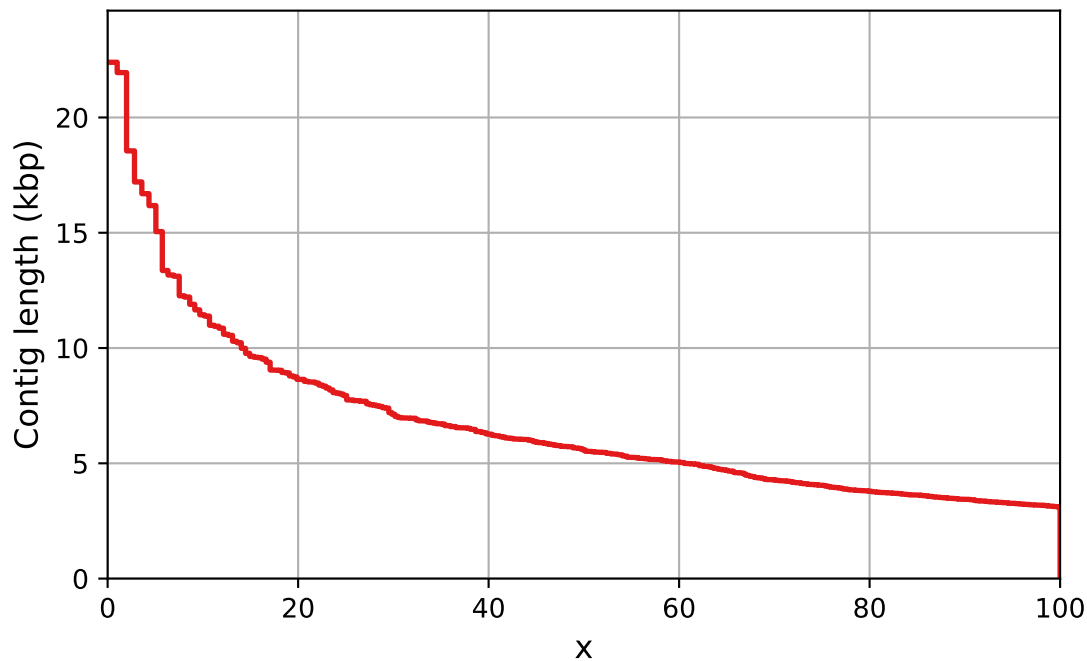


Report

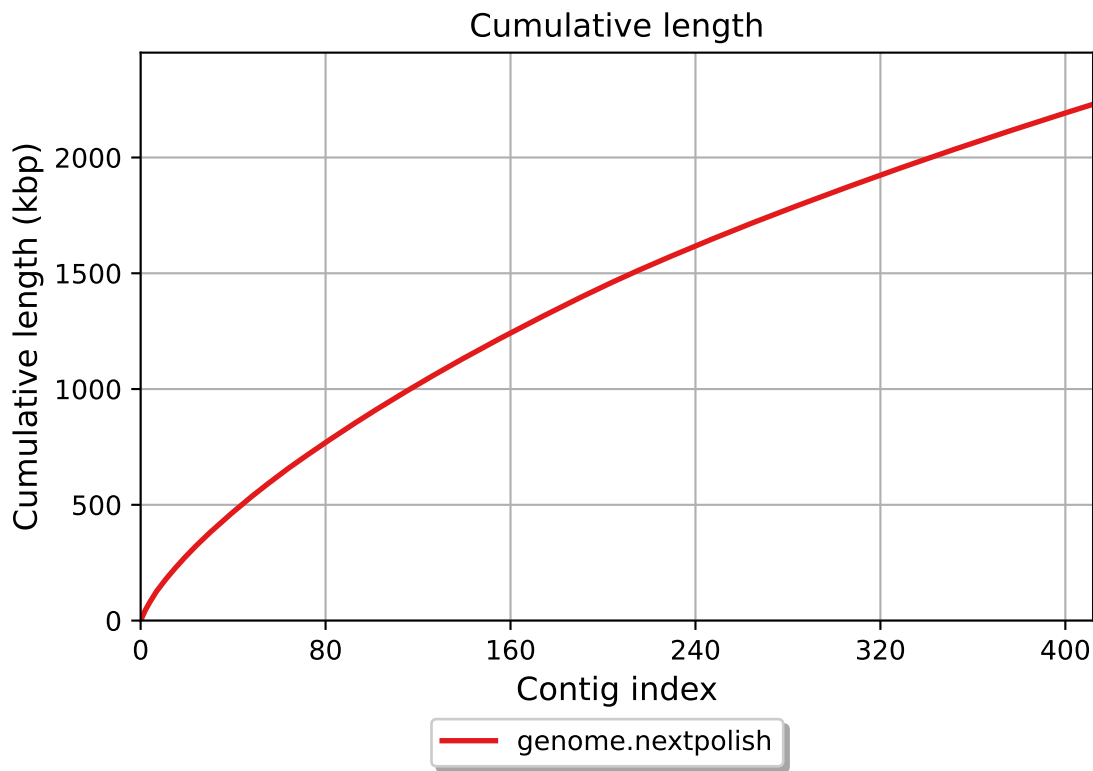
	genome.nextpolish
# contigs (>= 0 bp)	412
# contigs (>= 1000 bp)	412
# contigs (>= 5000 bp)	181
# contigs (>= 10000 bp)	23
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2229893
Total length (>= 1000 bp)	2229893
Total length (>= 5000 bp)	1349775
Total length (>= 10000 bp)	312976
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	412
Largest contig	22394
Total length	2229893
GC (%)	34.66
N50	5582
N90	3438
auN	6736.5
L50	137
L90	344
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 0 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).

