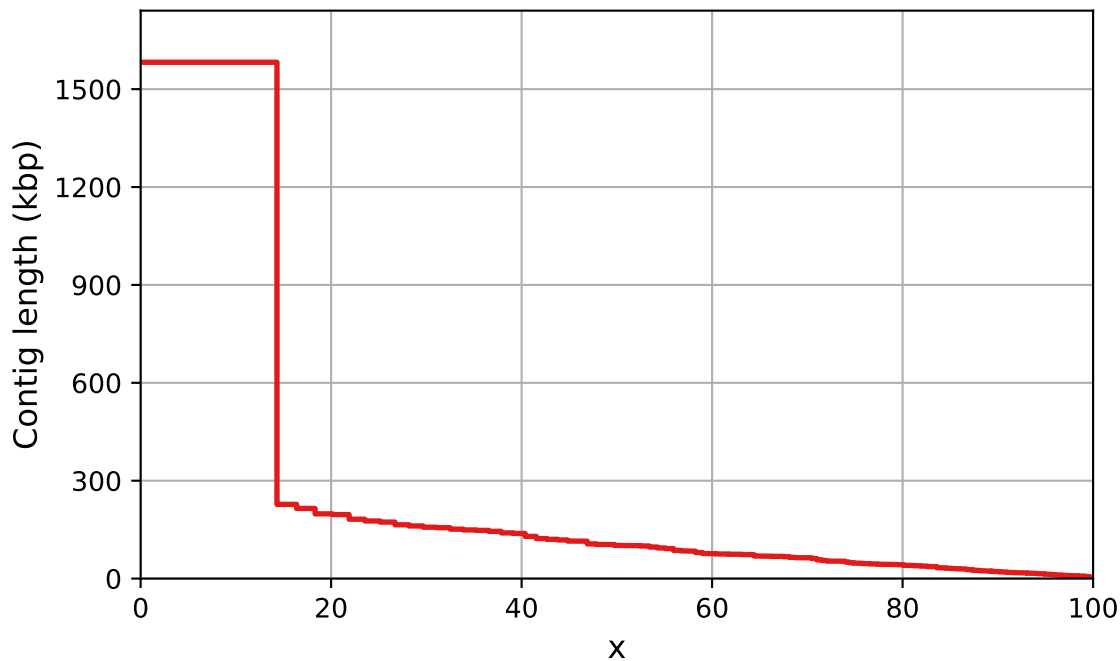


Report

	genome.nextpolish
# contigs (>= 0 bp)	213
# contigs (>= 1000 bp)	213
# contigs (>= 5000 bp)	200
# contigs (>= 10000 bp)	164
# contigs (>= 25000 bp)	105
# contigs (>= 50000 bp)	64
Total length (>= 0 bp)	11053795
Total length (>= 1000 bp)	11053795
Total length (>= 5000 bp)	11008310
Total length (>= 10000 bp)	10714173
Total length (>= 25000 bp)	9741471
Total length (>= 50000 bp)	8225872
# contigs	213
Largest contig	1582399
Total length	11053795
GC (%)	47.05
N50	101654
N90	21339
auN	308481.5
L50	28
L90	115
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	26 + 0 part

