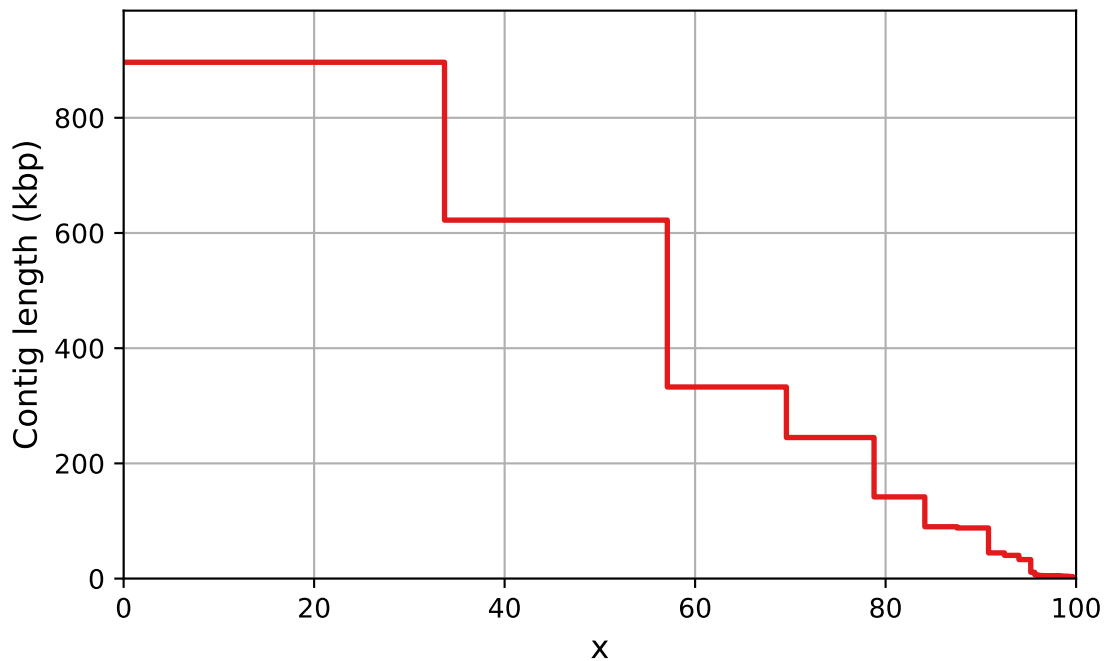


Report

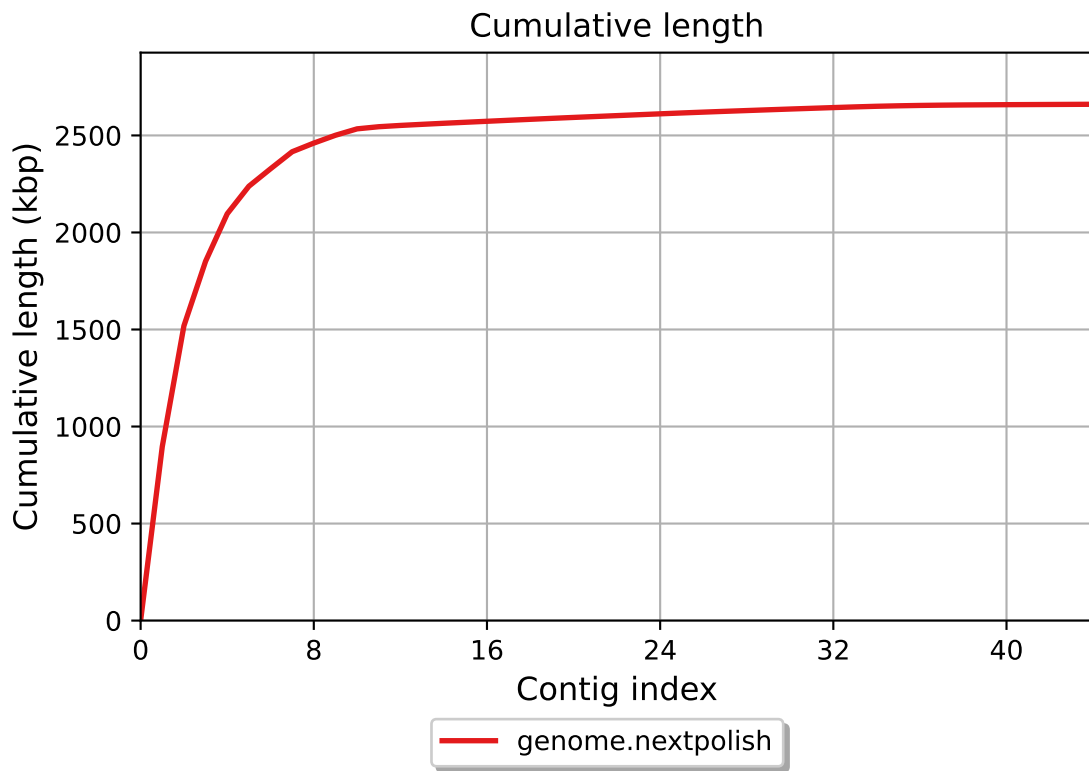
	genome.nextpolish
# contigs (>= 0 bp)	46
# contigs (>= 1000 bp)	38
# contigs (>= 5000 bp)	17
# contigs (>= 10000 bp)	11
# contigs (>= 25000 bp)	10
# contigs (>= 50000 bp)	7
Total length (>= 0 bp)	2661424
Total length (>= 1000 bp)	2657107
Total length (>= 5000 bp)	2578072
Total length (>= 10000 bp)	2544893
Total length (>= 25000 bp)	2533919
Total length (>= 50000 bp)	2415962
# contigs	44
Largest contig	896385
Total length	2660606
GC (%)	32.16
N50	622321
N90	87884
auN	527207.1
L50	2
L90	7
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	12 + 3 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).

