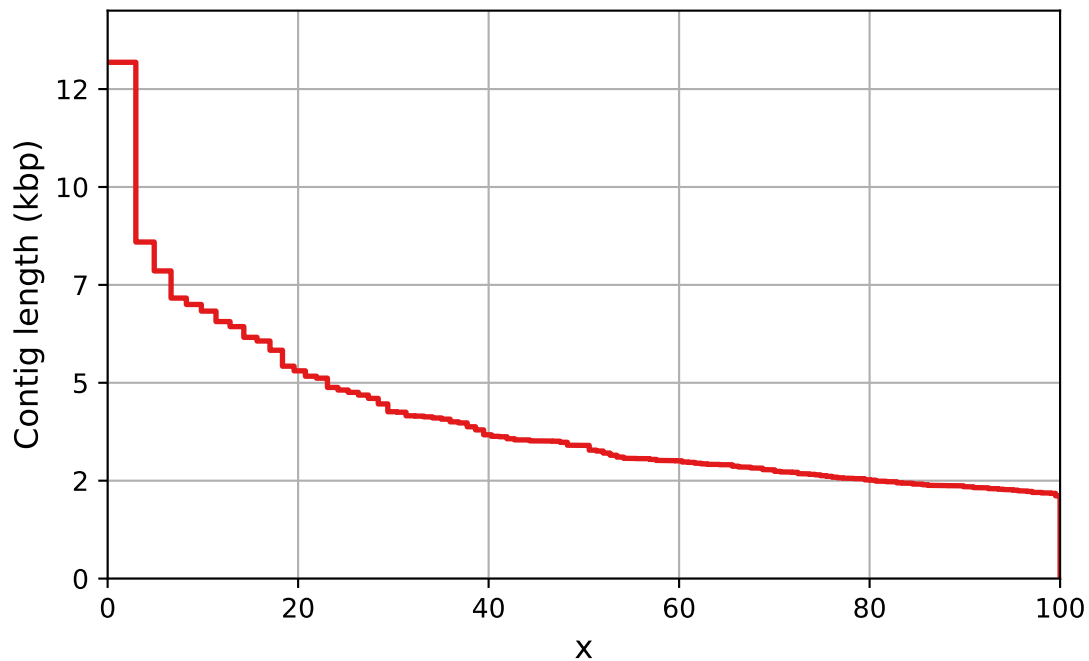


Report

	genome.nextpolish
# contigs (>= 0 bp)	130
# contigs (>= 1000 bp)	130
# contigs (>= 5000 bp)	15
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	444972
Total length (>= 1000 bp)	444972
Total length (>= 5000 bp)	102698
Total length (>= 10000 bp)	13184
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	130
Largest contig	13184
Total length	444972
GC (%)	48.04
N50	3400
N90	2348
auN	4119.4
L50	46
L90	111
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 1 part

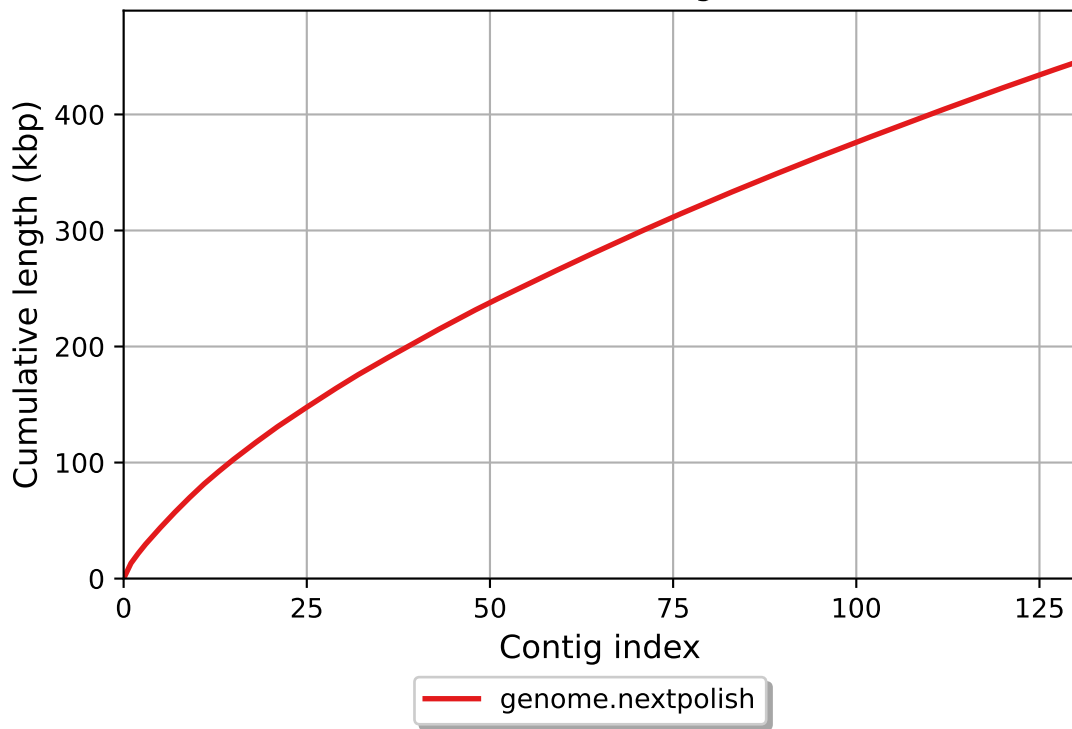
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

