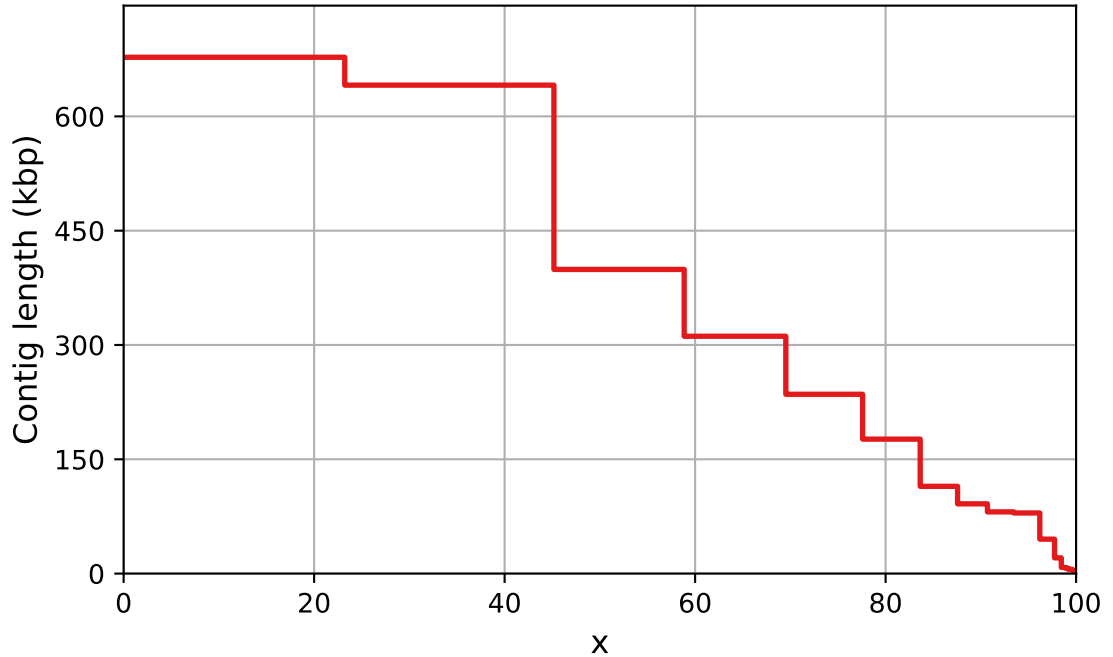


# Report

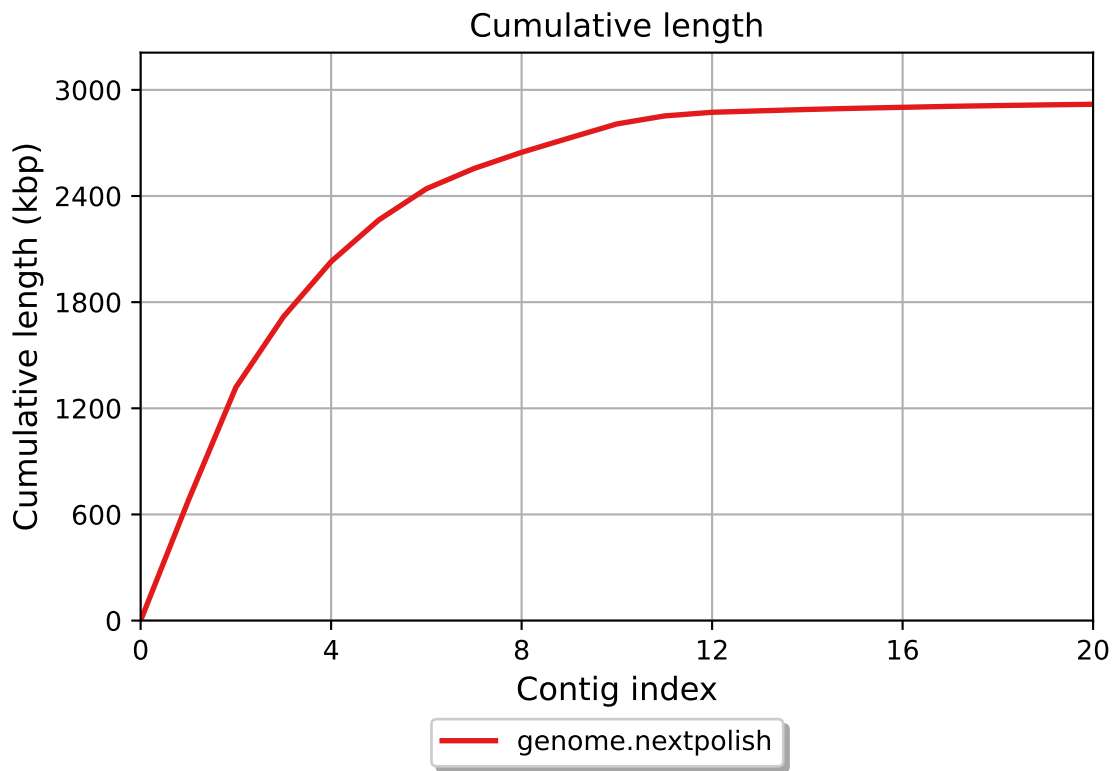
	genome.nextpolish
# contigs (>= 0 bp)	20
# contigs (>= 1000 bp)	20
# contigs (>= 5000 bp)	17
# contigs (>= 10000 bp)	12
# contigs (>= 25000 bp)	11
# contigs (>= 50000 bp)	10
Total length (>= 0 bp)	2918617
Total length (>= 1000 bp)	2918617
Total length (>= 5000 bp)	2906883
Total length (>= 10000 bp)	2873313
Total length (>= 25000 bp)	2852642
Total length (>= 50000 bp)	2807449
# contigs	20
Largest contig	677563
Total length	2918617
GC (%)	33.10
N50	399150
N90	91535
auN	428248.2
L50	3
L90	8
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 9 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).

