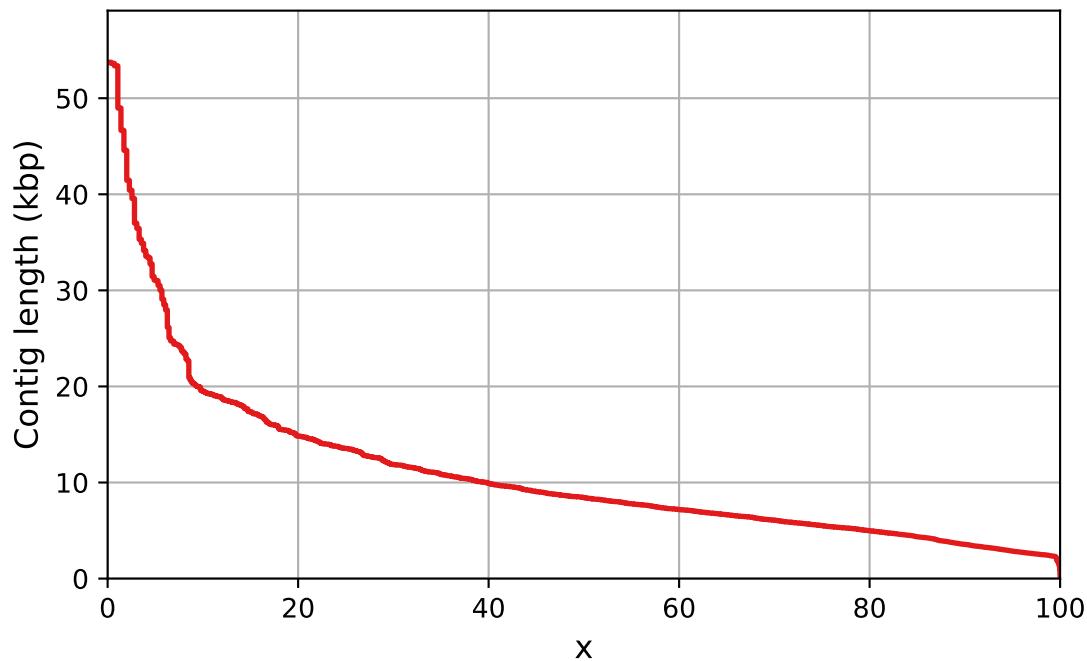


Report

	genome.nextpolish
# contigs (>= 0 bp)	2135
# contigs (>= 1000 bp)	2128
# contigs (>= 5000 bp)	1239
# contigs (>= 10000 bp)	384
# contigs (>= 25000 bp)	27
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	14984347
Total length (>= 1000 bp)	14979798
Total length (>= 5000 bp)	11970909
Total length (>= 10000 bp)	5967874
Total length (>= 25000 bp)	990638
Total length (>= 50000 bp)	160739
# contigs	2135
Largest contig	53741
Total length	14984347
GC (%)	41.88
N50	8427
N90	3556
auN	11100.9
L50	552
L90	1593
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	25 + 5 part