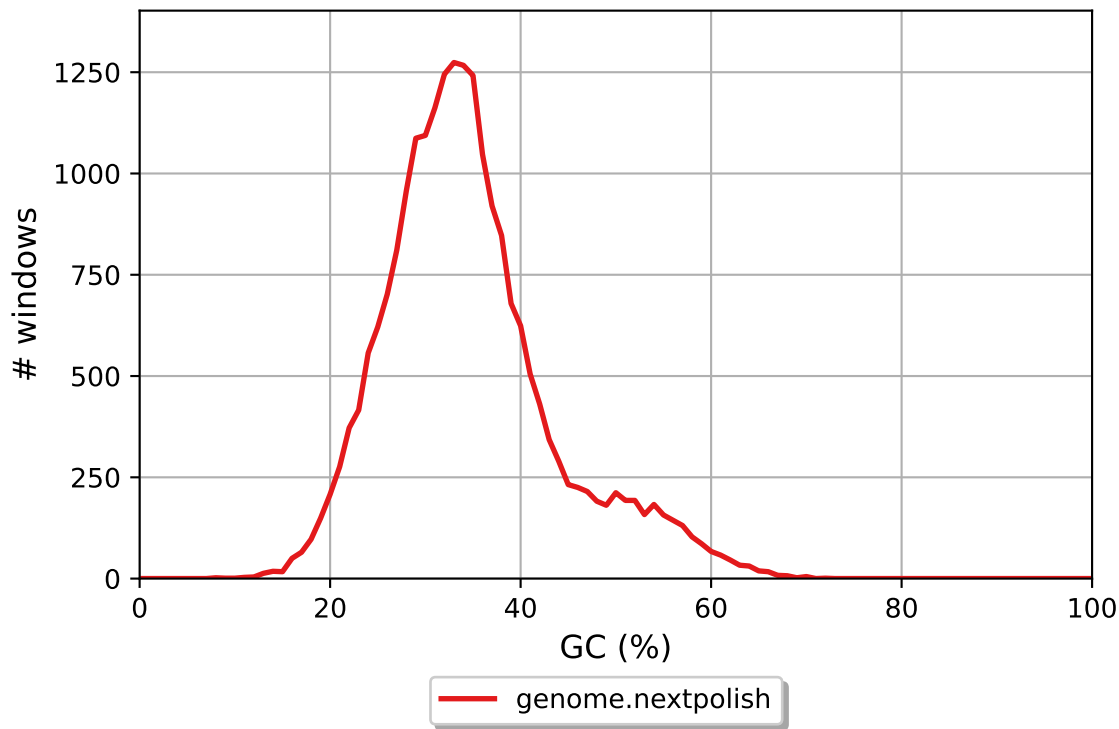
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

