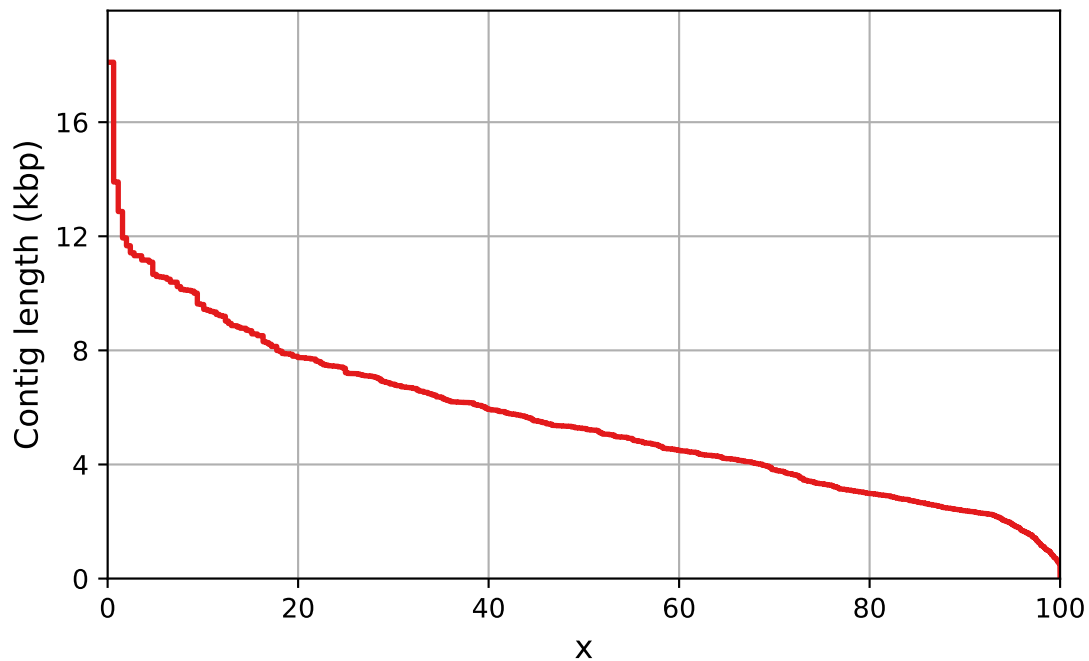


# Report

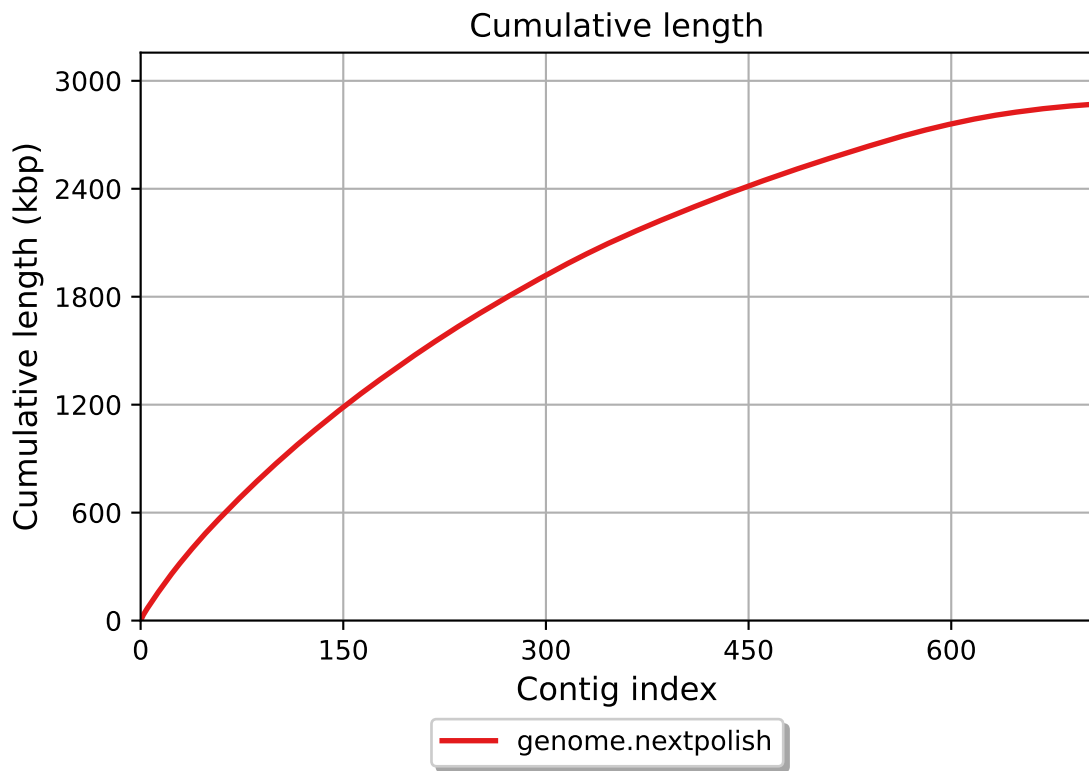
	genome.nextpolish
# contigs (>= 0 bp)	709
# contigs (>= 1000 bp)	653
# contigs (>= 5000 bp)	213
# contigs (>= 10000 bp)	24
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2870768
Total length (>= 1000 bp)	2831092
Total length (>= 5000 bp)	1526726
Total length (>= 10000 bp)	270301
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	705
Largest contig	18100
Total length	2869663
GC (%)	33.17
N50	5262
N90	2375
auN	5617.7
L50	196
L90	516
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 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).

