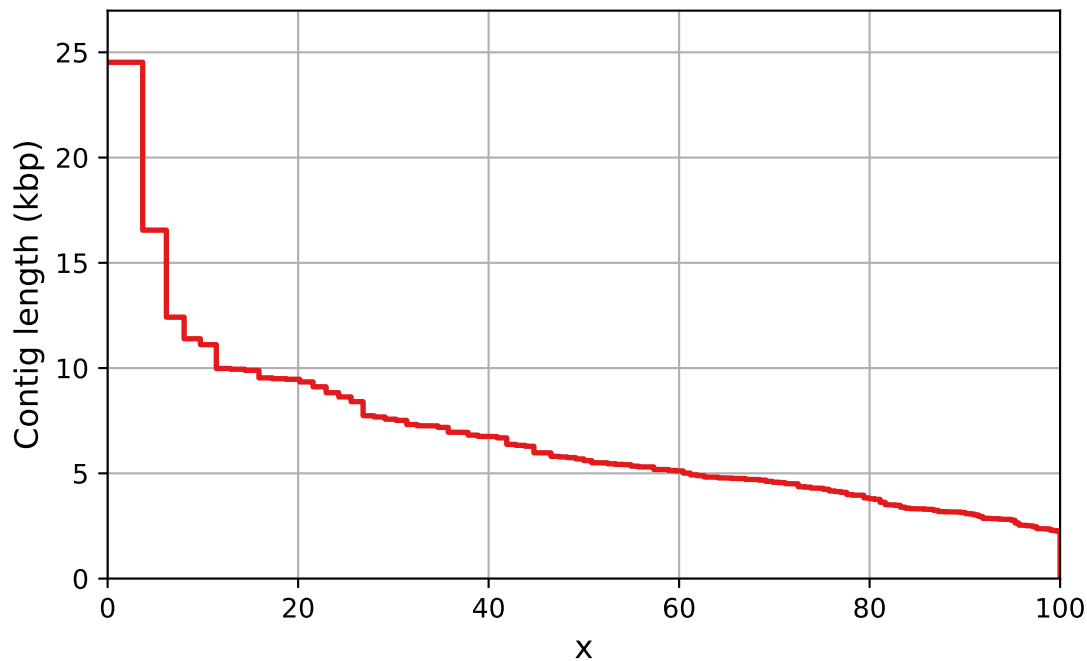


Report

	genome.nextpolish
# contigs (>= 0 bp)	126
# contigs (>= 1000 bp)	126
# contigs (>= 5000 bp)	53
# contigs (>= 10000 bp)	5
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	665965
Total length (>= 1000 bp)	665965
Total length (>= 5000 bp)	407469
Total length (>= 10000 bp)	76000
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	126
Largest contig	24526
Total length	665965
GC (%)	40.38
N50	5684
N90	3142
auN	6983.8
L50	39
L90	101
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 1 part

