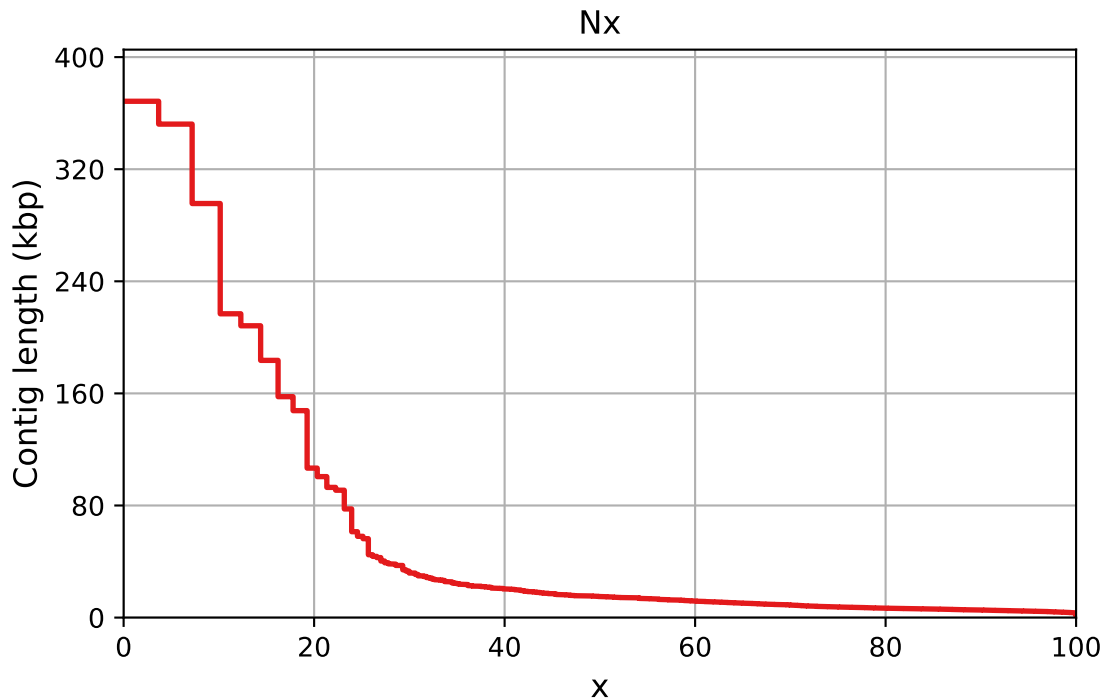


# Report

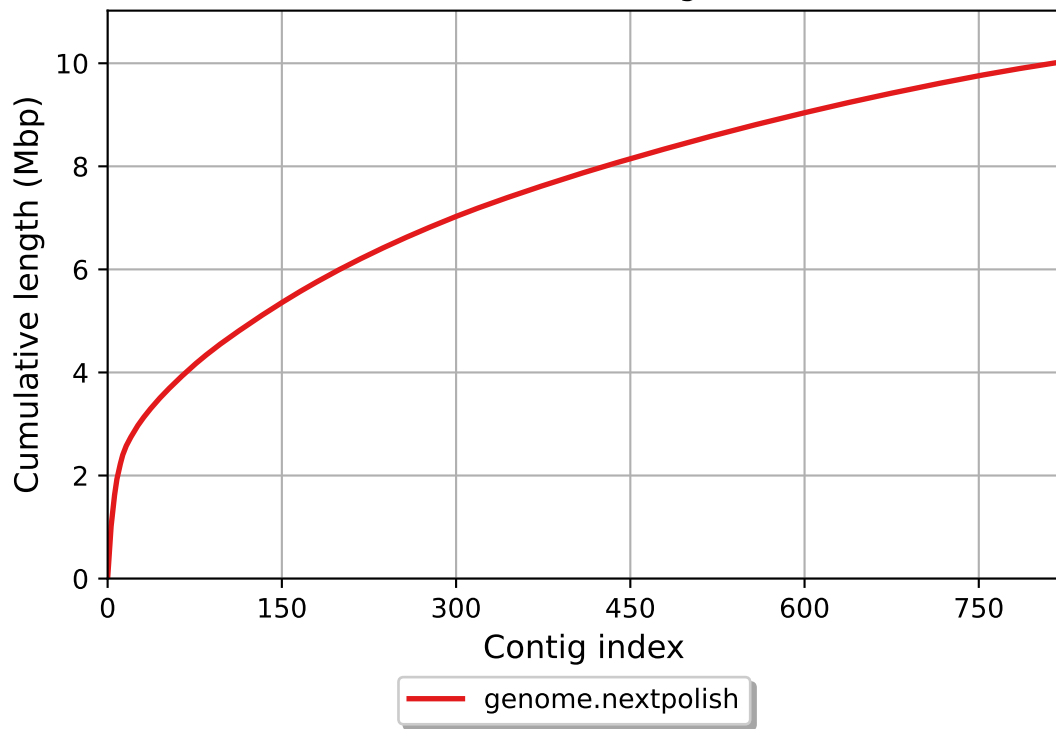
	genome.nextpolish
# contigs (>= 0 bp)	820
# contigs (>= 1000 bp)	820
# contigs (>= 5000 bp)	648
# contigs (>= 10000 bp)	259
# contigs (>= 25000 bp)	43
# contigs (>= 50000 bp)	16
Total length (>= 0 bp)	10019328
Total length (>= 1000 bp)	10019328
Total length (>= 5000 bp)	9288822
Total length (>= 10000 bp)	6641311
Total length (>= 25000 bp)	3454923
Total length (>= 50000 bp)	2574413
# contigs	820
Largest contig	368481
Total length	10019328
GC (%)	53.56
N50	15038
N90	5399
auN	67287.2
L50	127
L90	596
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	20 + 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).

