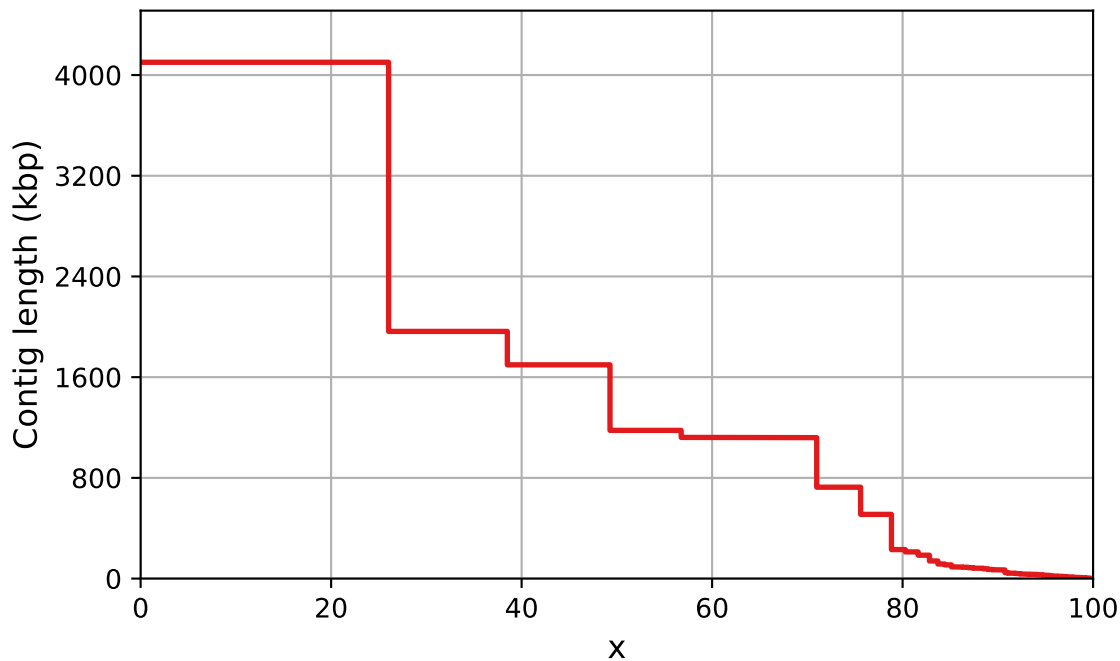


Report

	genome.nextpolish
# contigs (>= 0 bp)	136
# contigs (>= 1000 bp)	130
# contigs (>= 5000 bp)	102
# contigs (>= 10000 bp)	76
# contigs (>= 25000 bp)	46
# contigs (>= 50000 bp)	25
Total length (>= 0 bp)	15752895
Total length (>= 1000 bp)	15748432
Total length (>= 5000 bp)	15672833
Total length (>= 10000 bp)	15486420
Total length (>= 25000 bp)	15025794
Total length (>= 50000 bp)	14294970
# contigs	136
Largest contig	4101416
Total length	15752895
GC (%)	47.04
N50	1177388
N90	69378
auN	1811108.2
L50	4
L90	24
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	79 + 0 part