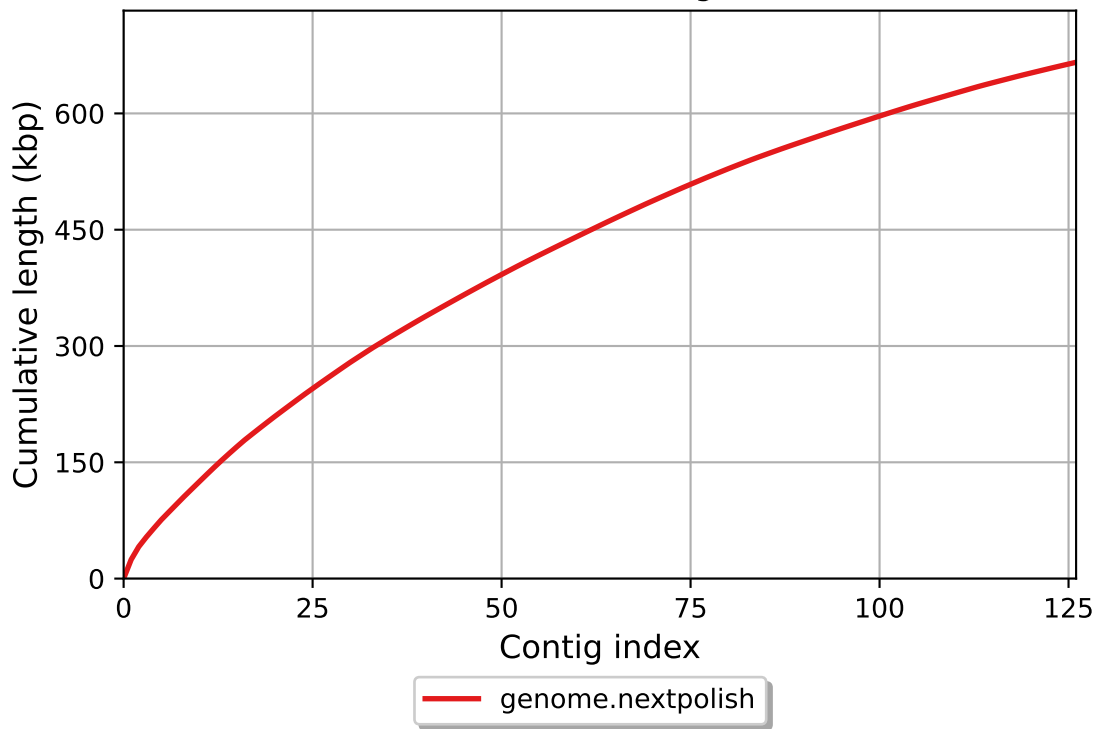
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

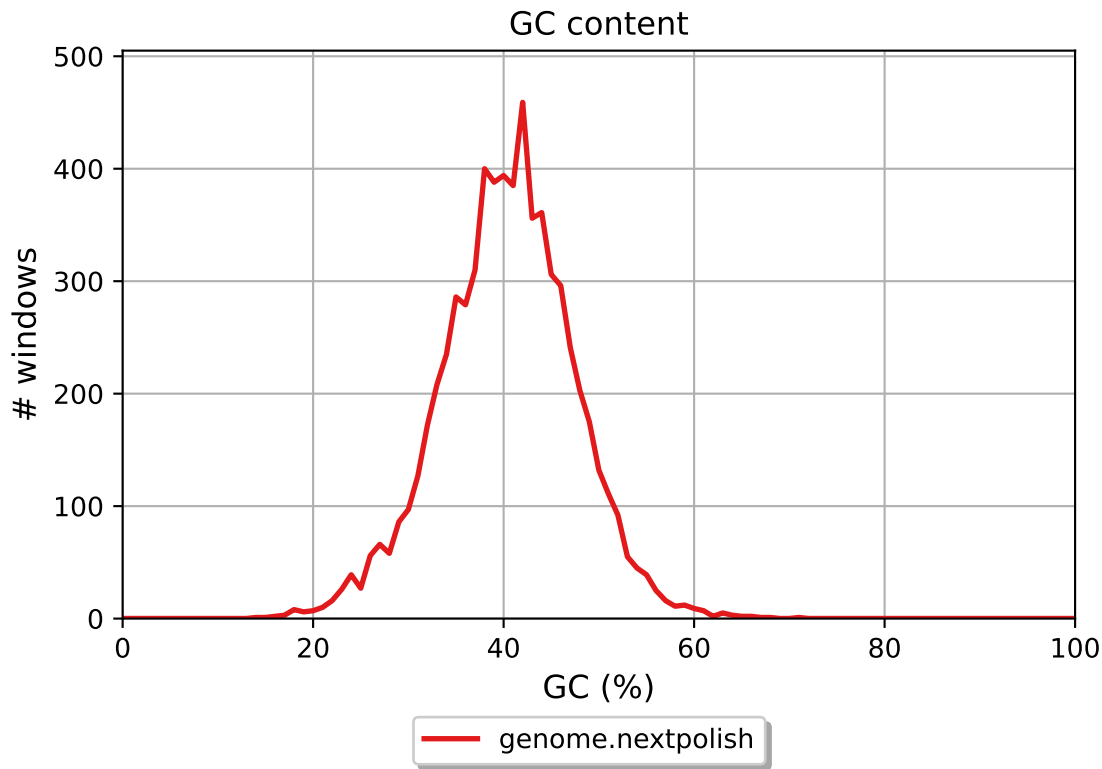
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

