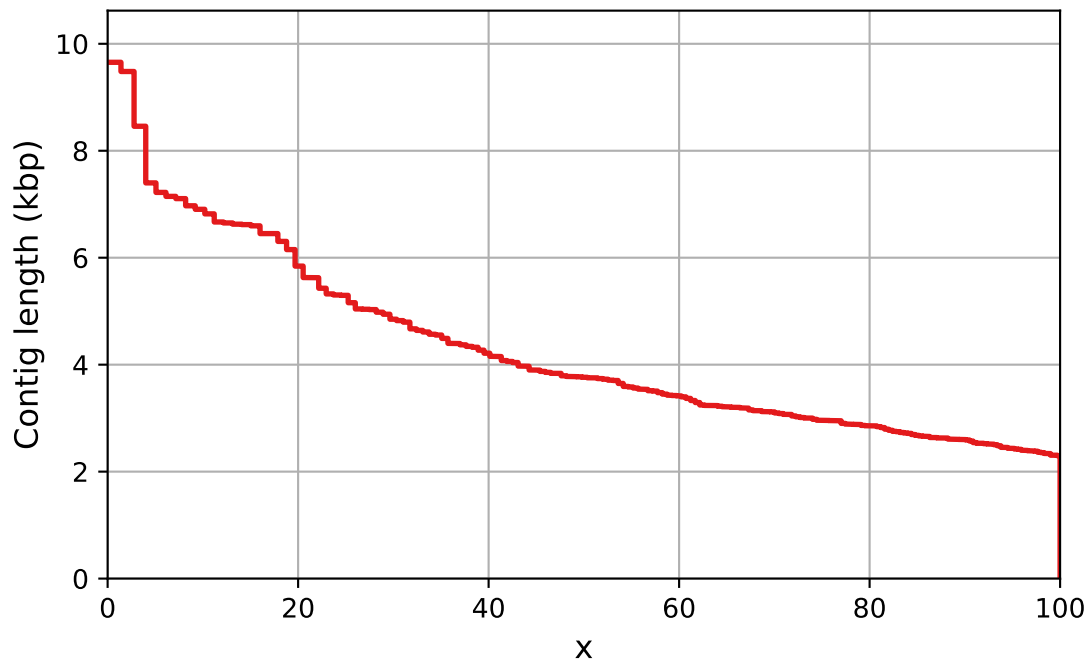


# Report

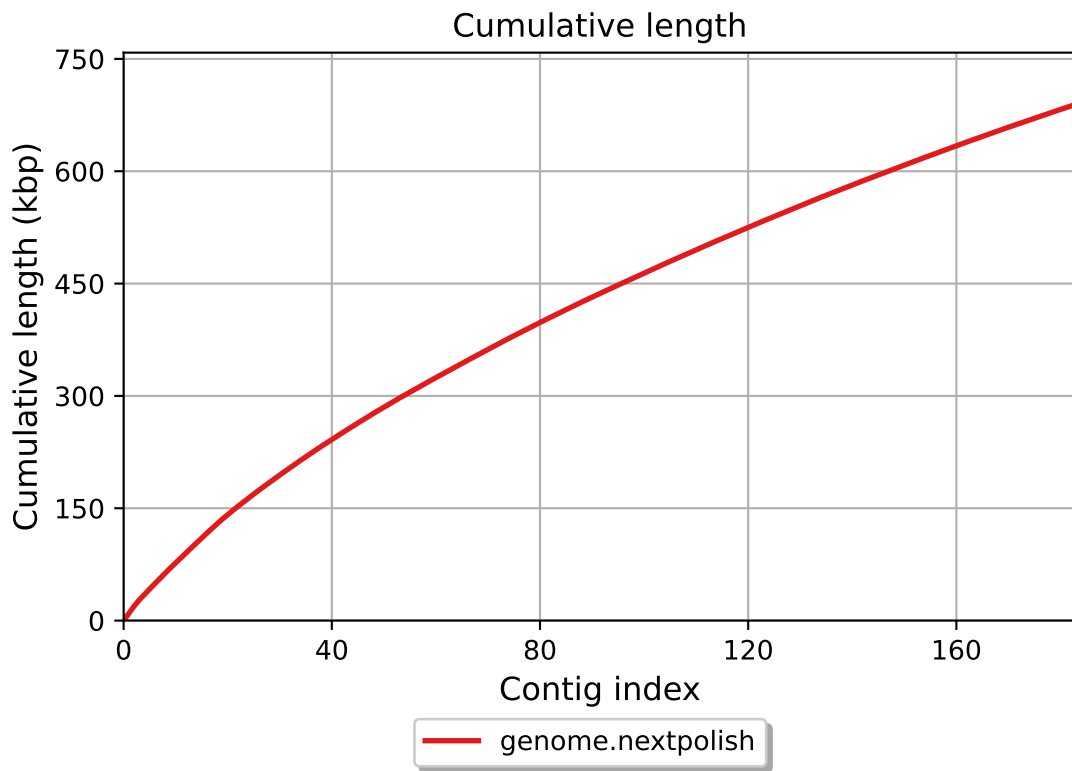
	genome.nextpolish
# contigs (>= 0 bp)	183
# contigs (>= 1000 bp)	183
# contigs (>= 5000 bp)	30
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	689423
Total length (>= 1000 bp)	689423
Total length (>= 5000 bp)	194397
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	183
Largest contig	9654
Total length	689423
GC (%)	50.19
N50	3764
N90	2599
auN	4314.2
L50	66
L90	155
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 1 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).

