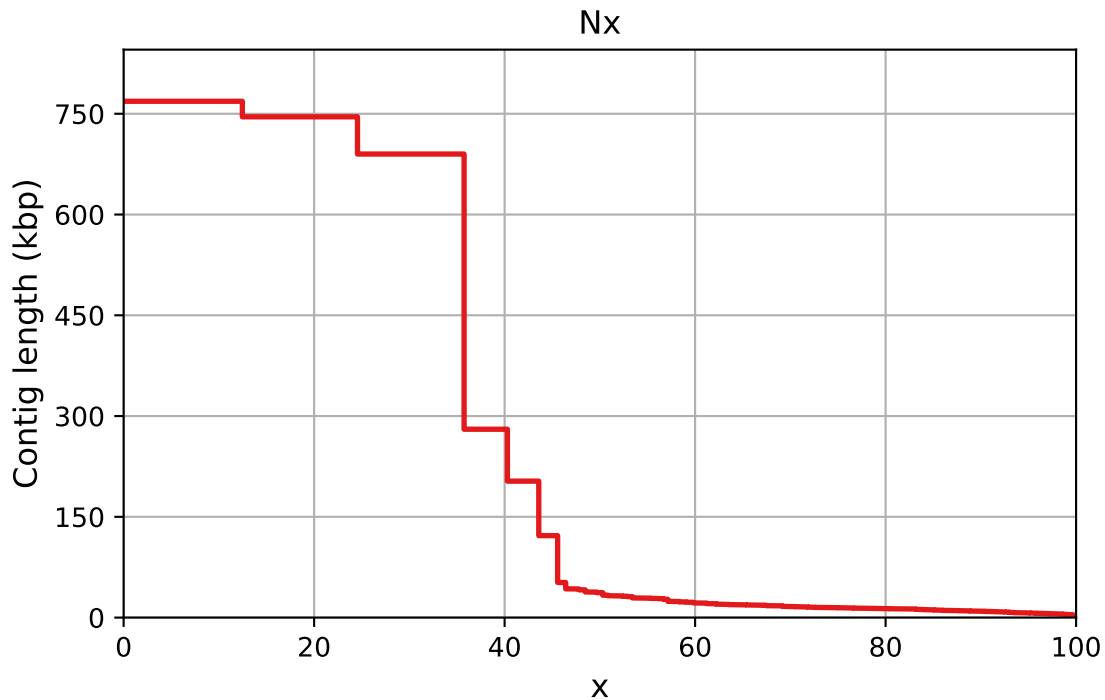


Report

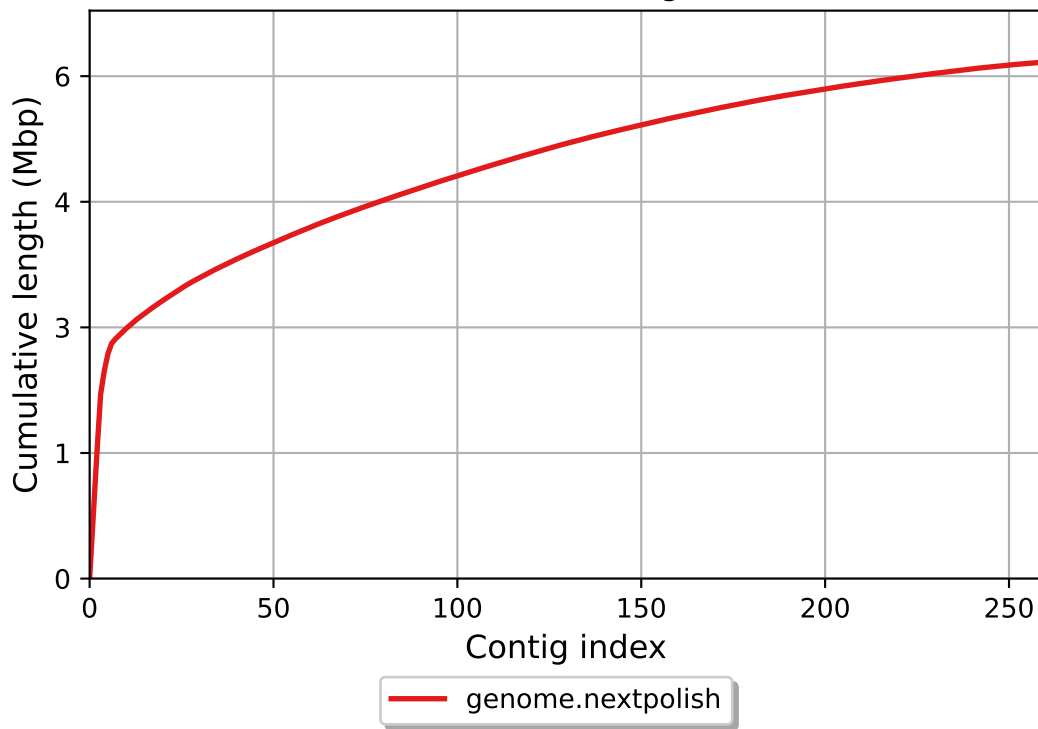
	genome.nextpolish
# contigs (>= 0 bp)	259
# contigs (>= 1000 bp)	259
# contigs (>= 5000 bp)	239
# contigs (>= 10000 bp)	155
# contigs (>= 25000 bp)	27
# contigs (>= 50000 bp)	7
Total length (>= 0 bp)	6166020
Total length (>= 1000 bp)	6166020
Total length (>= 5000 bp)	6082730
Total length (>= 10000 bp)	5468999
Total length (>= 25000 bp)	3523512
Total length (>= 50000 bp)	2861880
# contigs	259
Largest contig	768598
Total length	6166020
GC (%)	57.32
N50	37097
N90	9416
auN	294964.5
L50	13
L90	164
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	20 + 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).

