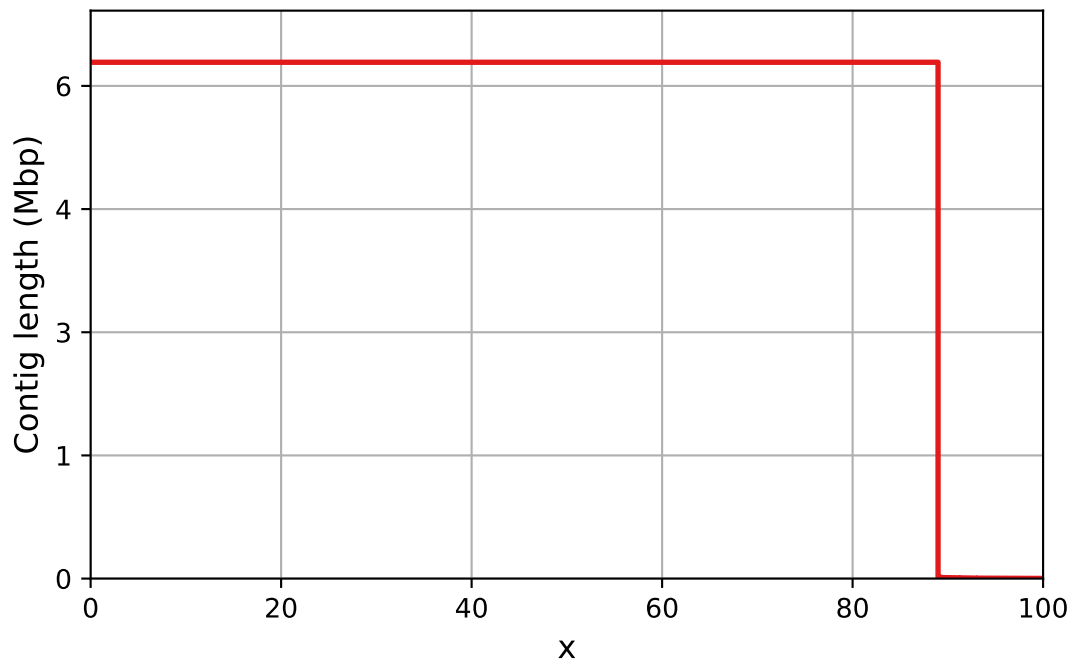


Report

	genome.nextpolish
# contigs (>= 0 bp)	191
# contigs (>= 1000 bp)	191
# contigs (>= 5000 bp)	40
# contigs (>= 10000 bp)	9
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	7067362
Total length (>= 1000 bp)	7067362
Total length (>= 5000 bp)	6601488
Total length (>= 10000 bp)	6384572
Total length (>= 25000 bp)	6287920
Total length (>= 50000 bp)	6287920
# contigs	191
Largest contig	6287920
Total length	7067362
GC (%)	58.17
N50	6287920
N90	11267
auN	5595053.2
L50	1
L90	7
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	23 + 0 part

