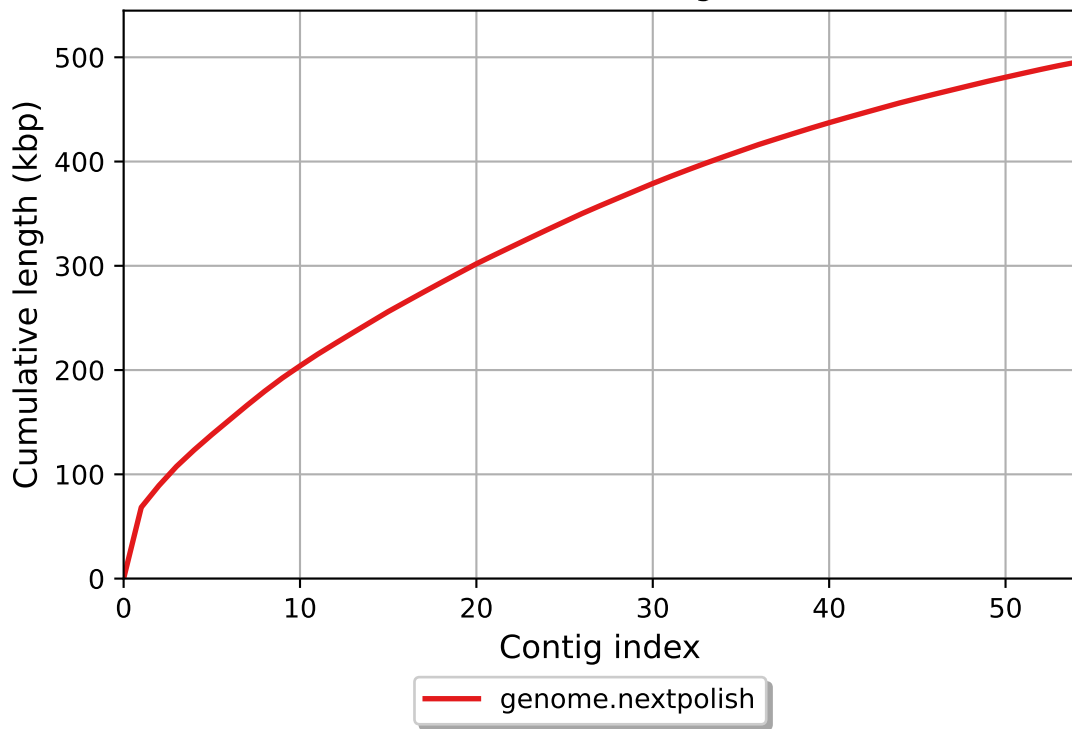
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

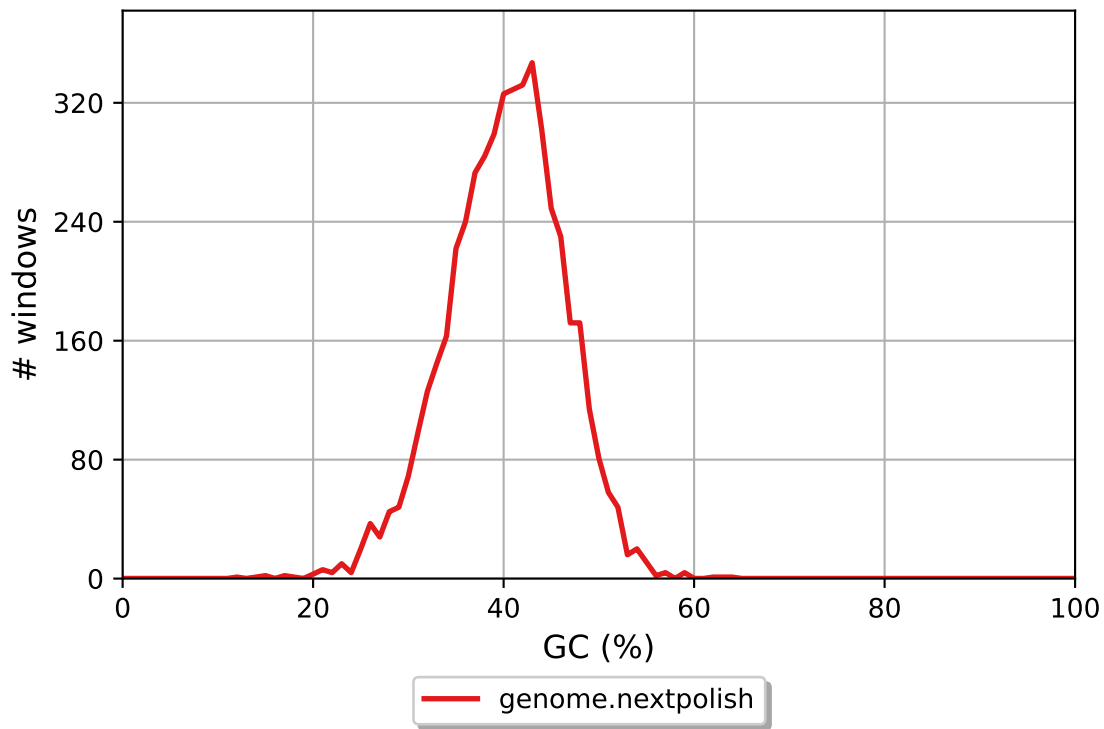


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

