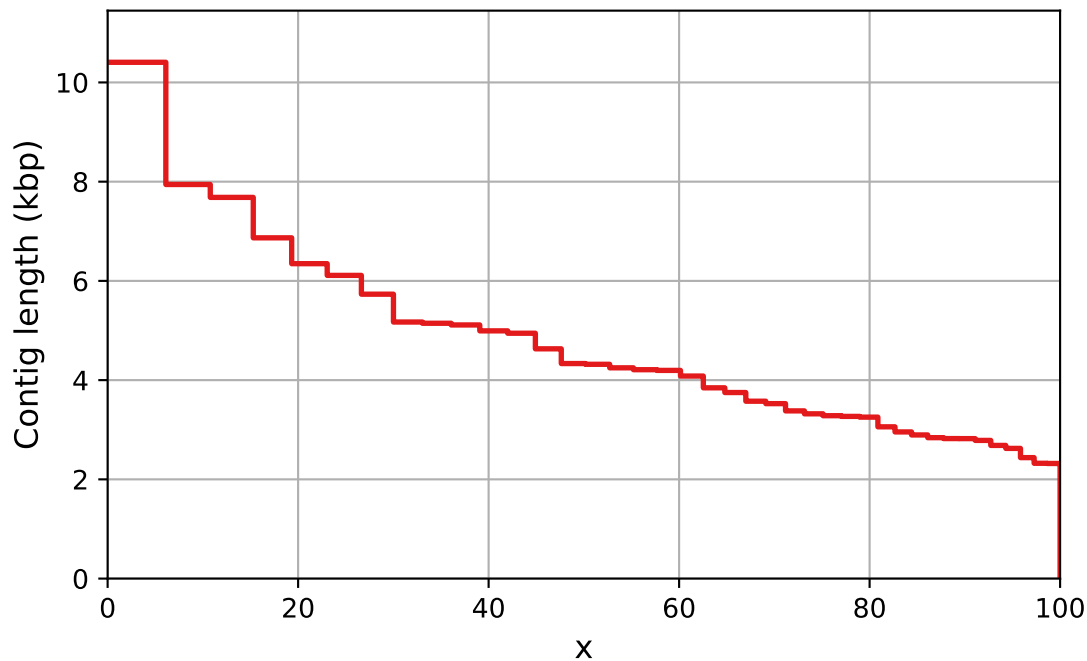


# Report

	genome.nextpolish
# contigs (>= 0 bp)	40
# contigs (>= 1000 bp)	40
# contigs (>= 5000 bp)	10
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	170248
Total length (>= 1000 bp)	170248
Total length (>= 5000 bp)	66519
Total length (>= 10000 bp)	10406
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	40
Largest contig	10406
Total length	170248
GC (%)	43.03
N50	4334
N90	2821
auN	4953.5
L50	14
L90	34
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 0 part

