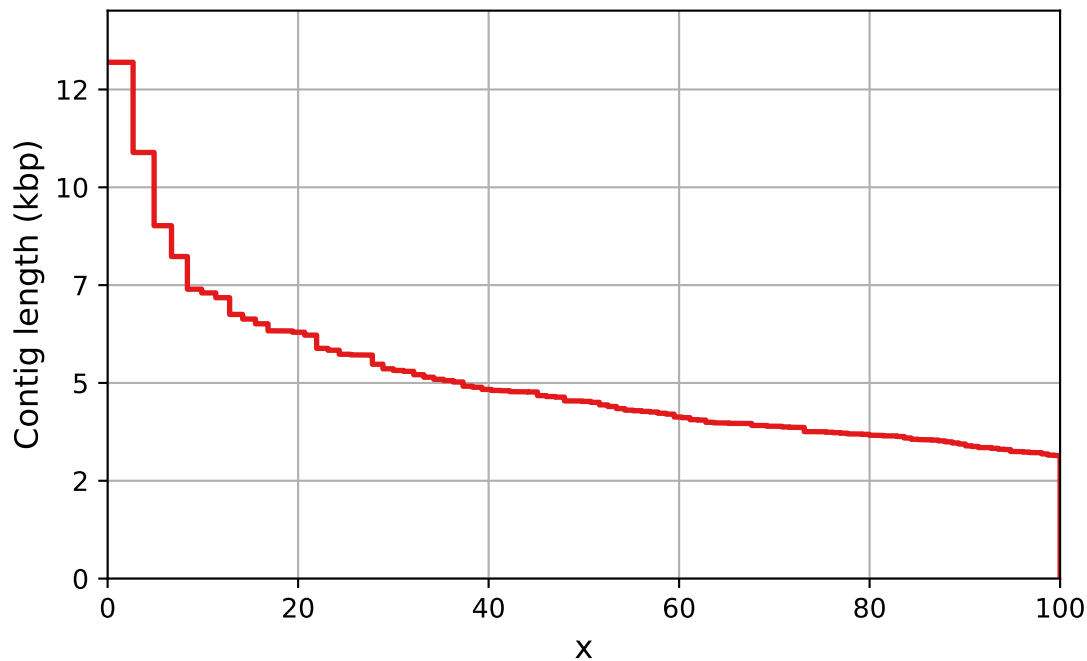


Report

	genome.nextpolish
# contigs (>= 0 bp)	107
# contigs (>= 1000 bp)	107
# contigs (>= 5000 bp)	28
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	493488
Total length (>= 1000 bp)	493488
Total length (>= 5000 bp)	184158
Total length (>= 10000 bp)	24085
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	107
Largest contig	13195
Total length	493488
GC (%)	49.32
N50	4528
N90	3439
auN	5137.1
L50	42
L90	92
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 1 part

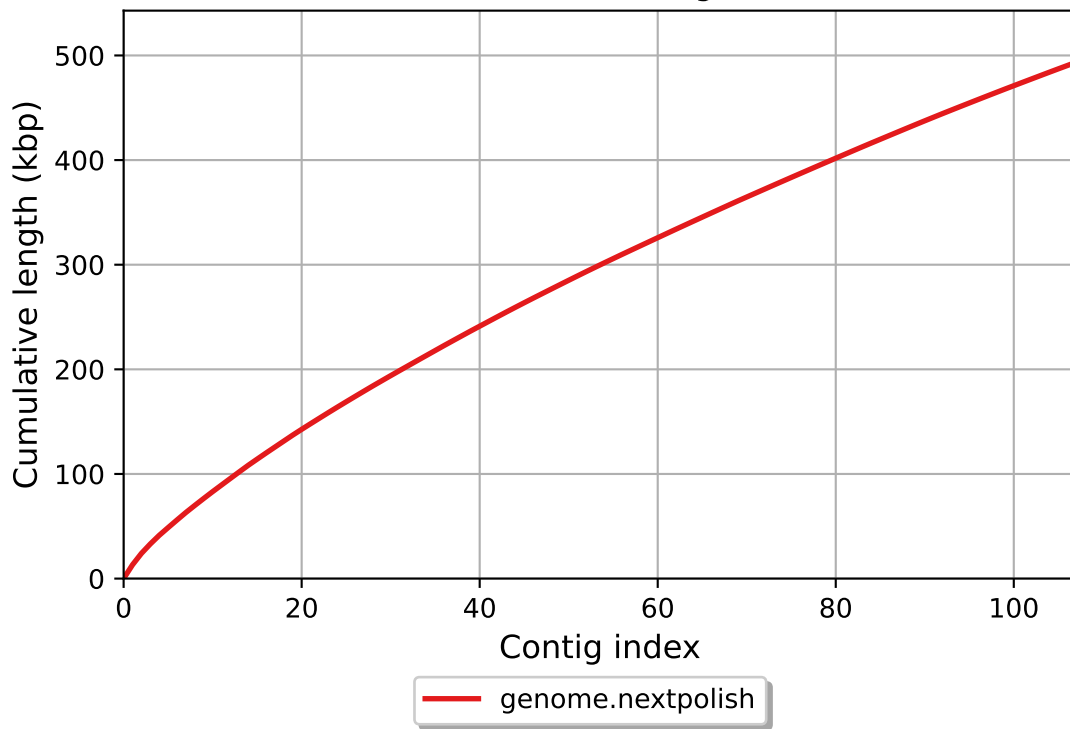
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

