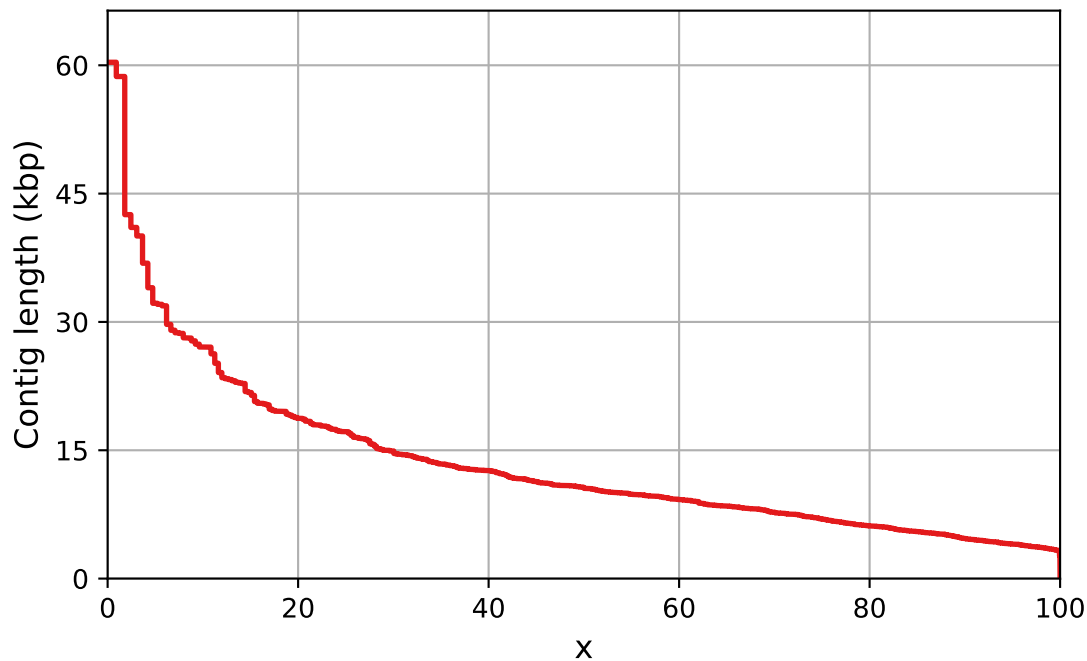


Report

	genome.nextpolish
# contigs (>= 0 bp)	725
# contigs (>= 1000 bp)	725
# contigs (>= 5000 bp)	540
# contigs (>= 10000 bp)	222
# contigs (>= 25000 bp)	23
# contigs (>= 50000 bp)	2
Total length (>= 0 bp)	6620440
Total length (>= 1000 bp)	6620440
Total length (>= 5000 bp)	5872472
Total length (>= 10000 bp)	3577618
Total length (>= 25000 bp)	769858
Total length (>= 50000 bp)	119039
# contigs	725
Largest contig	60351
Total length	6620440
GC (%)	41.08
N50	10665
N90	4689
auN	13634.5
L50	196
L90	558
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	24 + 3 part

