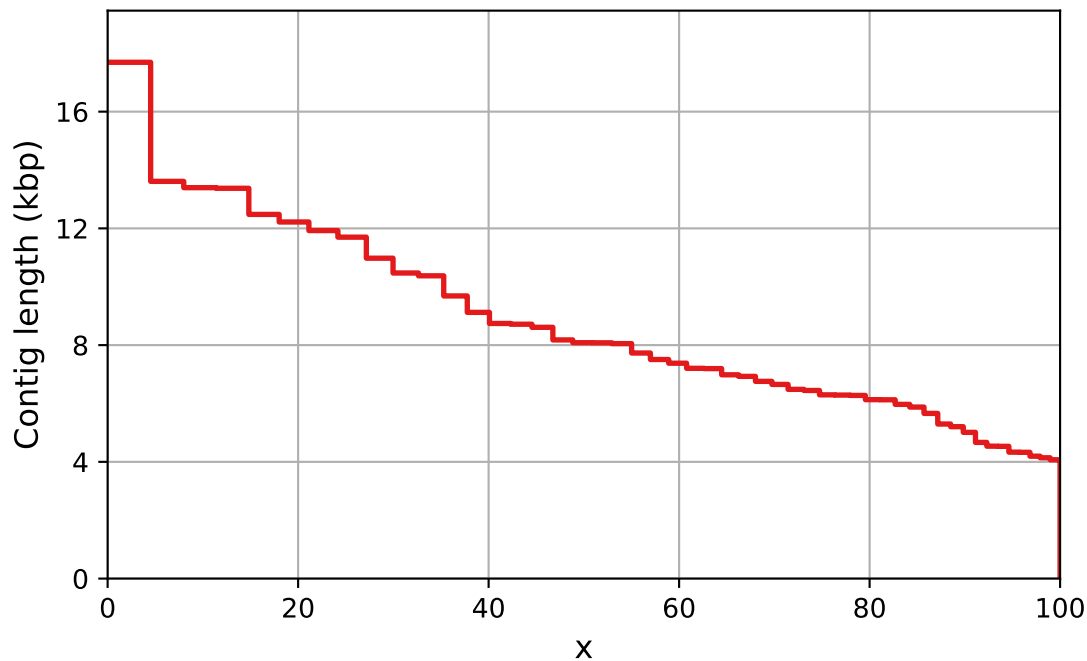


Report

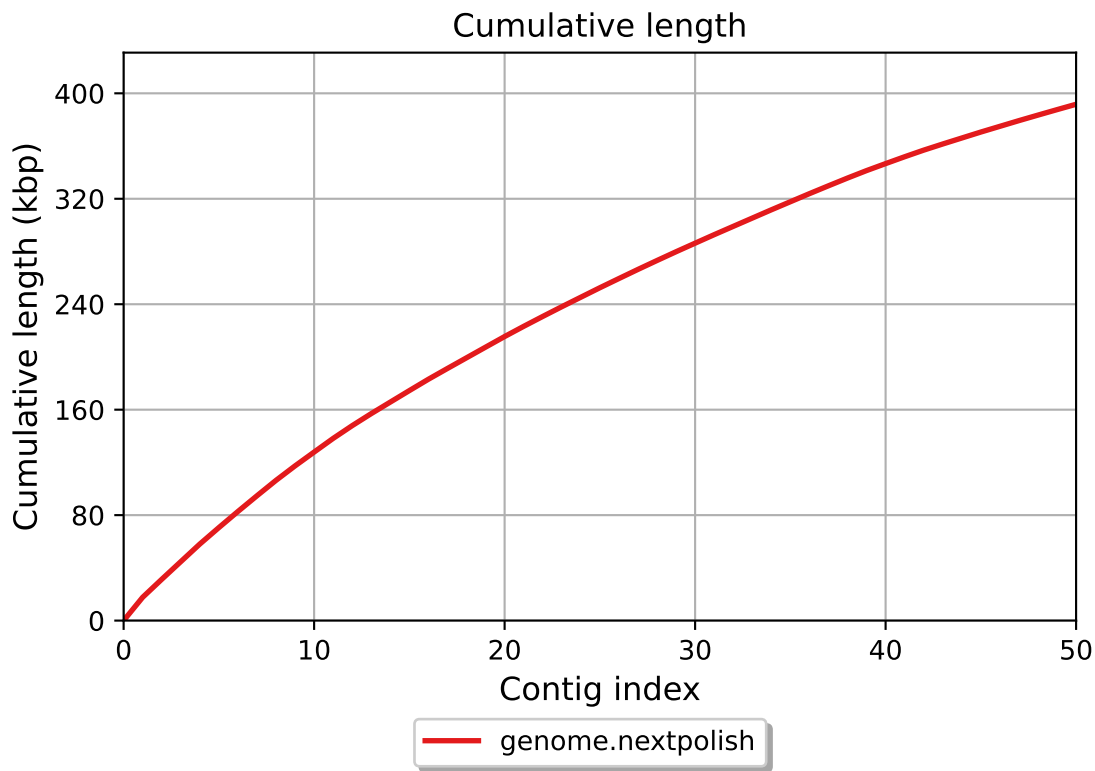
	genome.nextpolish
# contigs (>= 0 bp)	50
# contigs (>= 1000 bp)	50
# contigs (>= 5000 bp)	42
# contigs (>= 10000 bp)	11
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	391689
Total length (>= 1000 bp)	391689
Total length (>= 5000 bp)	356889
Total length (>= 10000 bp)	138217
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	50
Largest contig	17691
Total length	391689
GC (%)	34.13
N50	8083
N90	5011
auN	8978.2
L50	18
L90	42
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 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).