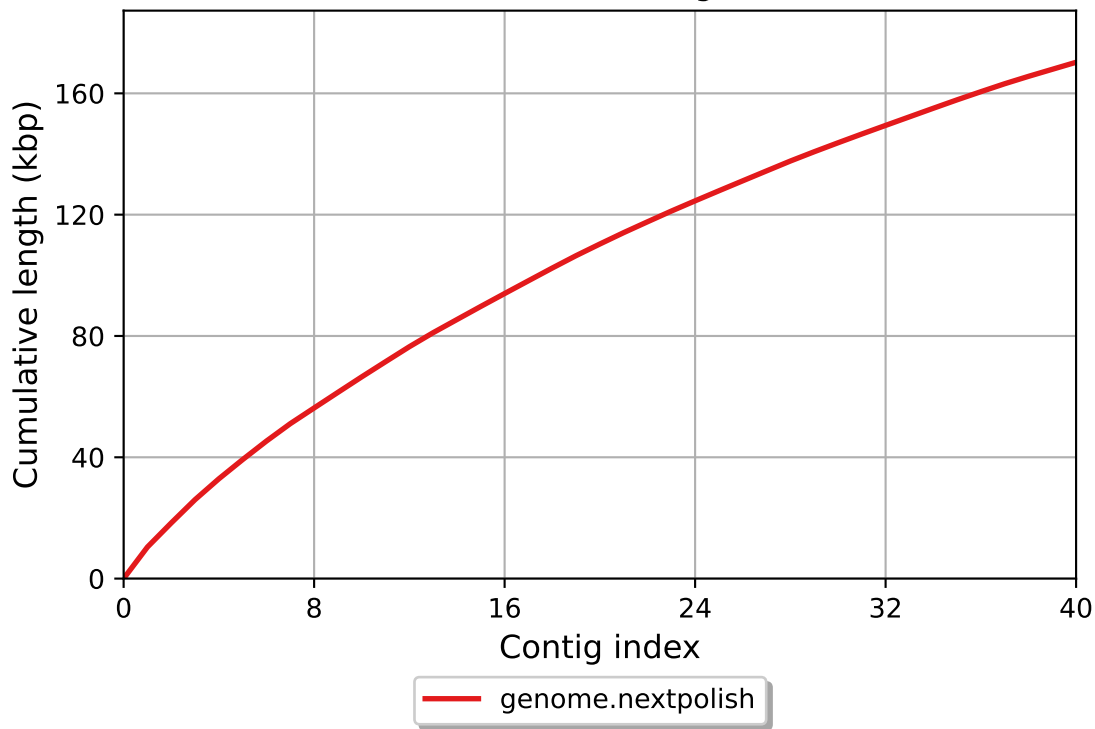
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

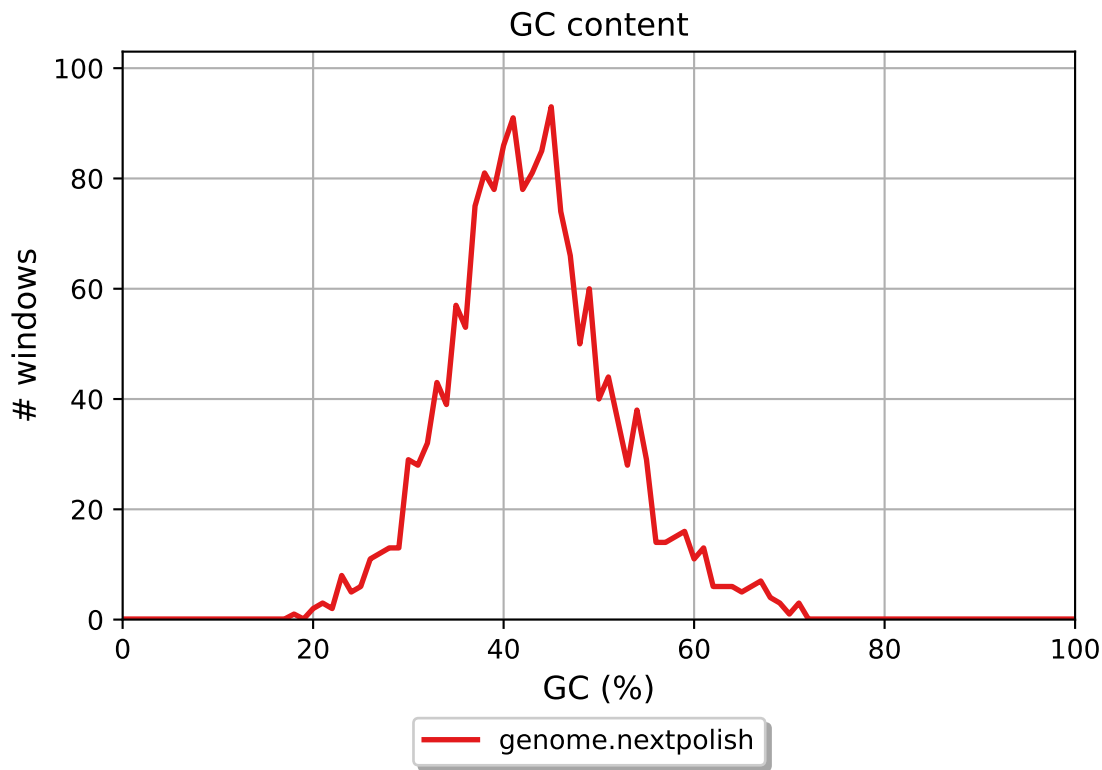
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

