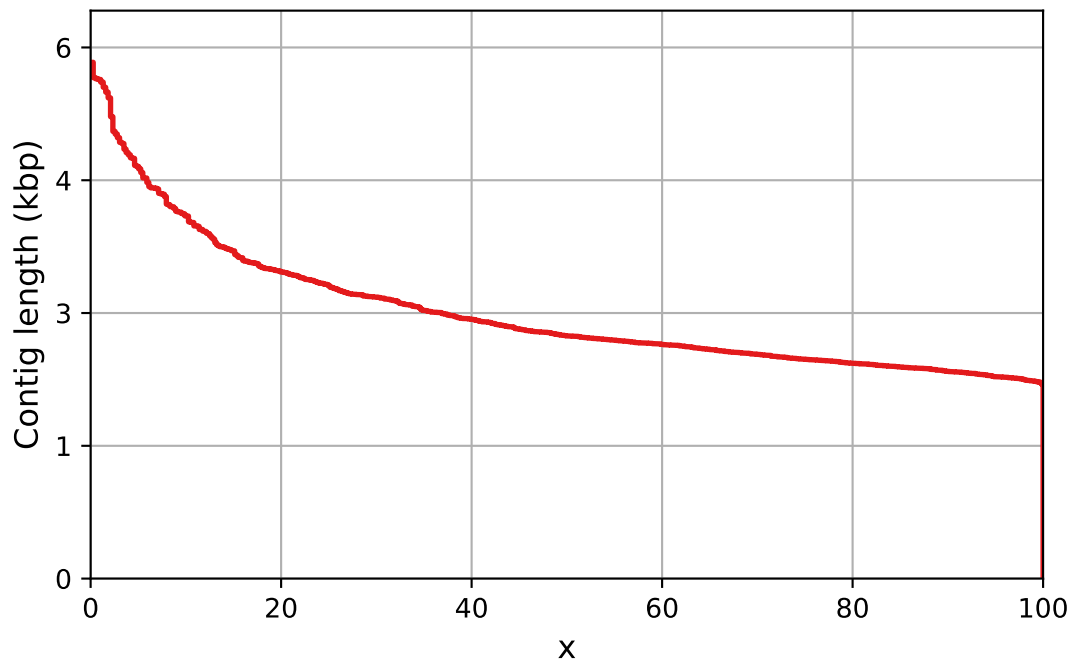


Report

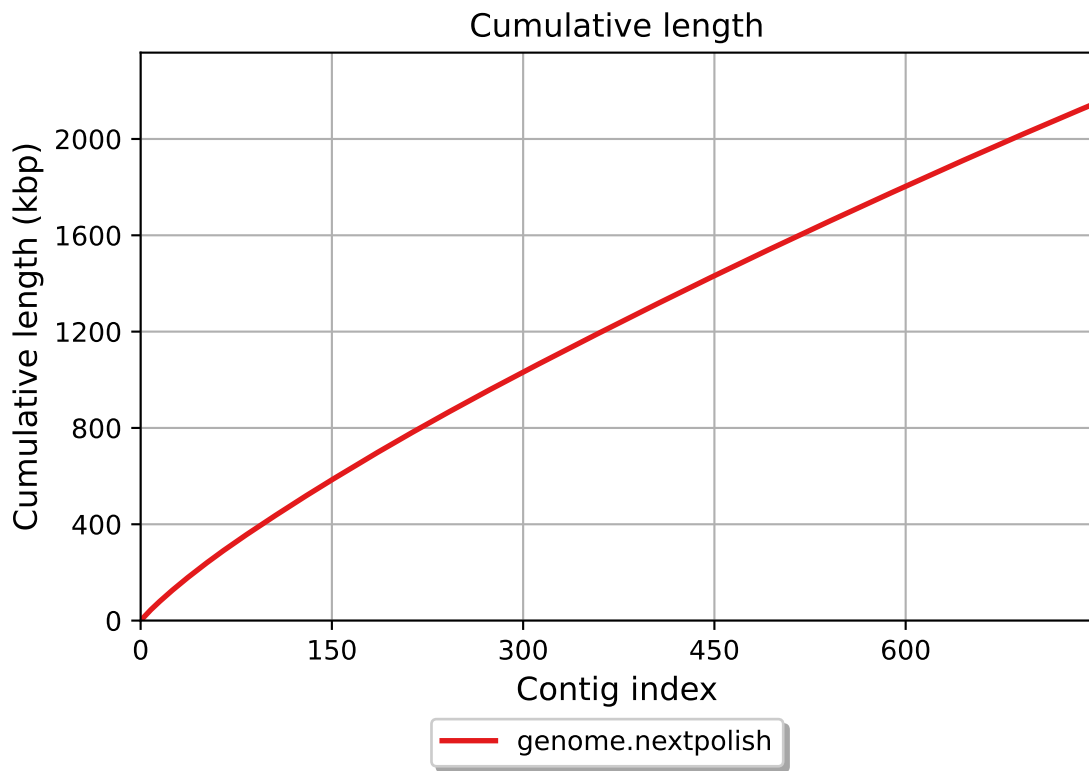
	genome.nextpolish
# contigs (>= 0 bp)	747
# contigs (>= 1000 bp)	747
# contigs (>= 5000 bp)	11
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2144115
Total length (>= 1000 bp)	2144115
Total length (>= 5000 bp)	60151
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	747
Largest contig	5832
Total length	2144115
GC (%)	55.34
N50	2742
N90	2340
auN	3009.5
L50	315
L90	654
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 2 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).