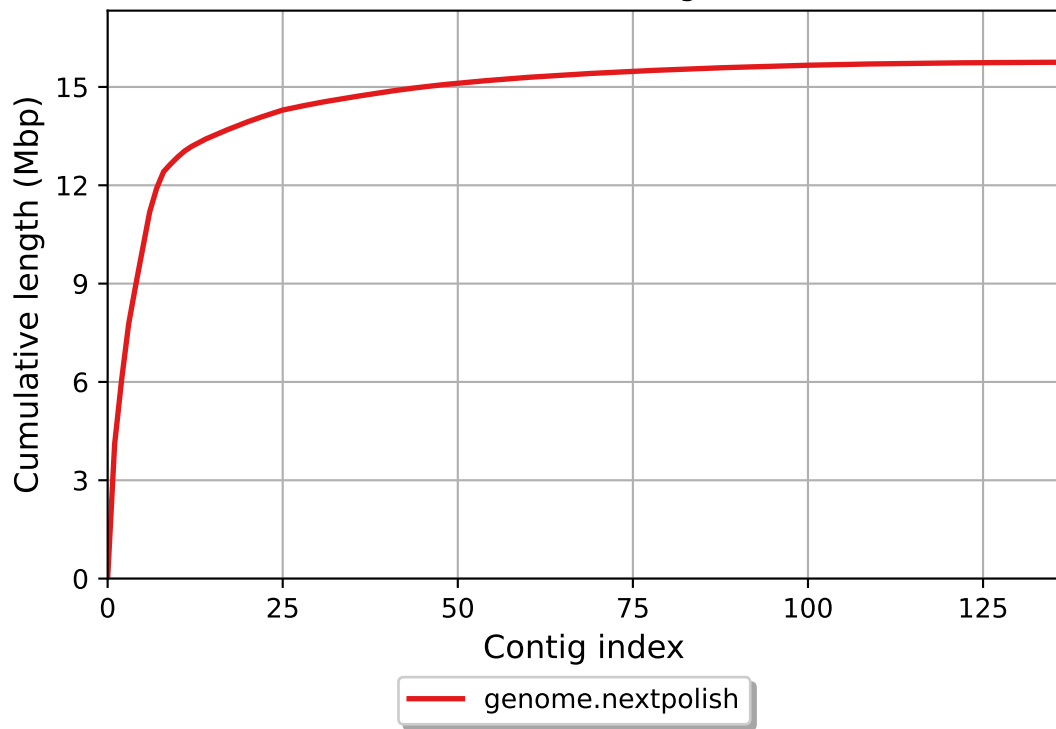
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

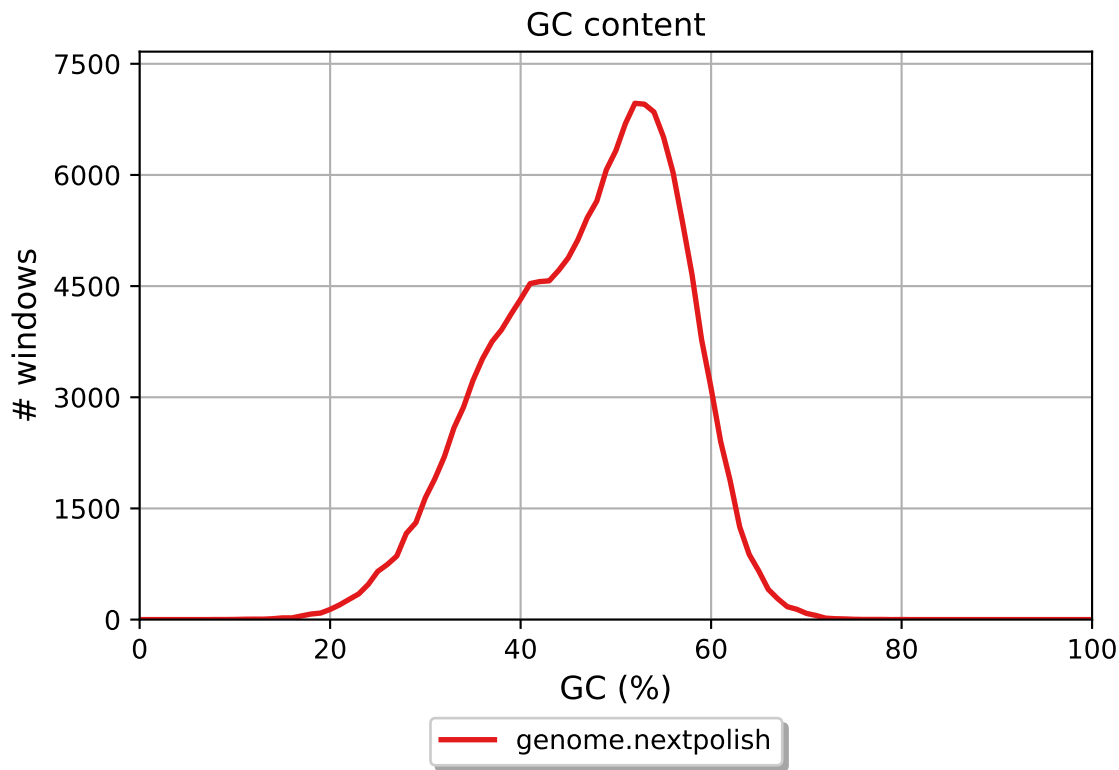
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

