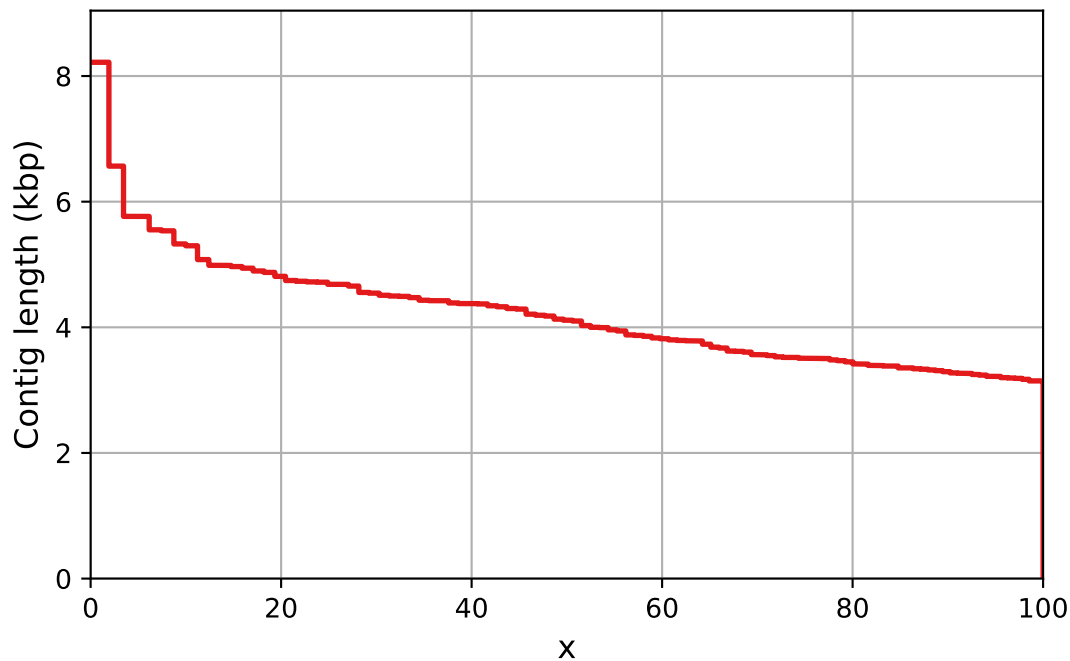


Report

	genome.nextpolish
# contigs (>= 0 bp)	105
# contigs (>= 1000 bp)	105
# contigs (>= 5000 bp)	9
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	427965
Total length (>= 1000 bp)	427965
Total length (>= 5000 bp)	53107
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	105
Largest contig	8219
Total length	427965
GC (%)	36.55
N50	4113
N90	3294
auN	4238.6
L50	45
L90	92
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 1 part

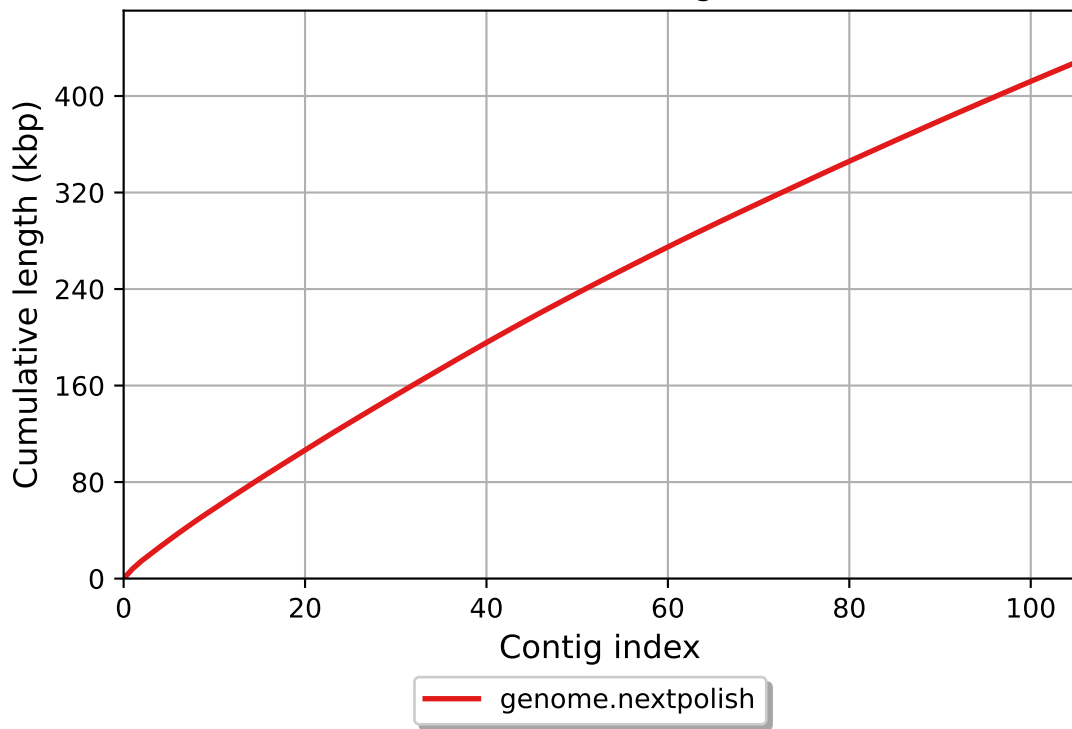
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