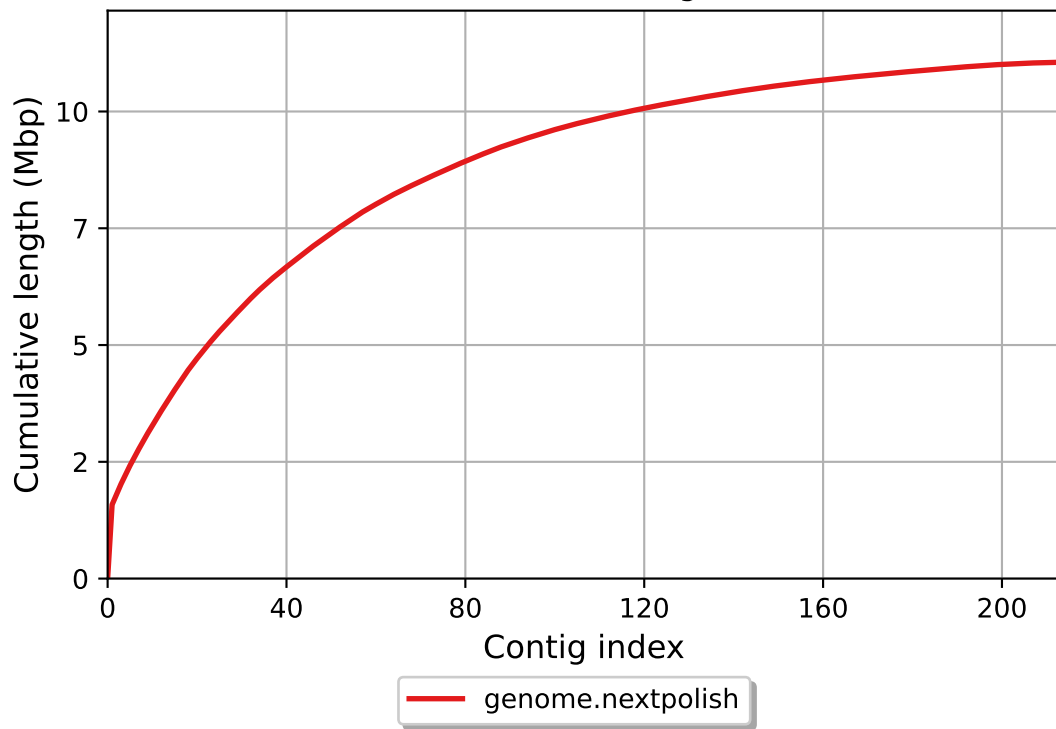
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

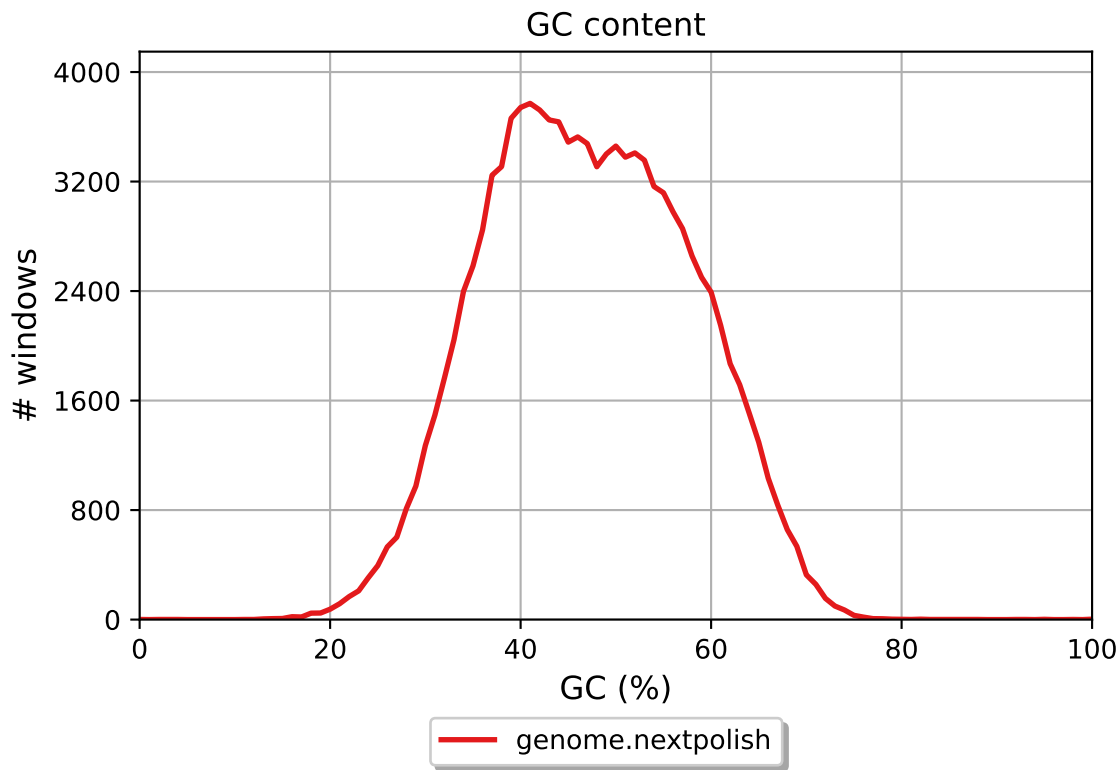
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

