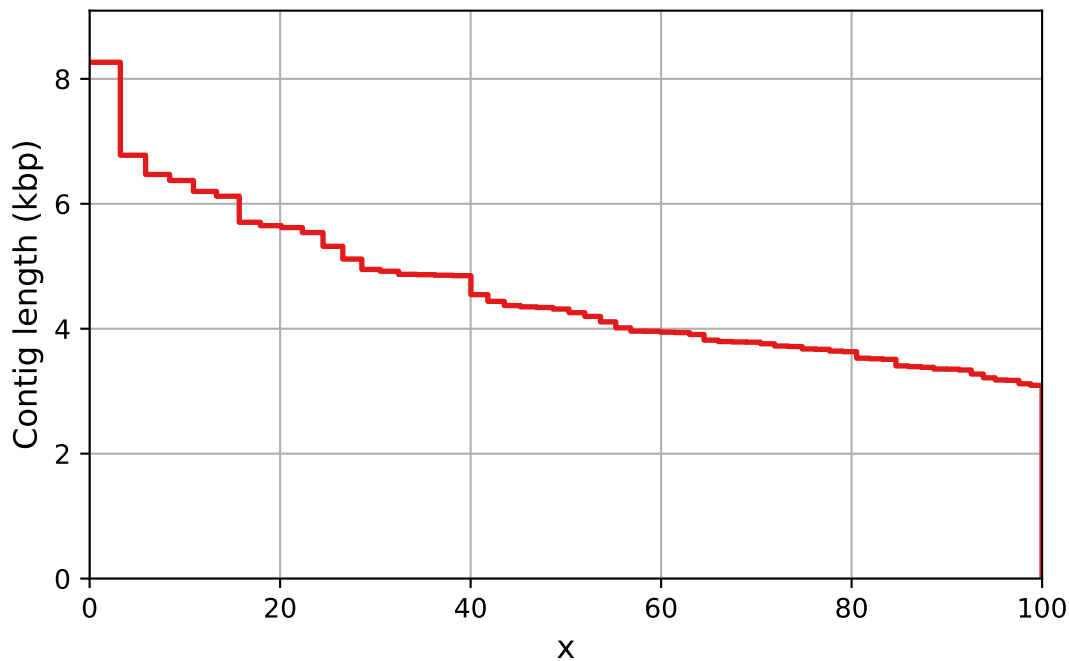


Report

	genome.nextpolish
# contigs (>= 0 bp)	59
# contigs (>= 1000 bp)	59
# contigs (>= 5000 bp)	12
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	255942
Total length (>= 1000 bp)	255942
Total length (>= 5000 bp)	73149
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	59
Largest contig	8265
Total length	255942
GC (%)	42.57
N50	4315
N90	3353
auN	4602.3
L50	24
L90	52
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 0 part

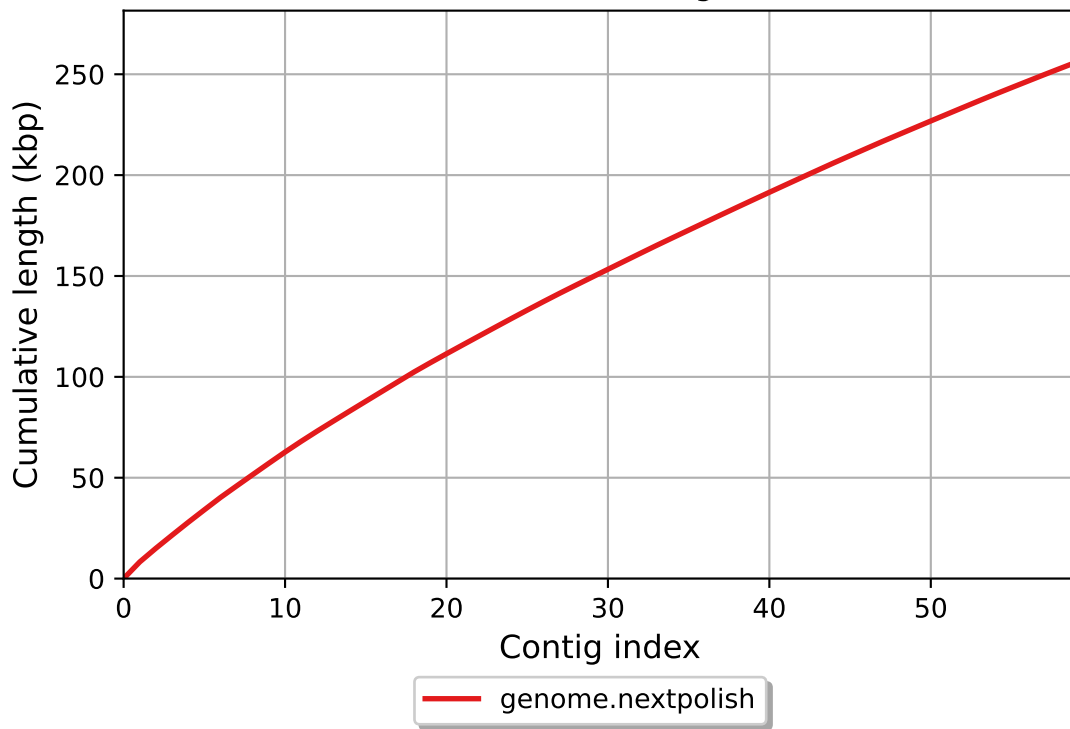
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

