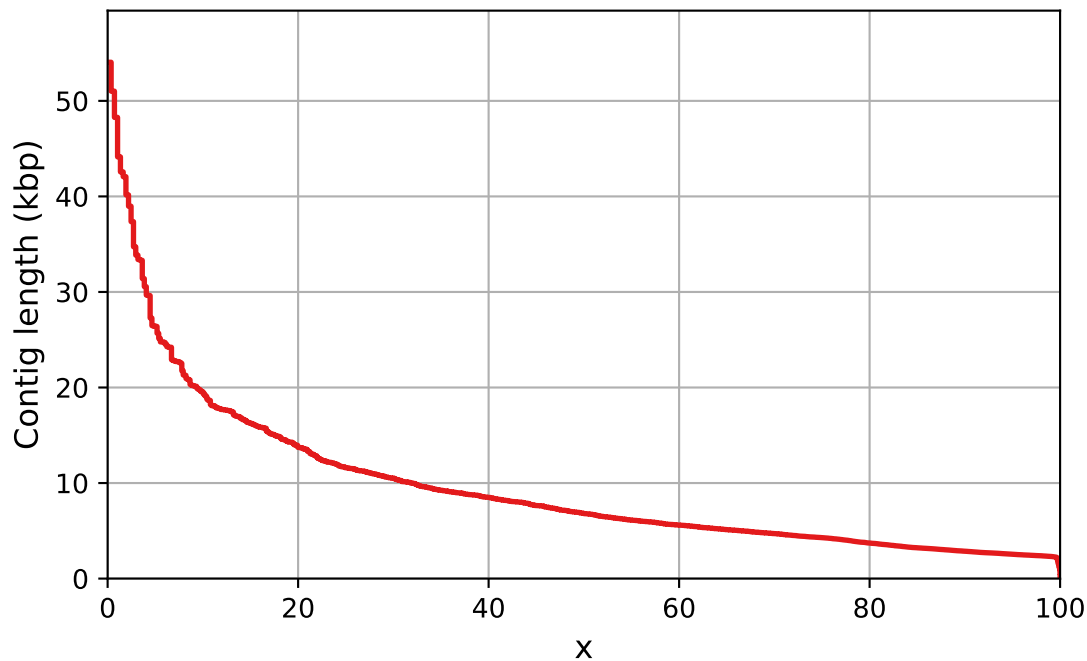


Report

	genome.nextpolish
# contigs (>= 0 bp)	2538
# contigs (>= 1000 bp)	2528
# contigs (>= 5000 bp)	1041
# contigs (>= 10000 bp)	296
# contigs (>= 25000 bp)	23
# contigs (>= 50000 bp)	2
Total length (>= 0 bp)	14677178
Total length (>= 1000 bp)	14670230
Total length (>= 5000 bp)	9776400
Total length (>= 10000 bp)	4685952
Total length (>= 25000 bp)	812516
Total length (>= 50000 bp)	105056
# contigs	2538
Largest contig	54040
Total length	14677178
GC (%)	43.26
N50	6805
N90	2861
auN	9698.5
L50	619
L90	1945
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	23 + 6 part

