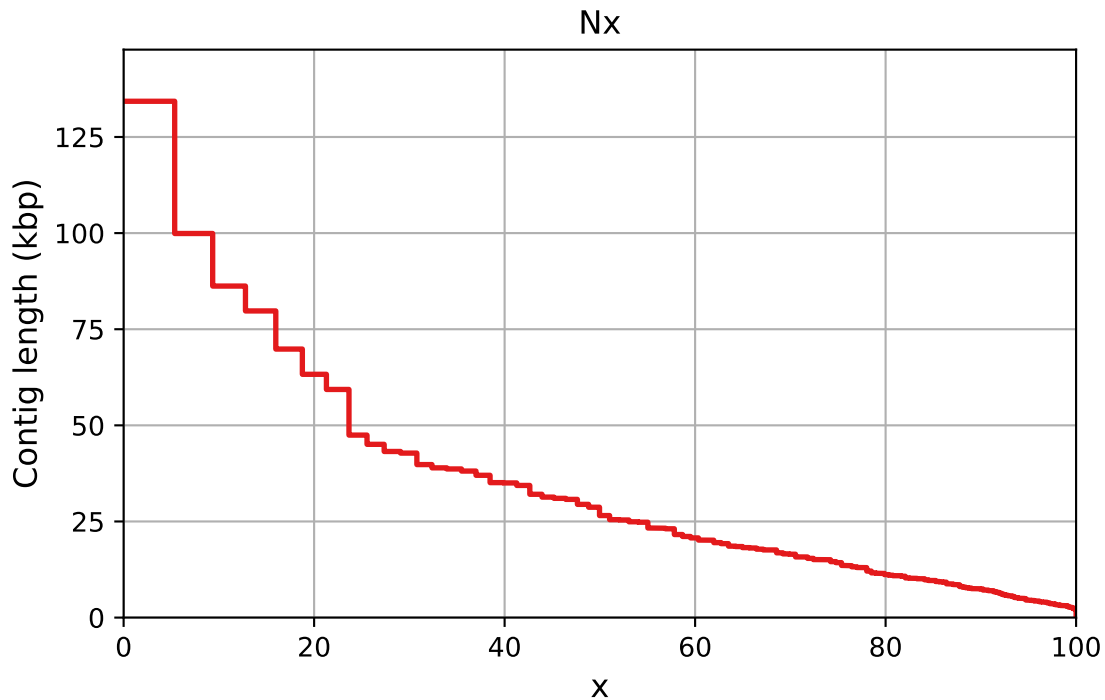


# Report

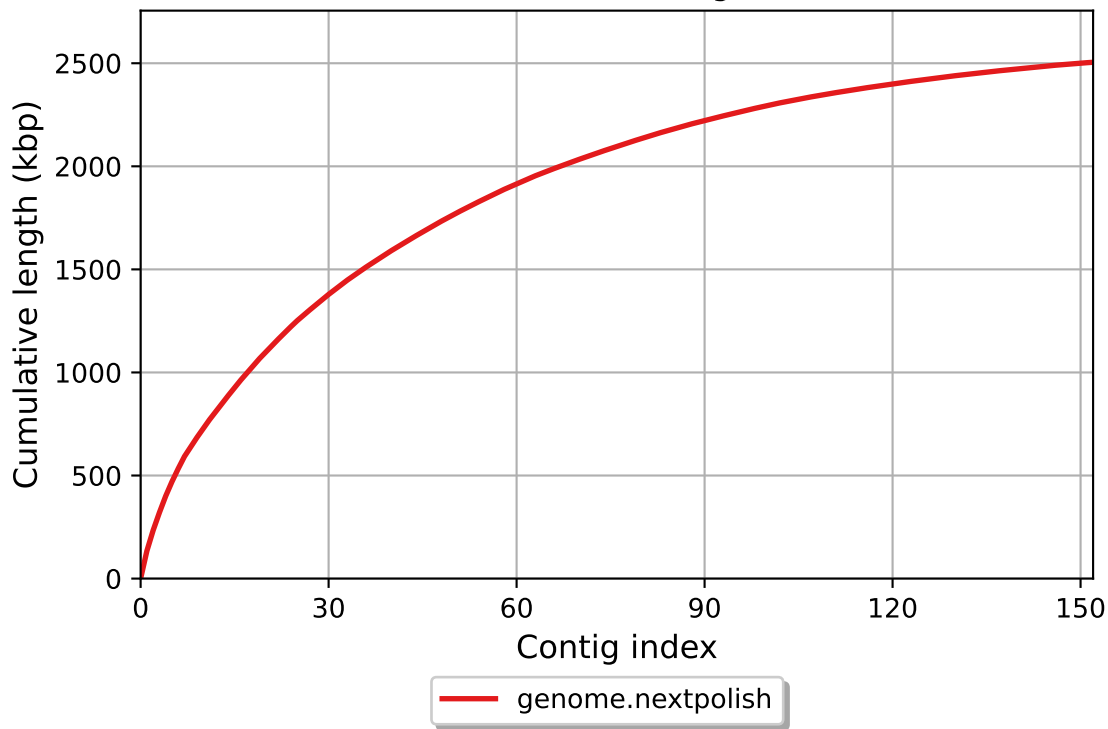
	genome.nextpolish
# contigs (>= 0 bp)	152
# contigs (>= 1000 bp)	152
# contigs (>= 5000 bp)	112
# contigs (>= 10000 bp)	77
# contigs (>= 25000 bp)	28
# contigs (>= 50000 bp)	7
Total length (>= 0 bp)	2504715
Total length (>= 1000 bp)	2504715
Total length (>= 5000 bp)	2362688
Total length (>= 10000 bp)	2106573
Total length (>= 25000 bp)	1328846
Total length (>= 50000 bp)	592617
# contigs	152
Largest contig	134289
Total length	2504715
GC (%)	32.23
N50	26536
N90	7450
auN	38555.7
L50	26
L90	95
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