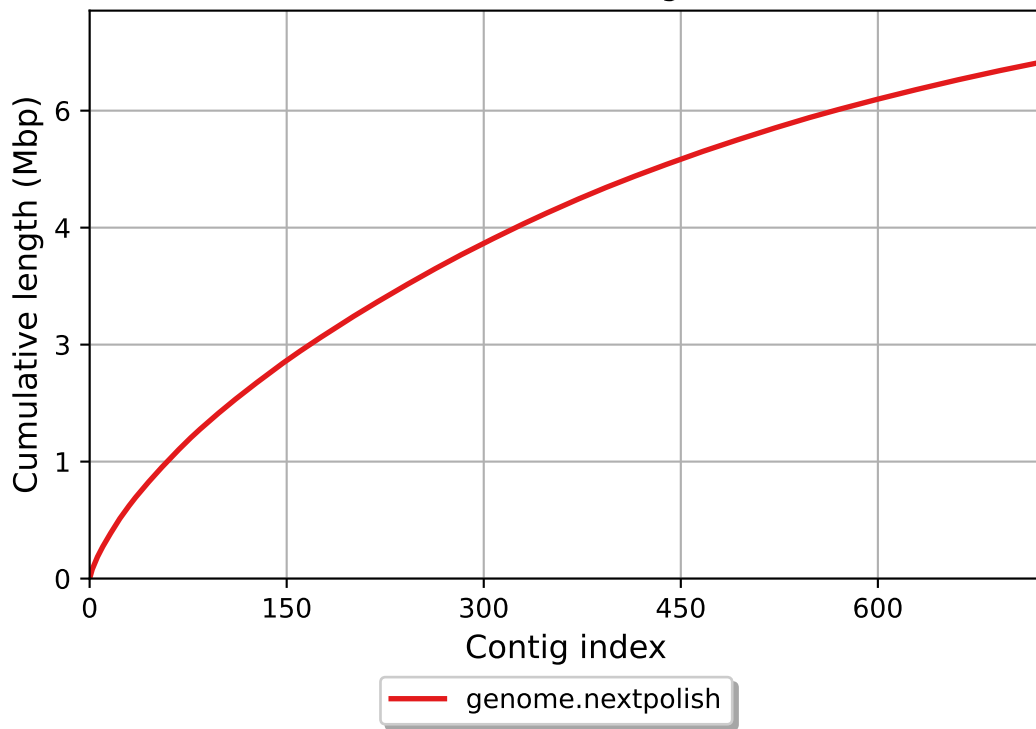
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

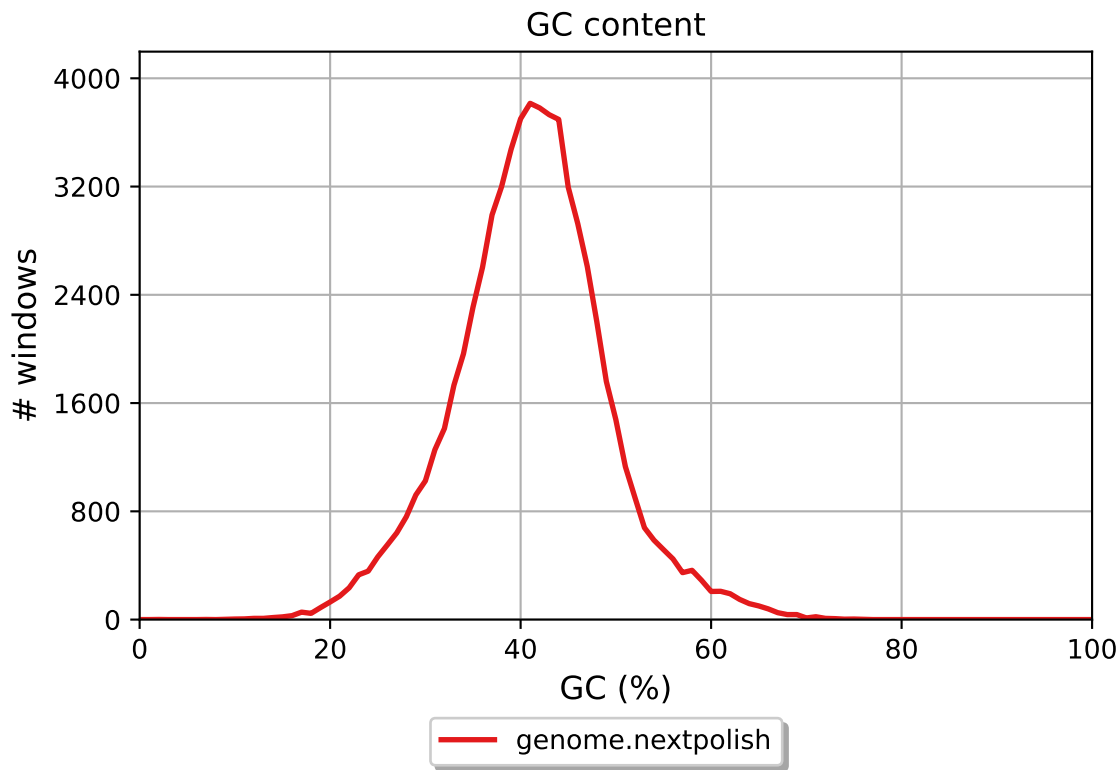
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