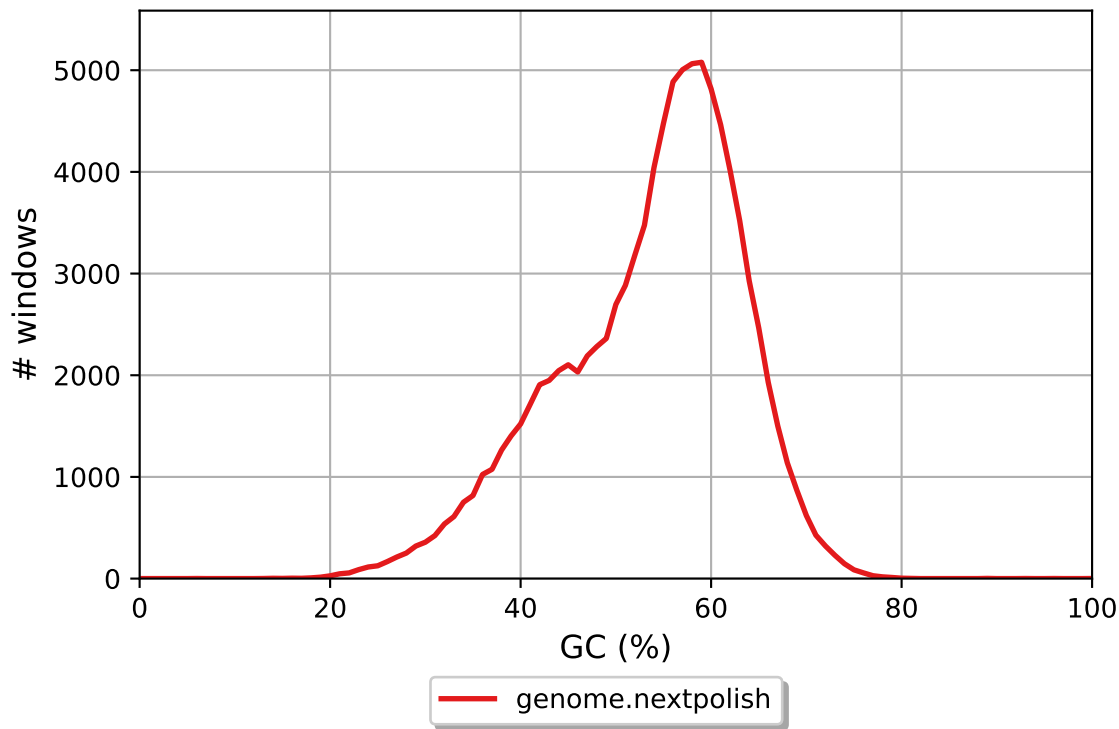


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