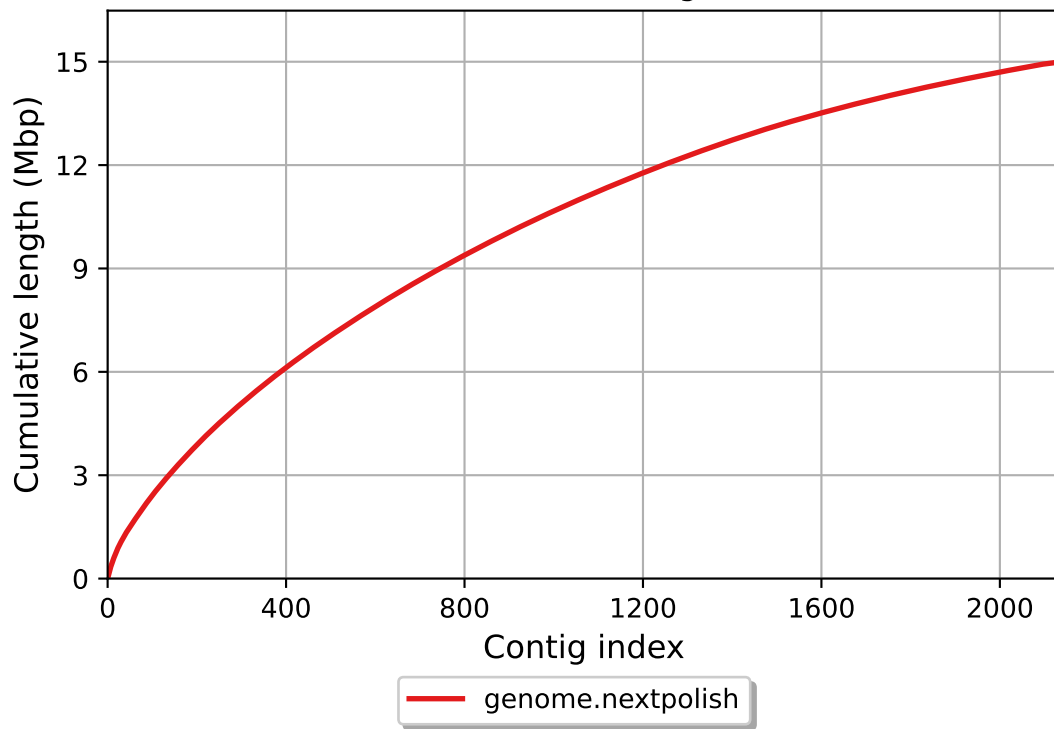
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

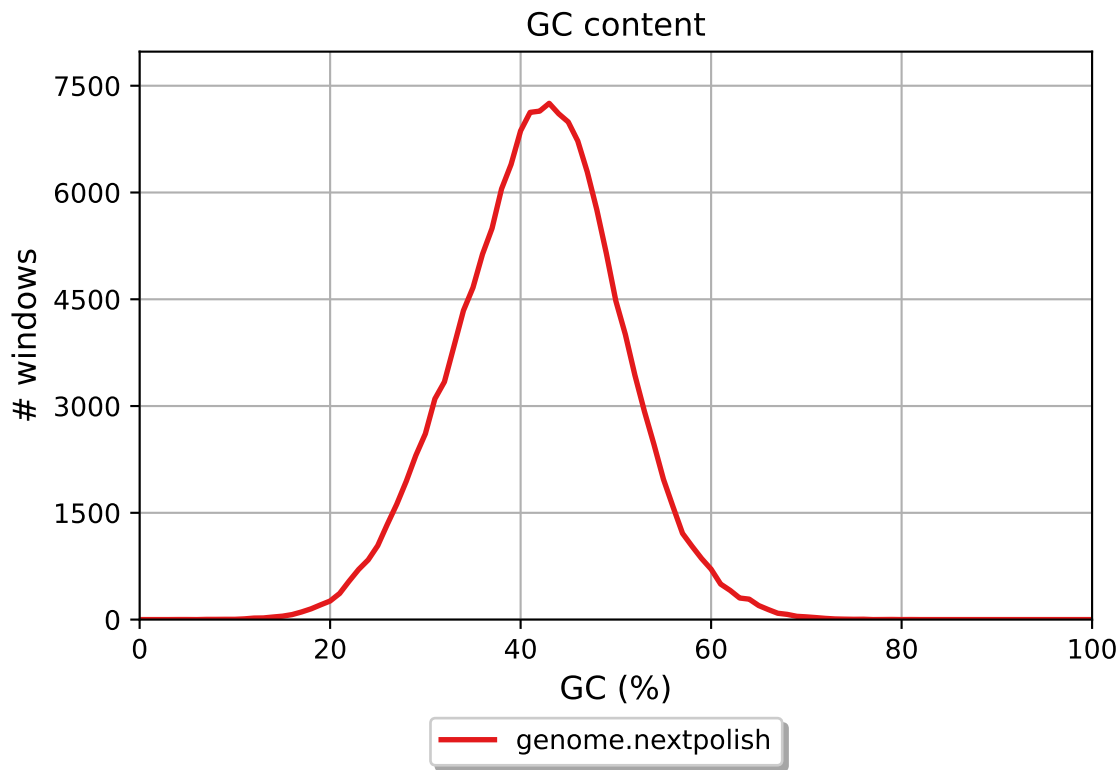
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