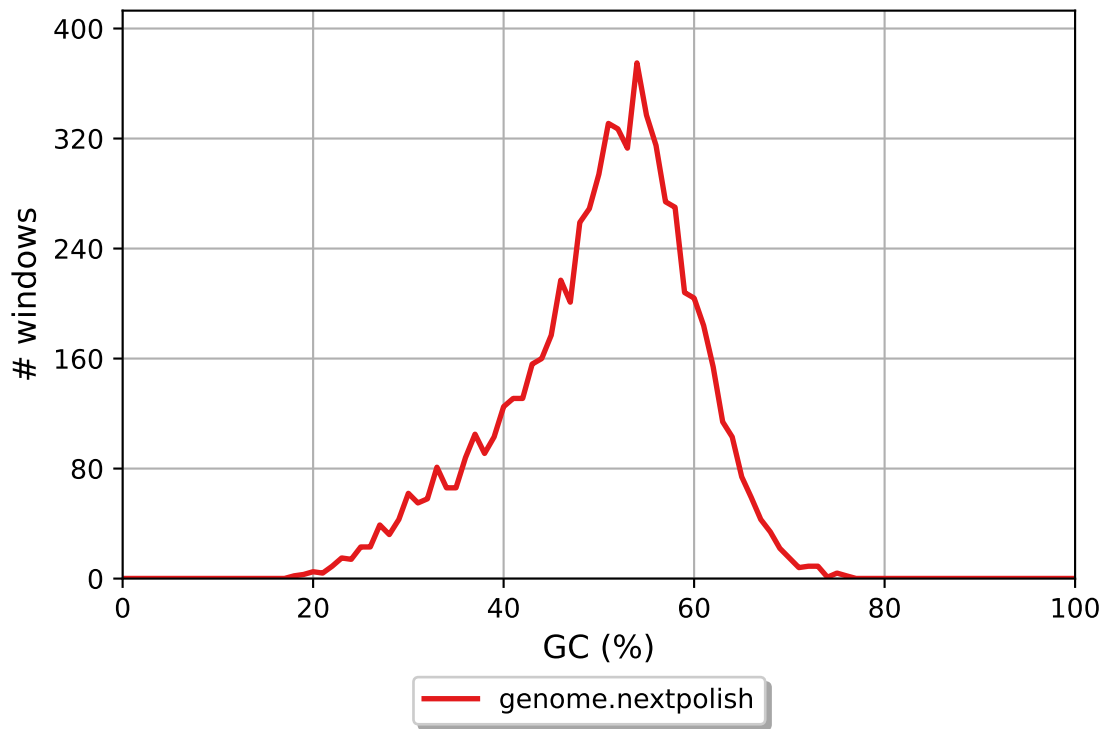
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