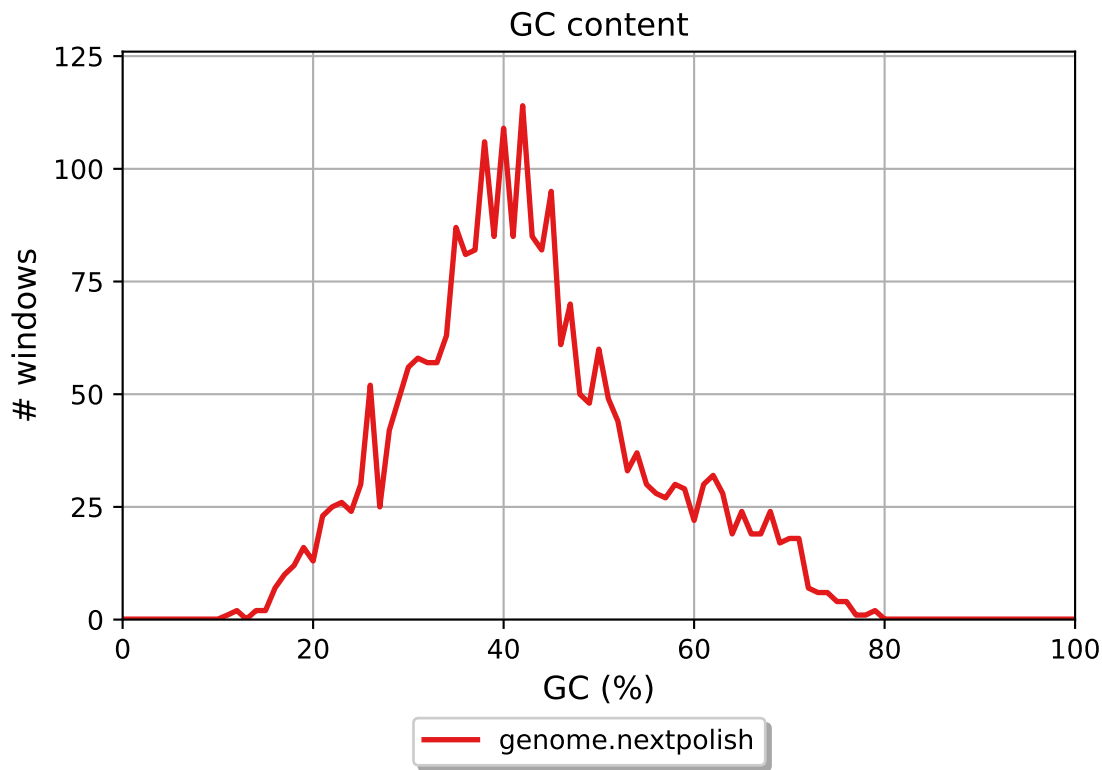
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

