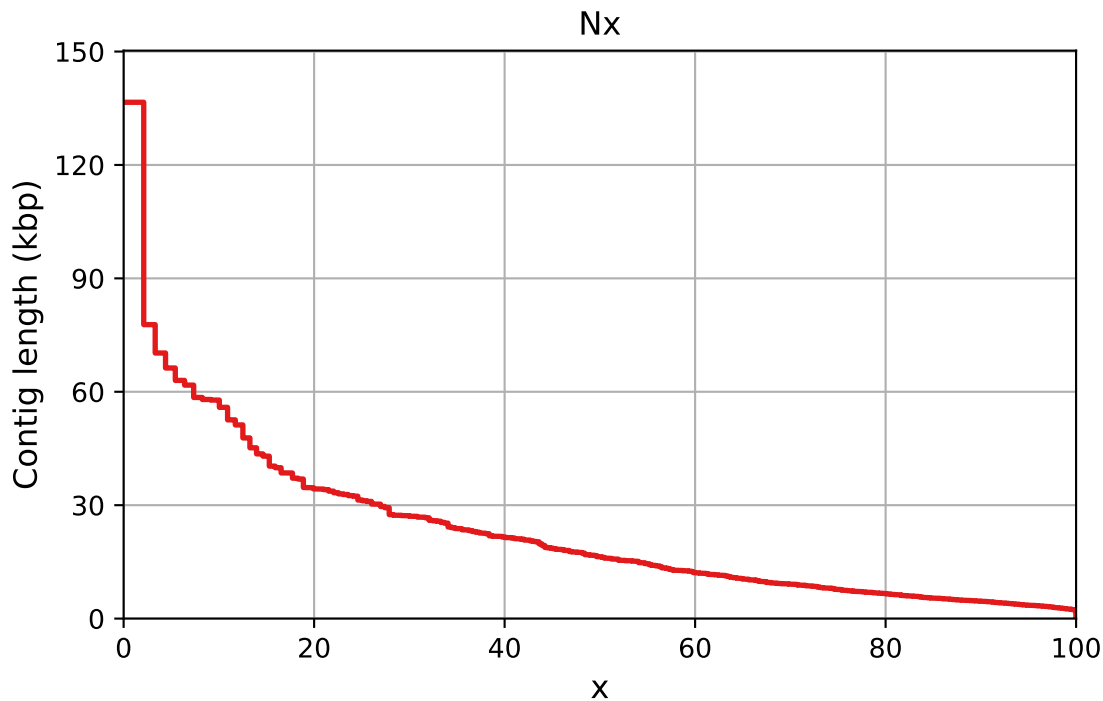


# Report

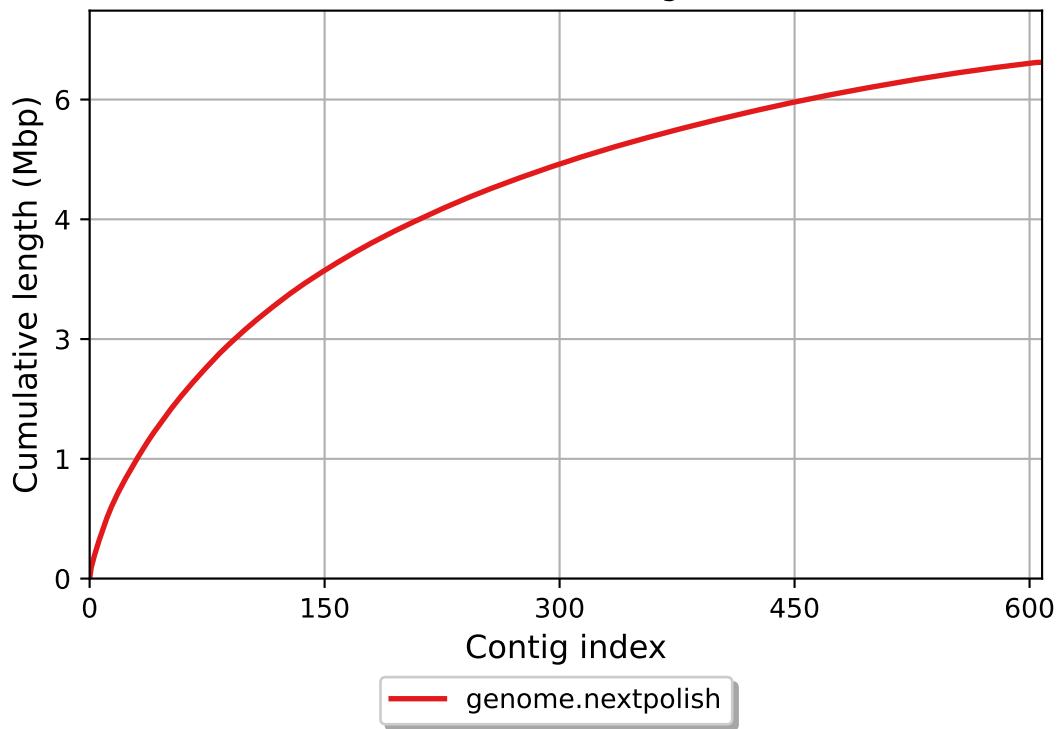
	genome.nextpolish
# contigs (>= 0 bp)	608
# contigs (>= 1000 bp)	605
# contigs (>= 5000 bp)	380
# contigs (>= 10000 bp)	190
# contigs (>= 25000 bp)	55
# contigs (>= 50000 bp)	12
Total length (>= 0 bp)	6466921
Total length (>= 1000 bp)	6464949
Total length (>= 5000 bp)	5647874
Total length (>= 10000 bp)	4306667
Total length (>= 25000 bp)	2202564
Total length (>= 50000 bp)	809590
# contigs	608
Largest contig	136583
Total length	6466921
GC (%)	35.81
N50	16339
N90	4597
auN	23978.8
L50	107
L90	416
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 2 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).

