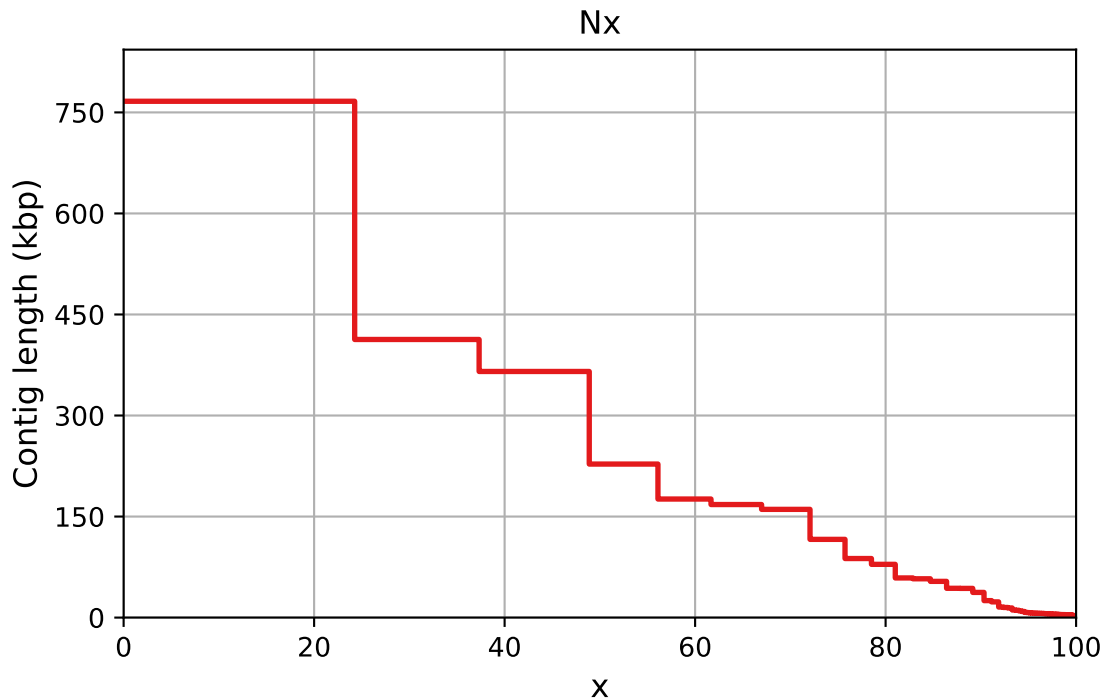


# Report

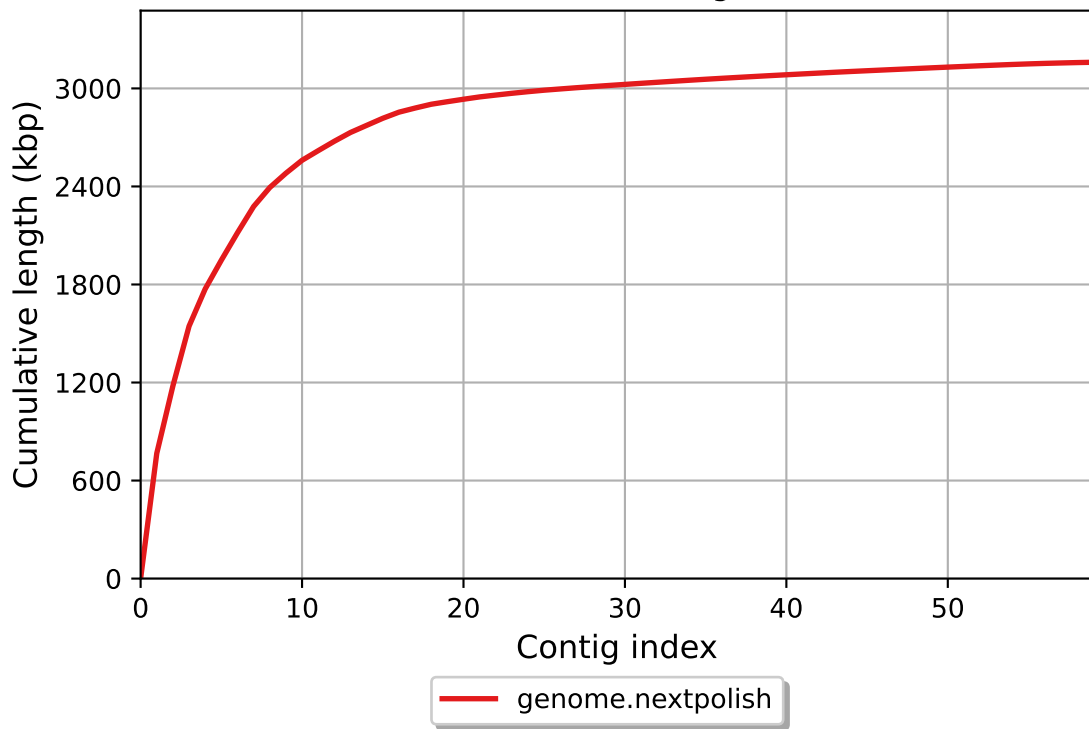
	genome.nextpolish
# contigs (>= 0 bp)	59
# contigs (>= 1000 bp)	59
# contigs (>= 5000 bp)	43
# contigs (>= 10000 bp)	23
# contigs (>= 25000 bp)	17
# contigs (>= 50000 bp)	13
Total length (>= 0 bp)	3160032
Total length (>= 1000 bp)	3160032
Total length (>= 5000 bp)	3098873
Total length (>= 10000 bp)	2970181
Total length (>= 25000 bp)	2879753
Total length (>= 50000 bp)	2730347
# contigs	59
Largest contig	766539
Total length	3160032
GC (%)	33.16
N50	227920
N90	37374
auN	339855.1
L50	4
L90	16
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 5 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).

