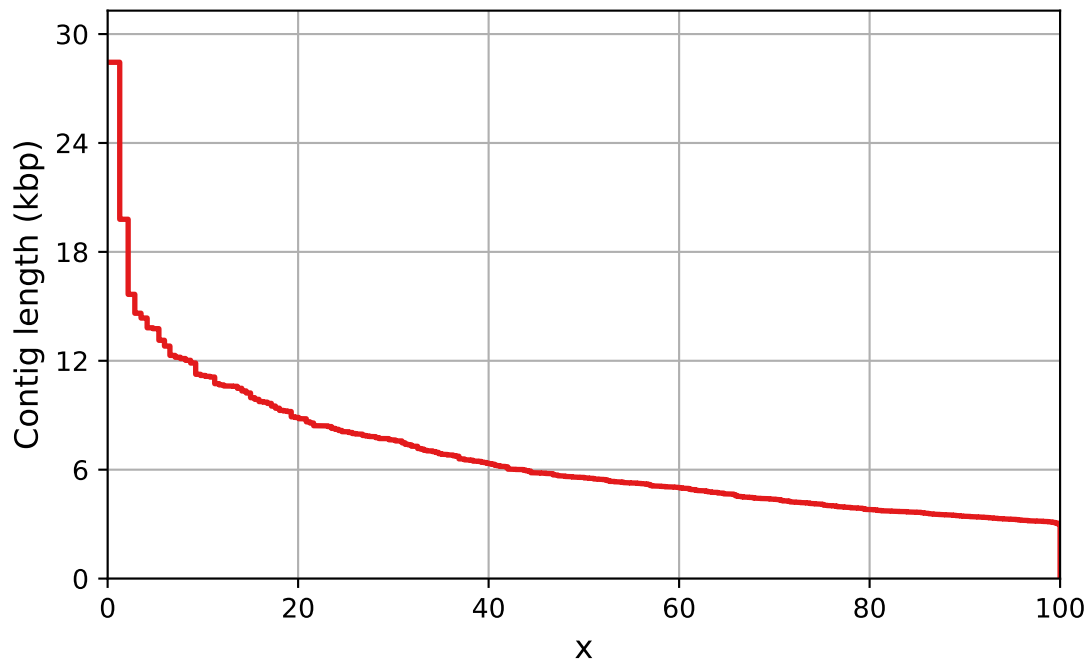


# Report

	genome.nextpolish
# contigs (>= 0 bp)	412
# contigs (>= 1000 bp)	412
# contigs (>= 5000 bp)	179
# contigs (>= 10000 bp)	26
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2238142
Total length (>= 1000 bp)	2238142
Total length (>= 5000 bp)	1346083
Total length (>= 10000 bp)	335874
Total length (>= 25000 bp)	28446
Total length (>= 50000 bp)	0
# contigs	412
Largest contig	28446
Total length	2238142
GC (%)	35.27
N50	5574
N90	3432
auN	6758.7
L50	136
L90	344
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 0 part