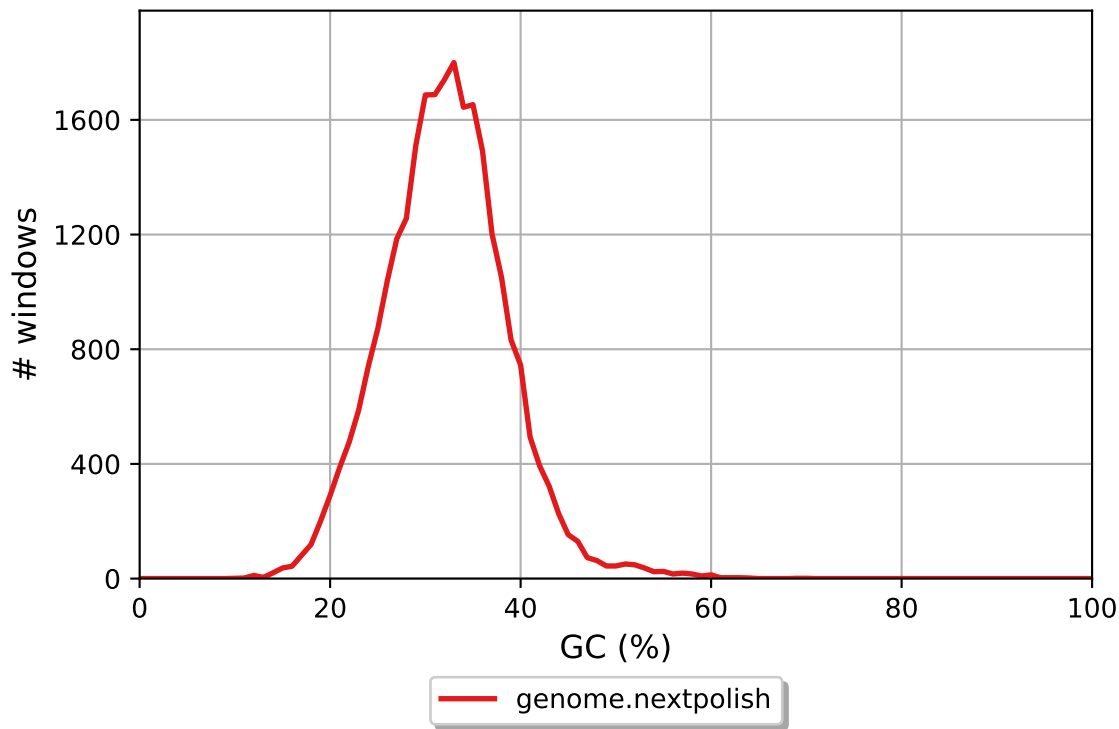
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