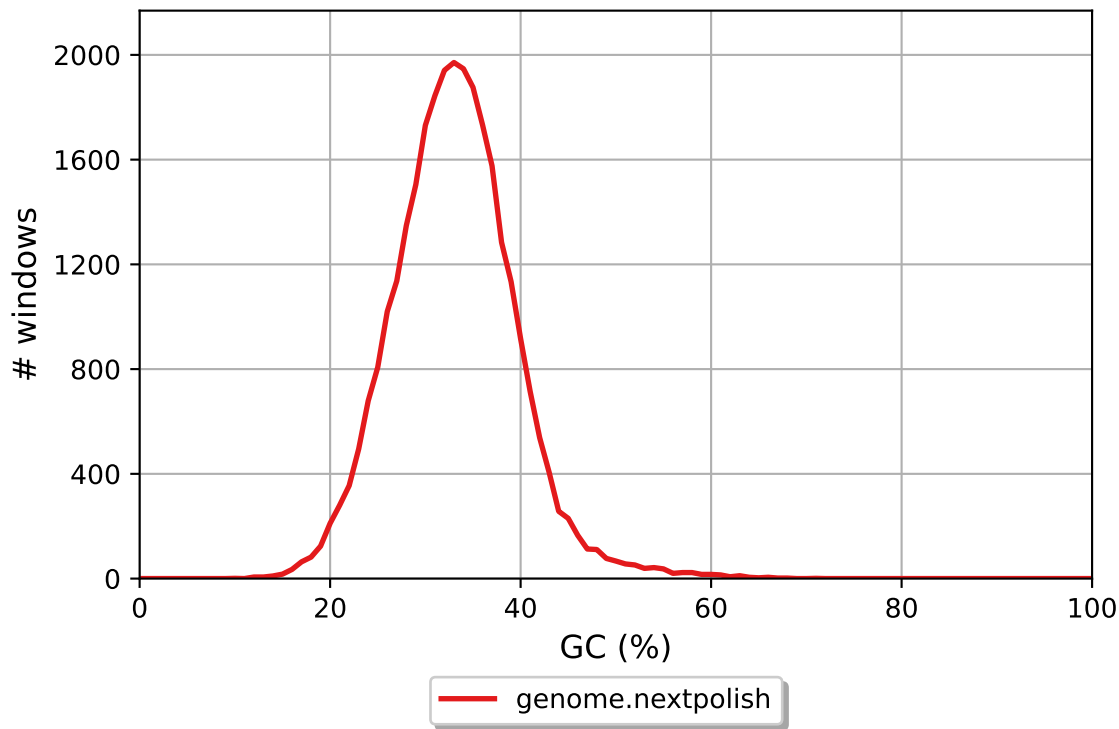
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