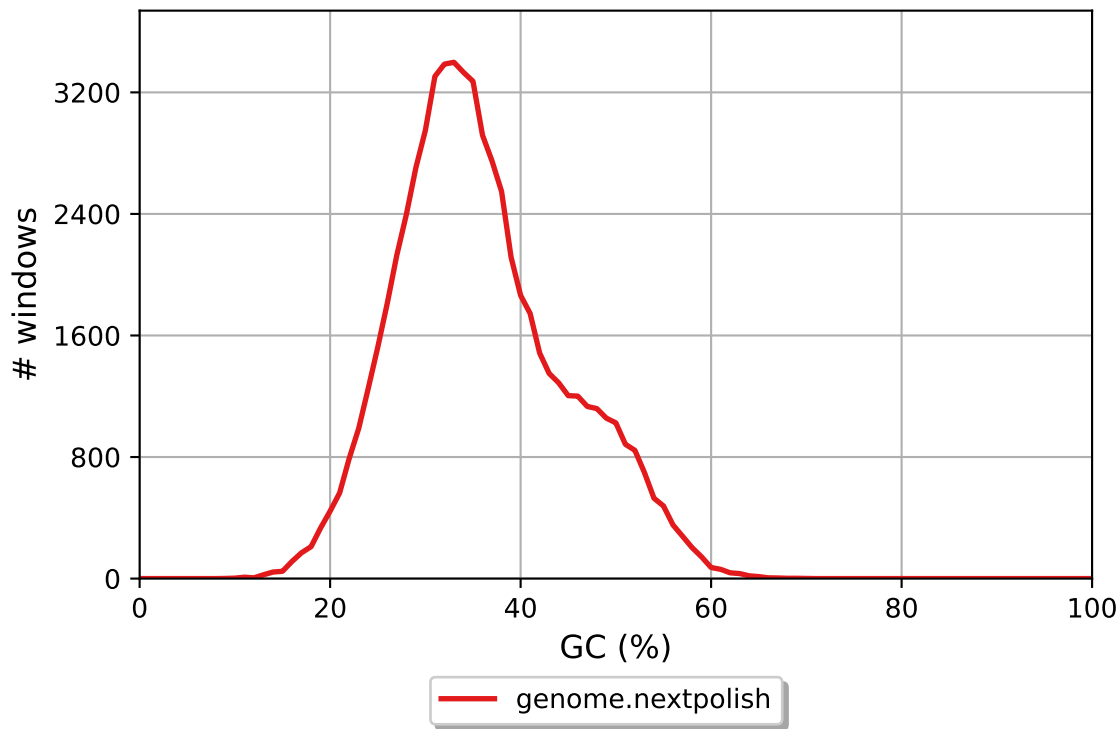


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