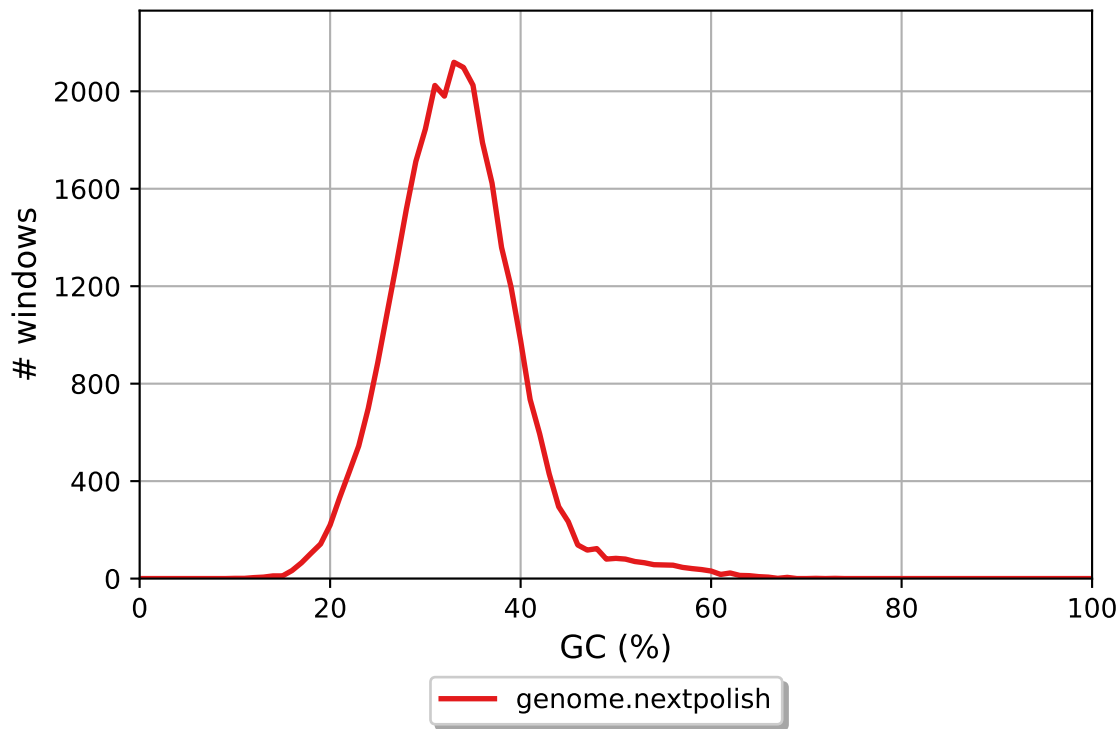


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

