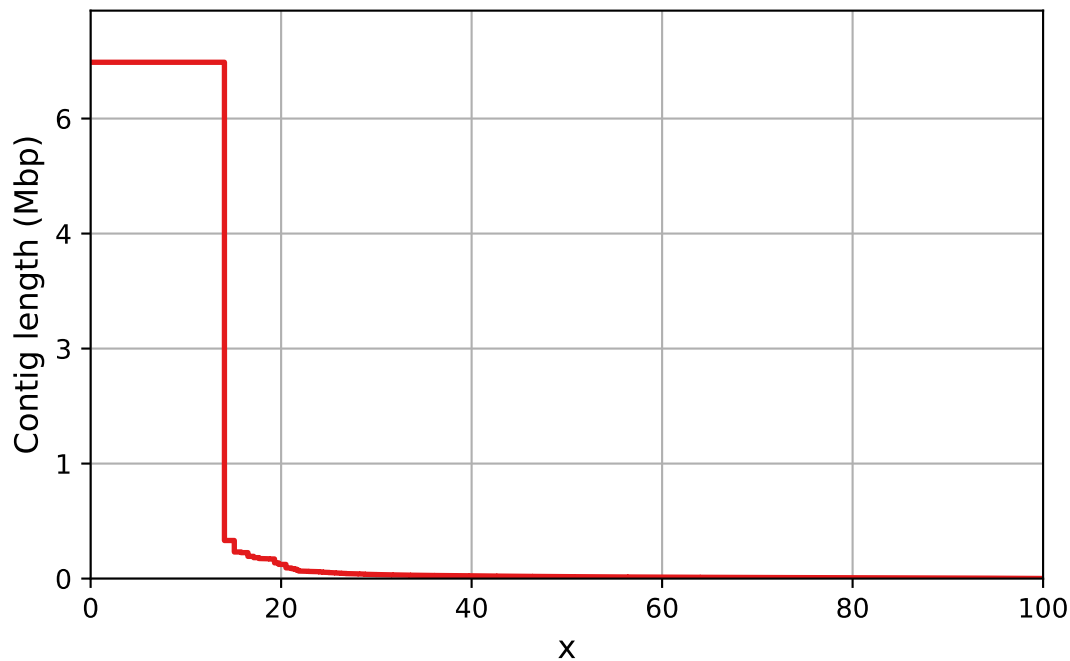


Report

	genome.nextpolish
# contigs (>= 0 bp)	2920
# contigs (>= 1000 bp)	2901
# contigs (>= 5000 bp)	2190
# contigs (>= 10000 bp)	1350
# contigs (>= 25000 bp)	352
# contigs (>= 50000 bp)	77
Total length (>= 0 bp)	47933268
Total length (>= 1000 bp)	47918907
Total length (>= 5000 bp)	45615321
Total length (>= 10000 bp)	39442558
Total length (>= 25000 bp)	24144919
Total length (>= 50000 bp)	14642331
# contigs	2919
Largest contig	6734089
Total length	47932771
GC (%)	54.62
N50	25473
N90	7168
auN	987497.6
L50	345
L90	1781
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	61 + 4 part

