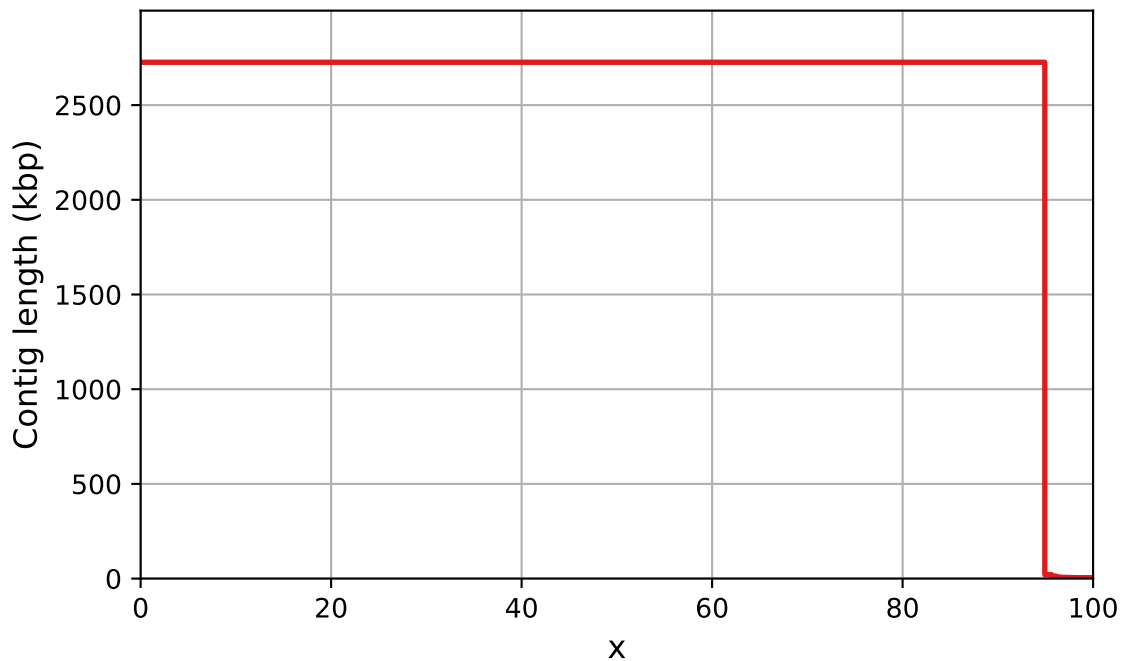


# Report

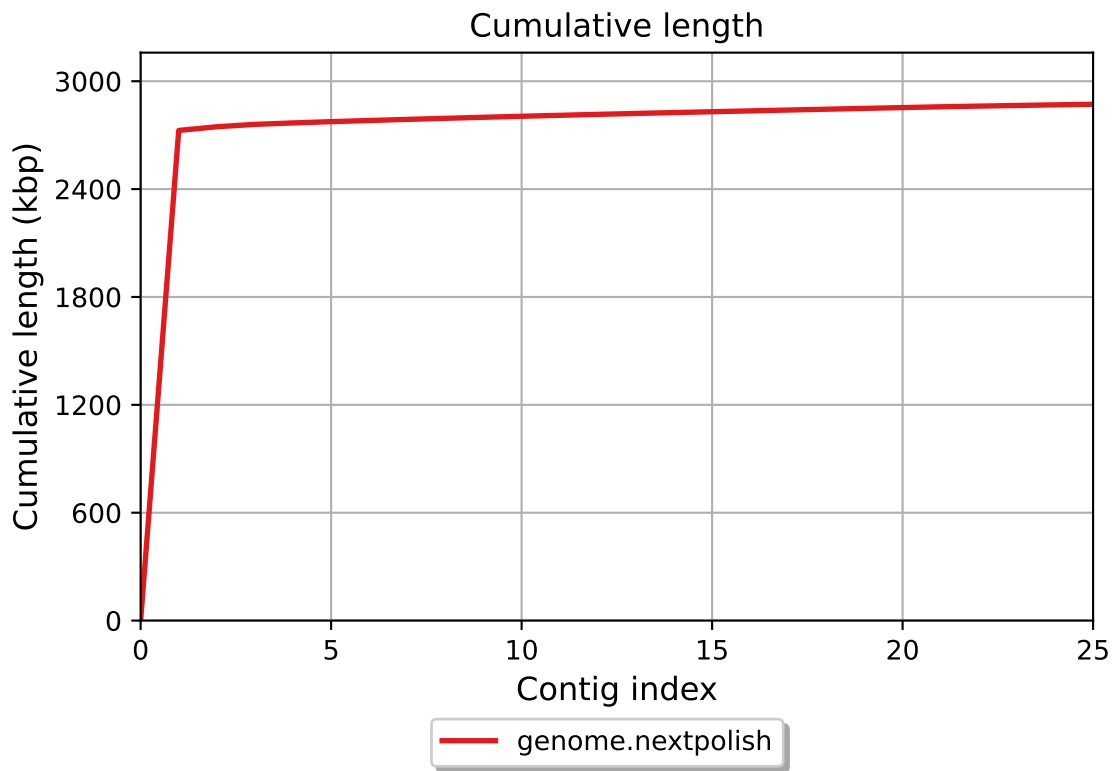
	genome.nextpolish
# contigs (>= 0 bp)	25
# contigs (>= 1000 bp)	25
# contigs (>= 5000 bp)	13
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	2871822
Total length (>= 1000 bp)	2871822
Total length (>= 5000 bp)	2820758
Total length (>= 10000 bp)	2760039
Total length (>= 25000 bp)	2726221
Total length (>= 50000 bp)	2726221
# contigs	25
Largest contig	2726221
Total length	2871822
GC (%)	33.67
N50	2726221
N90	2726221
auN	2588418.4
L50	1
L90	1
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	10 + 6 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).