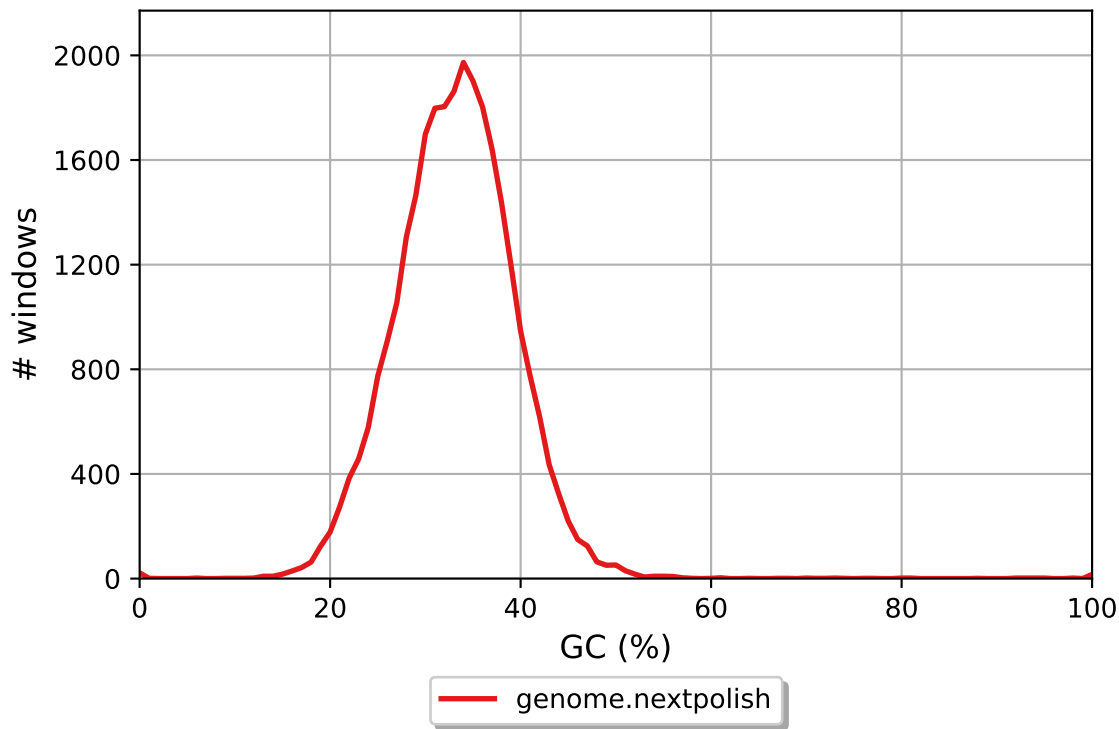
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