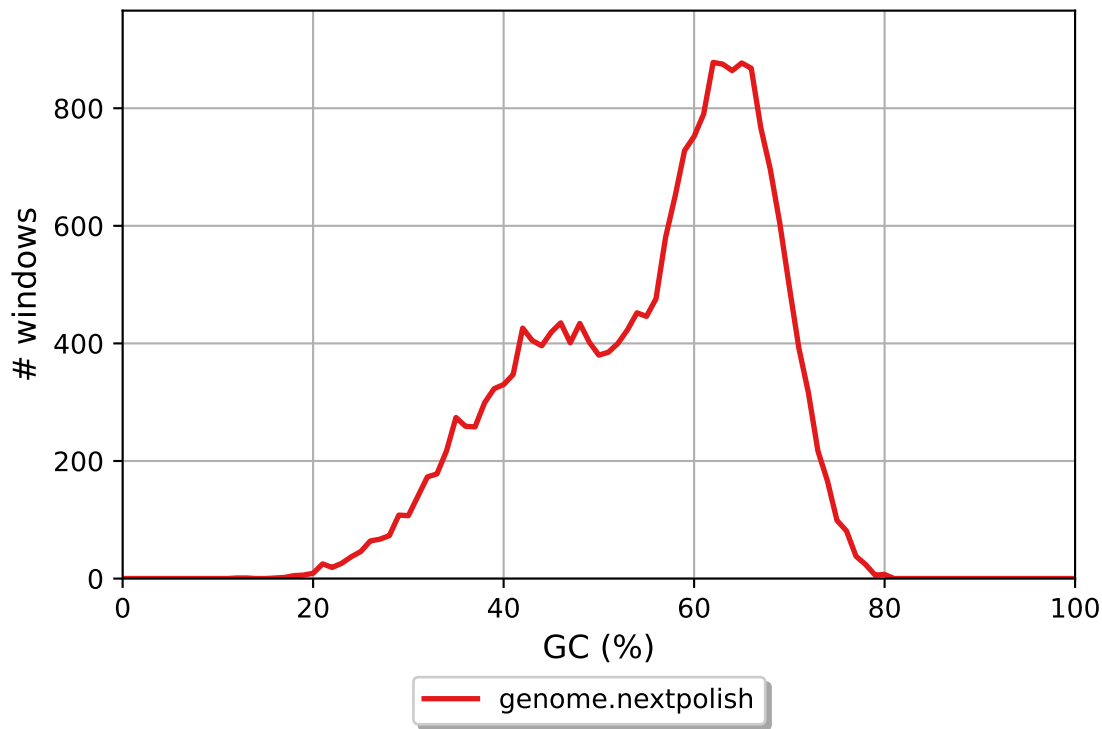
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