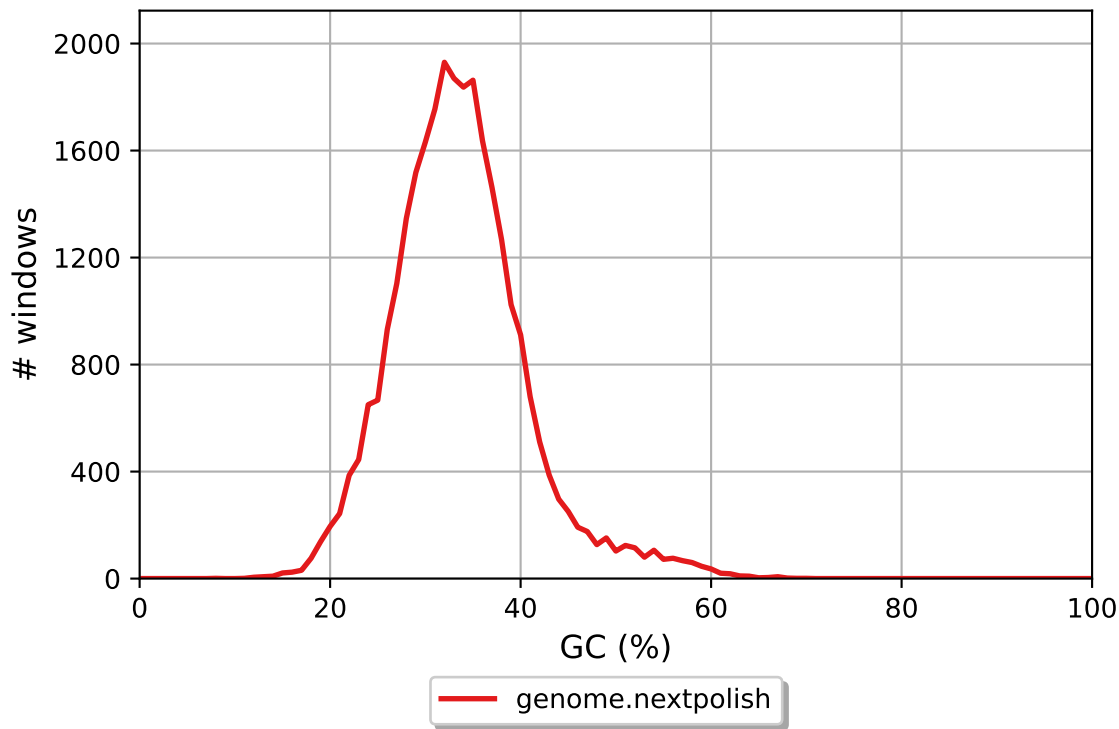
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

