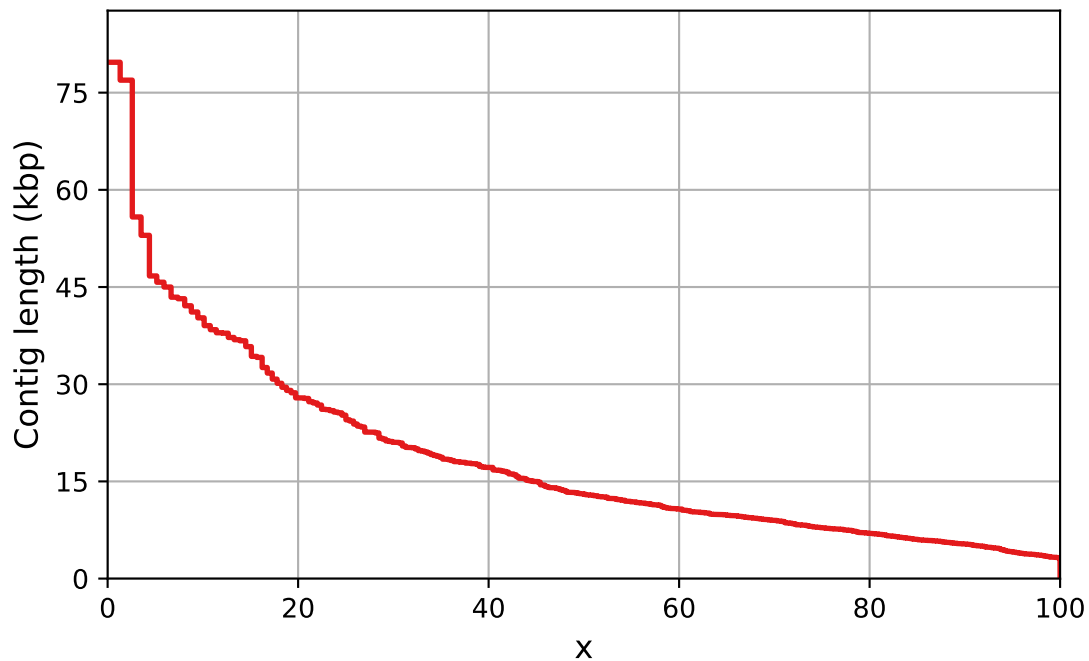


Report

	genome.nextpolish
# contigs (>= 0 bp)	564
# contigs (>= 1000 bp)	563
# contigs (>= 5000 bp)	438
# contigs (>= 10000 bp)	197
# contigs (>= 25000 bp)	41
# contigs (>= 50000 bp)	4
Total length (>= 0 bp)	6048941
Total length (>= 1000 bp)	6048393
Total length (>= 5000 bp)	5554769
Total length (>= 10000 bp)	3823165
Total length (>= 25000 bp)	1513134
Total length (>= 50000 bp)	265411
# contigs	564
Largest contig	79689
Total length	6048941
GC (%)	57.03
N50	13037
N90	5331
auN	18741.8
L50	128
L90	417
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 2 part