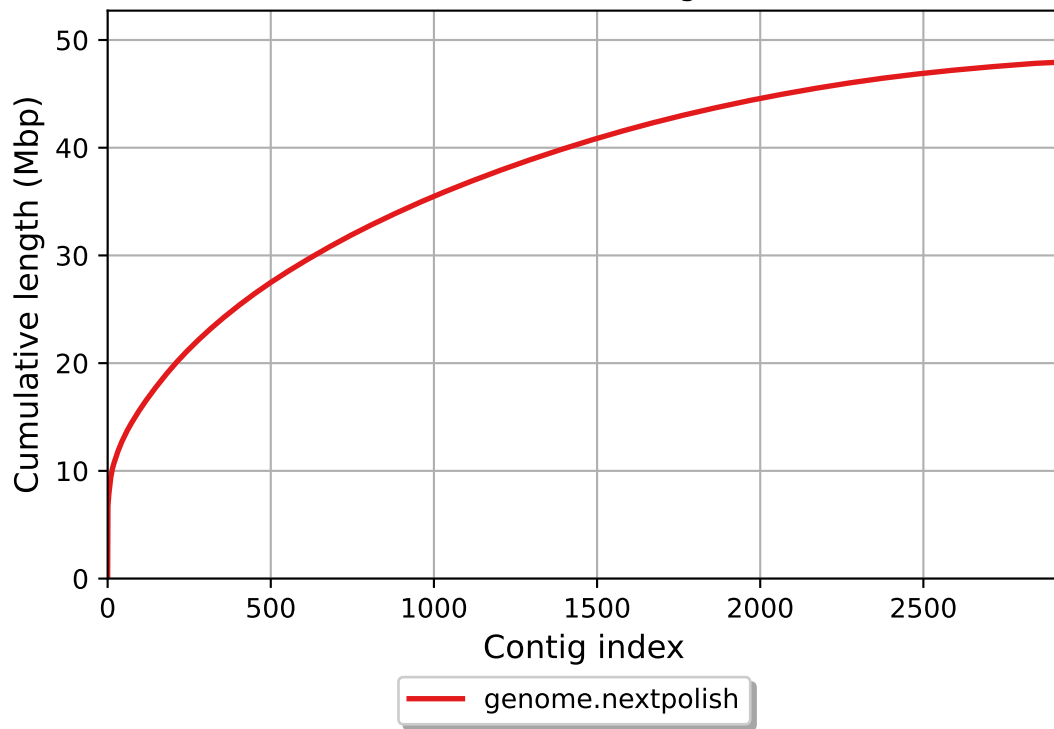
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

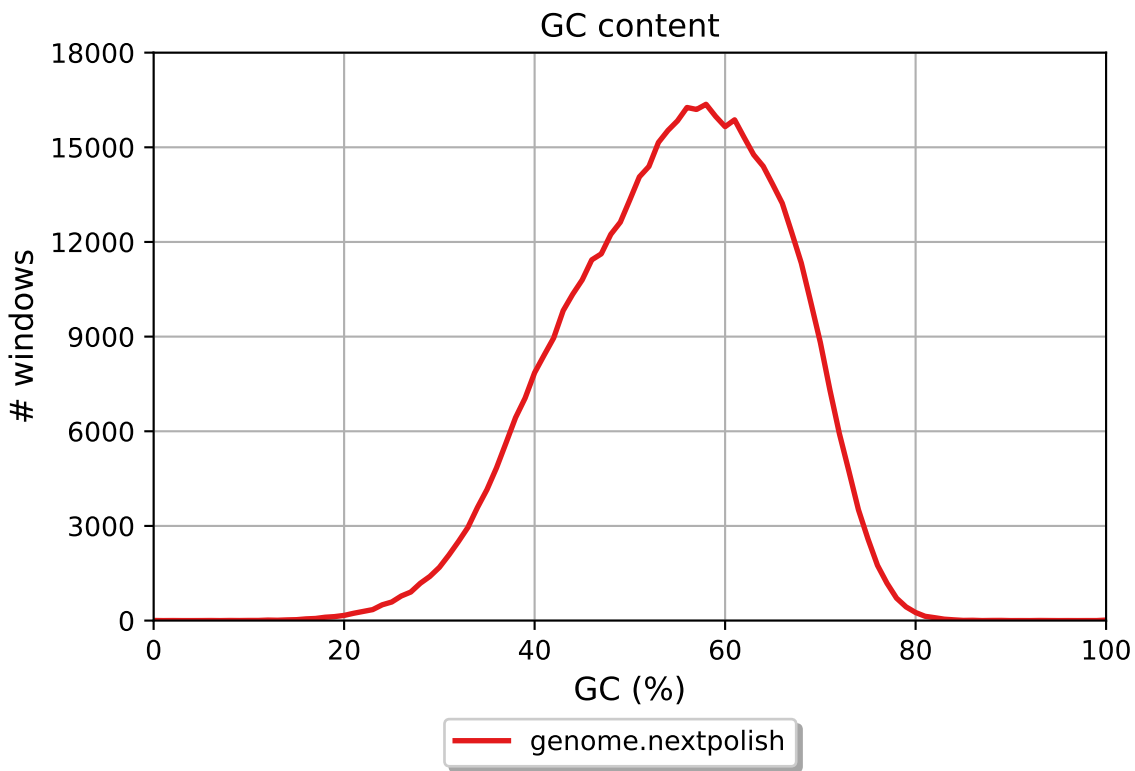
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