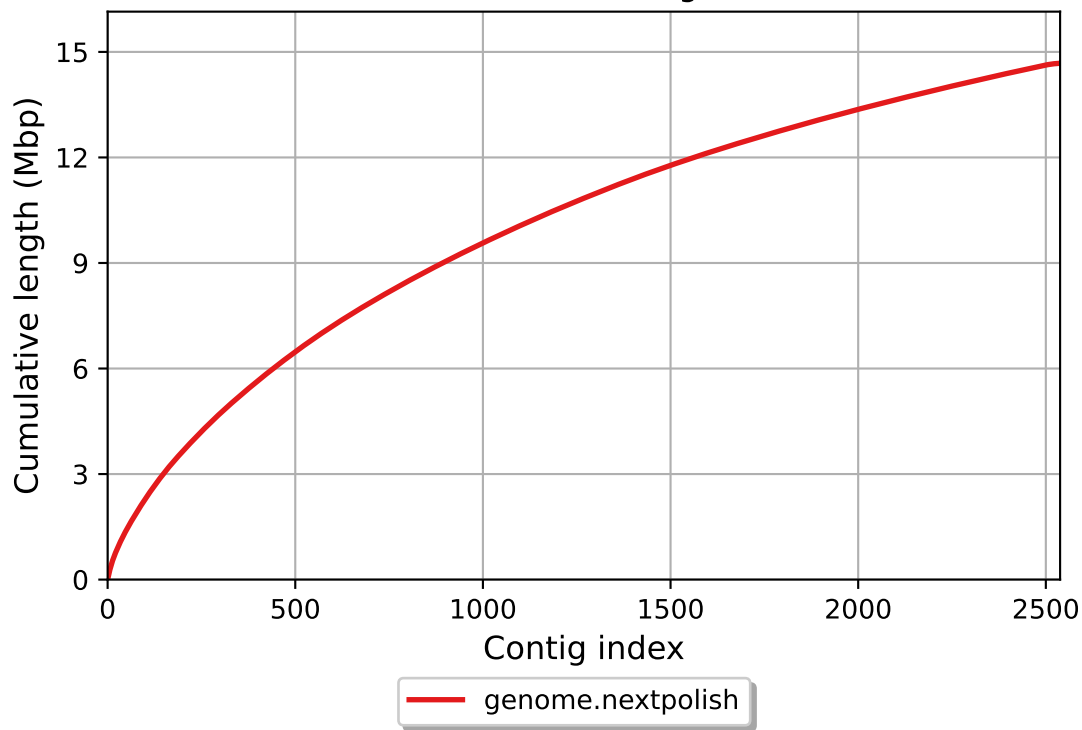
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

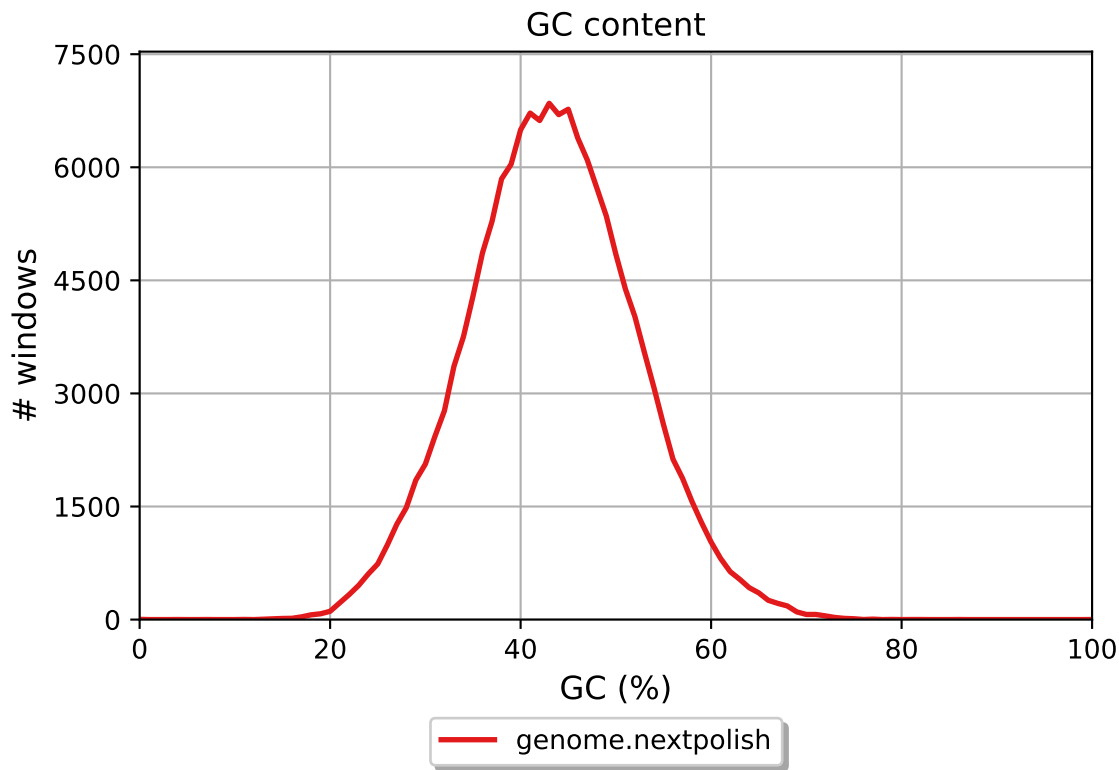
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