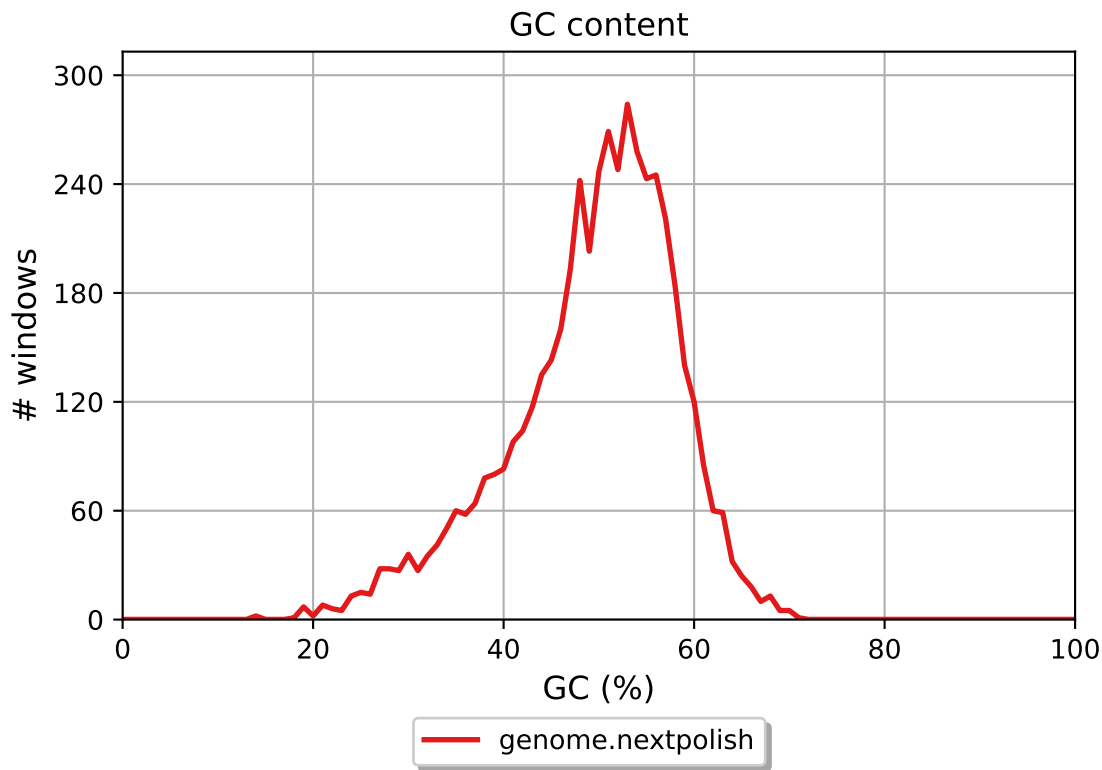
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