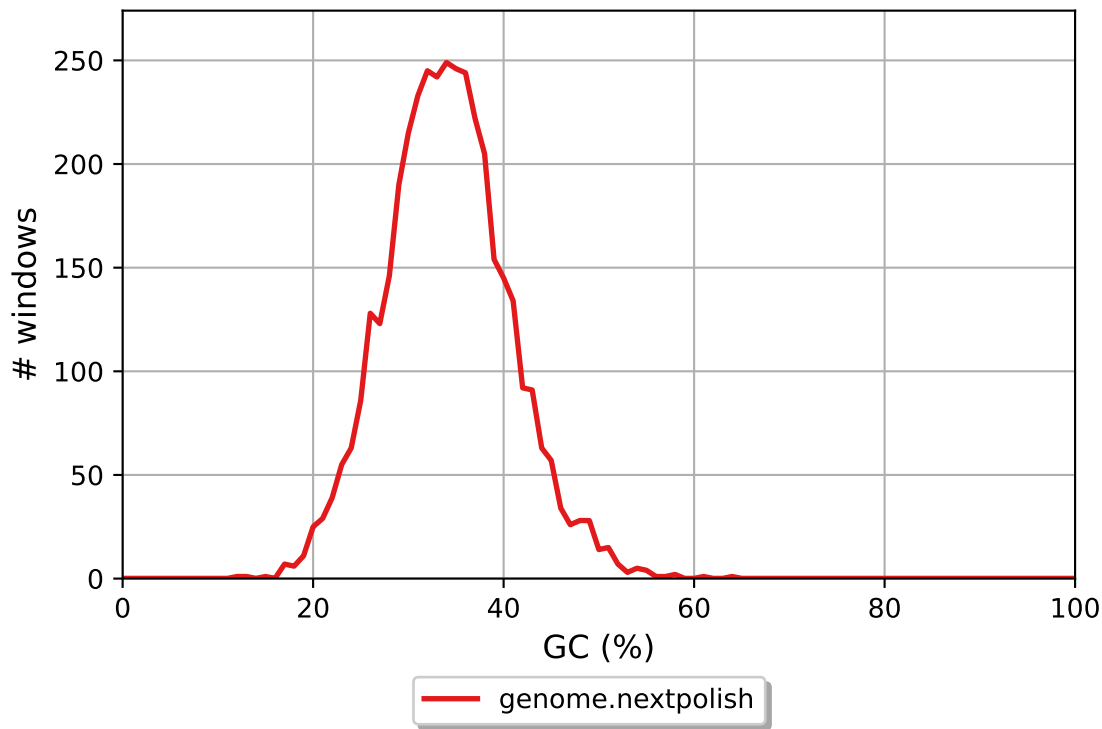
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