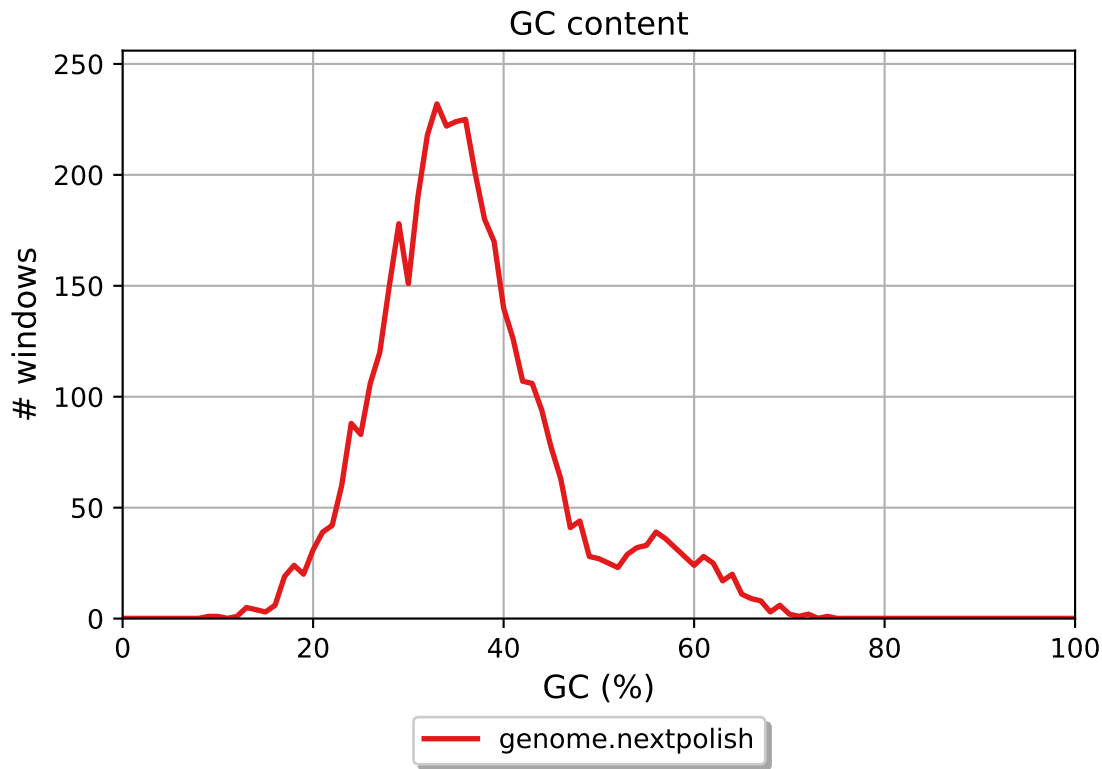
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