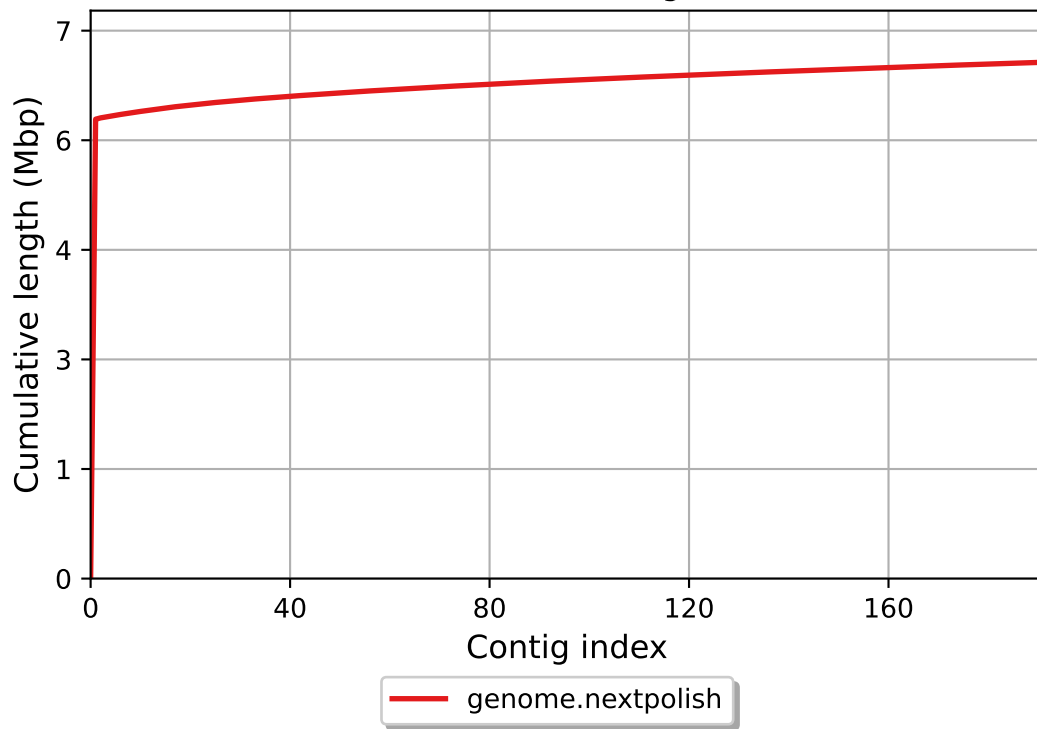
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

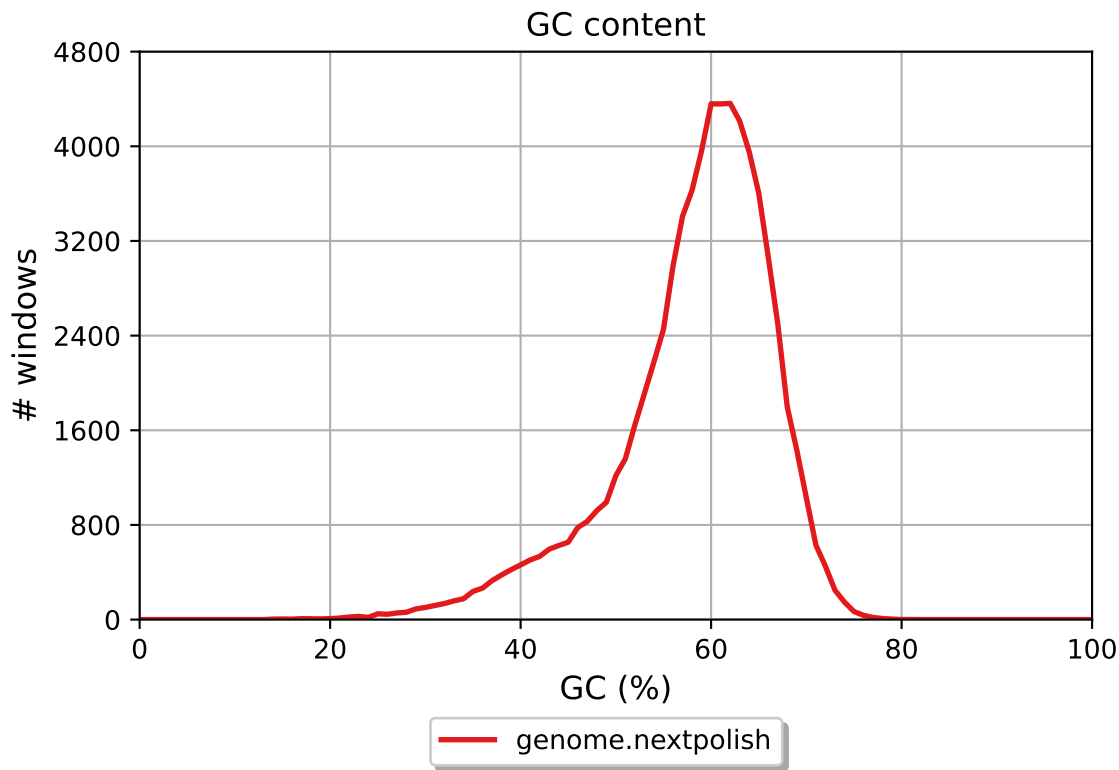
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