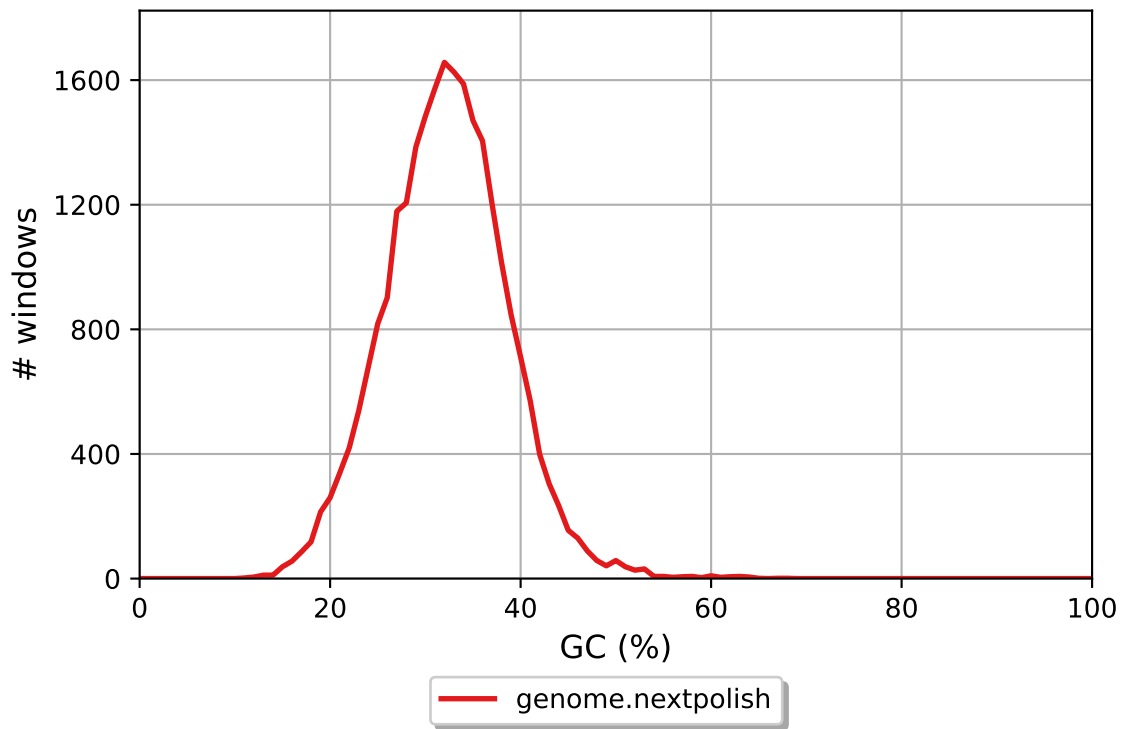


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