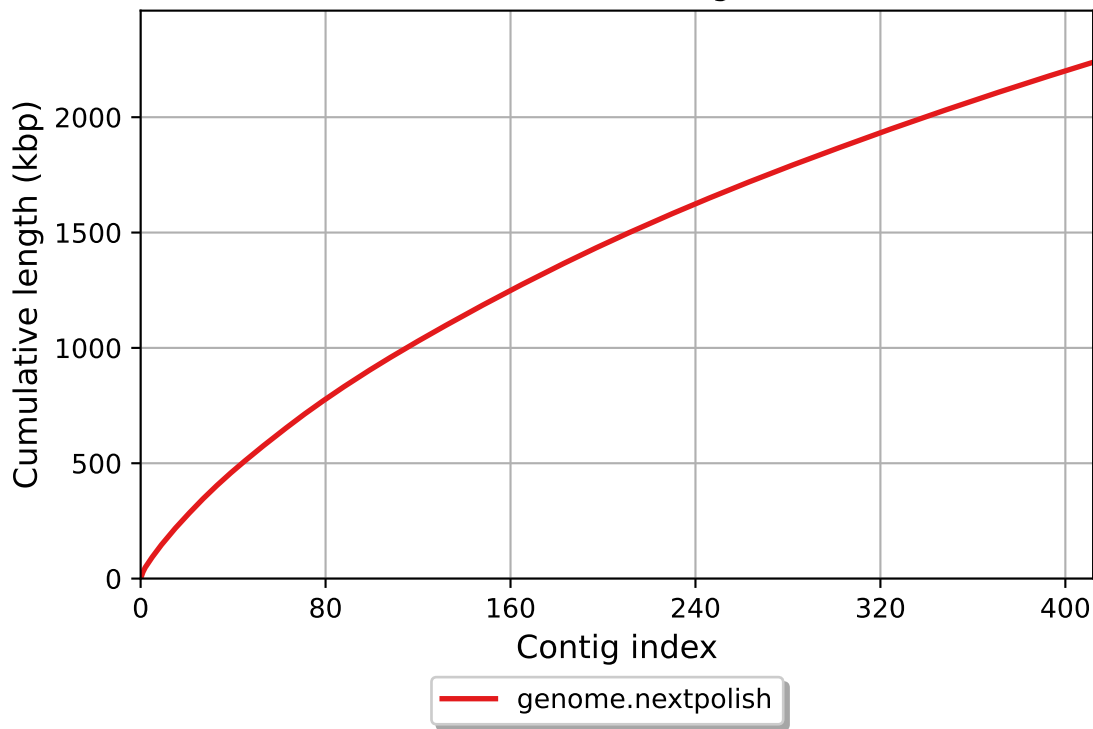
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

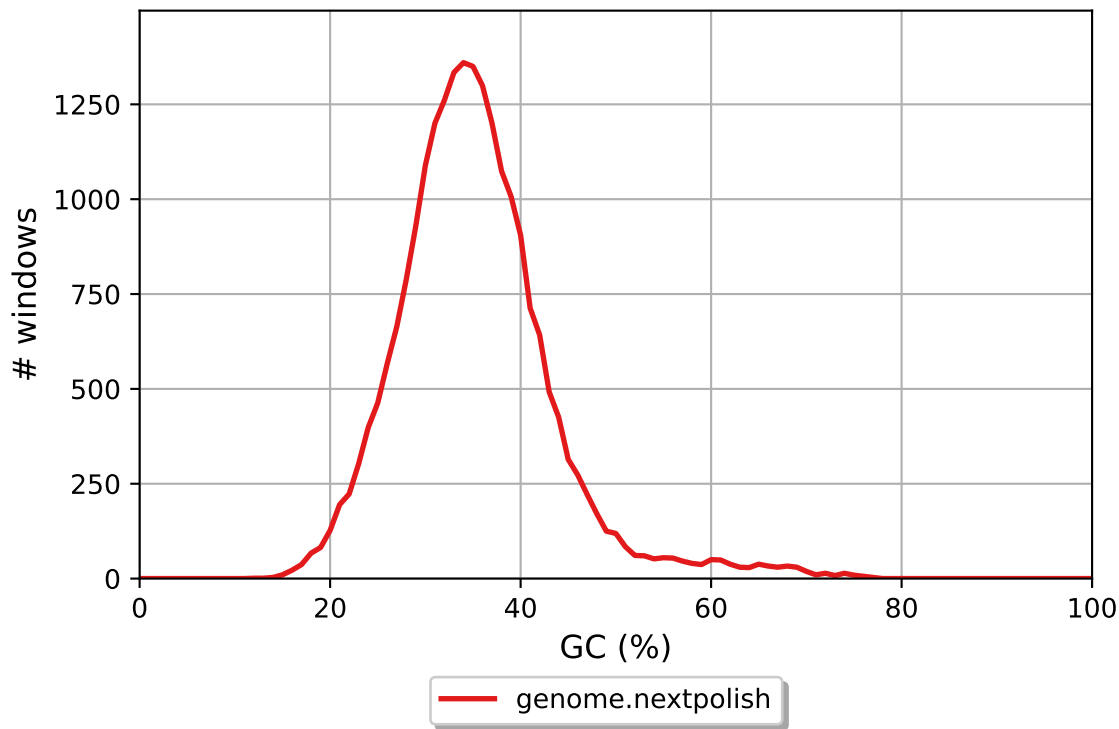


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