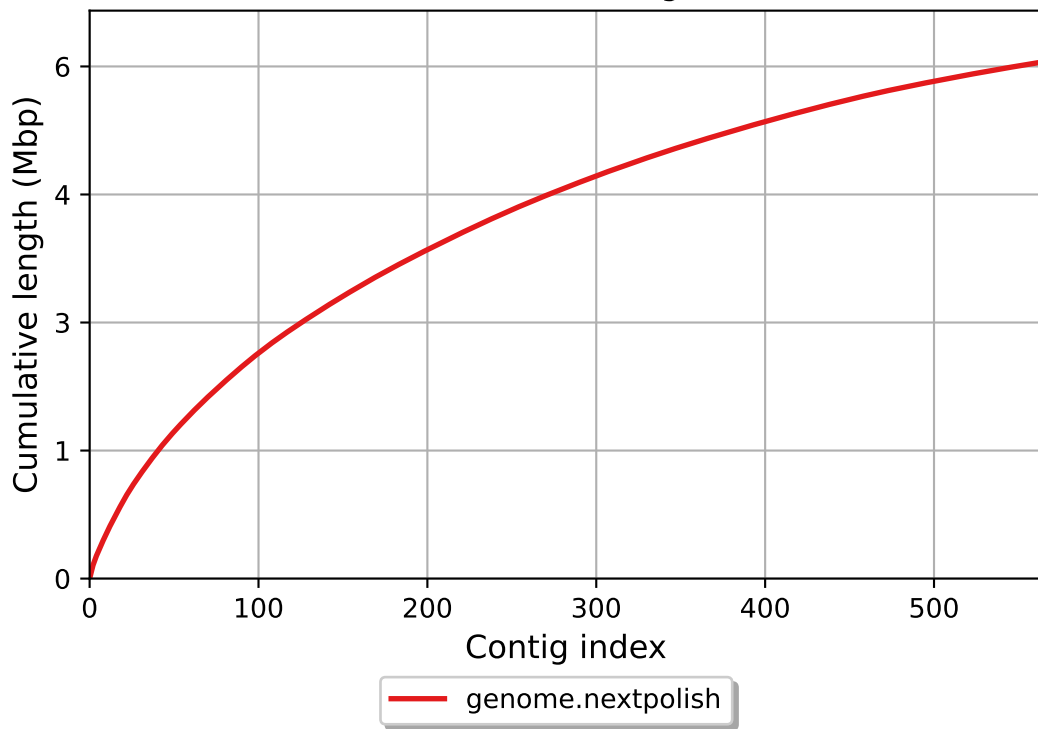
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

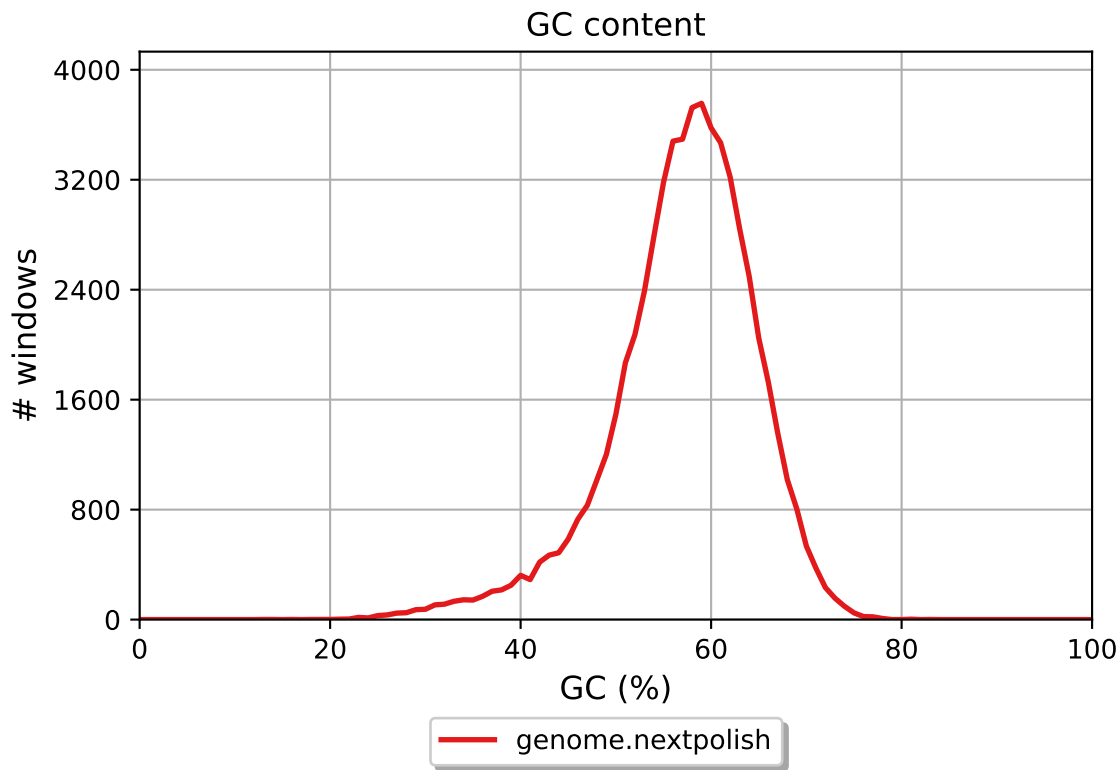
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