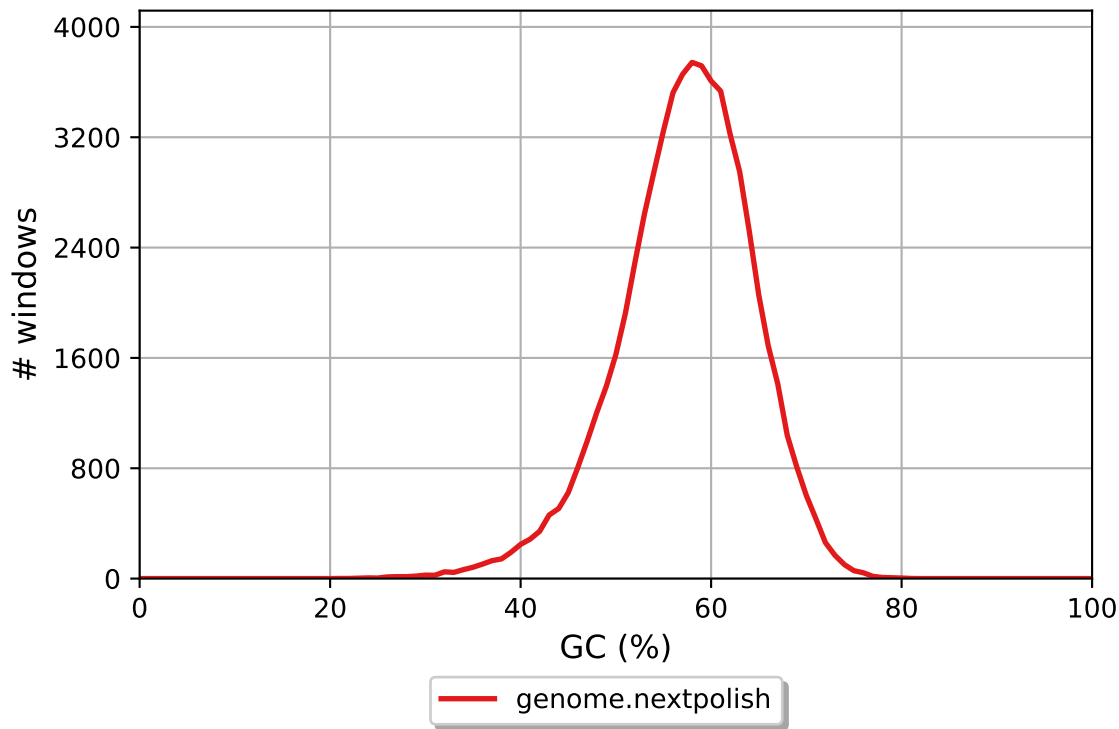


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