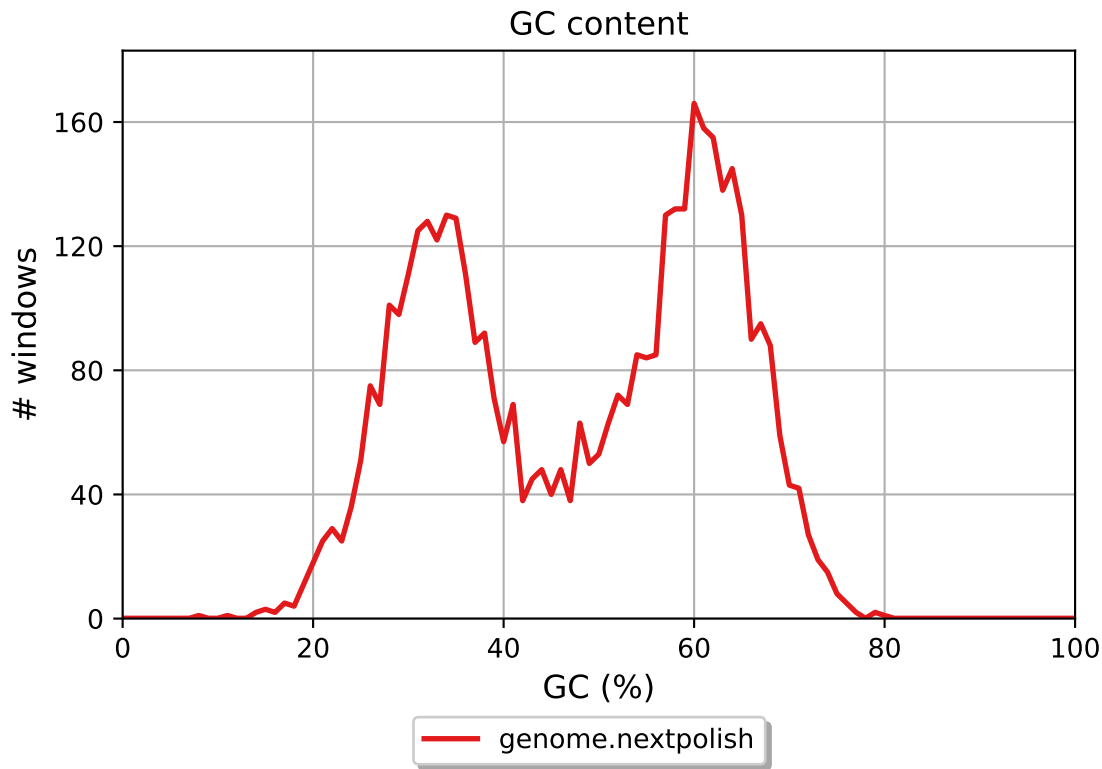
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

