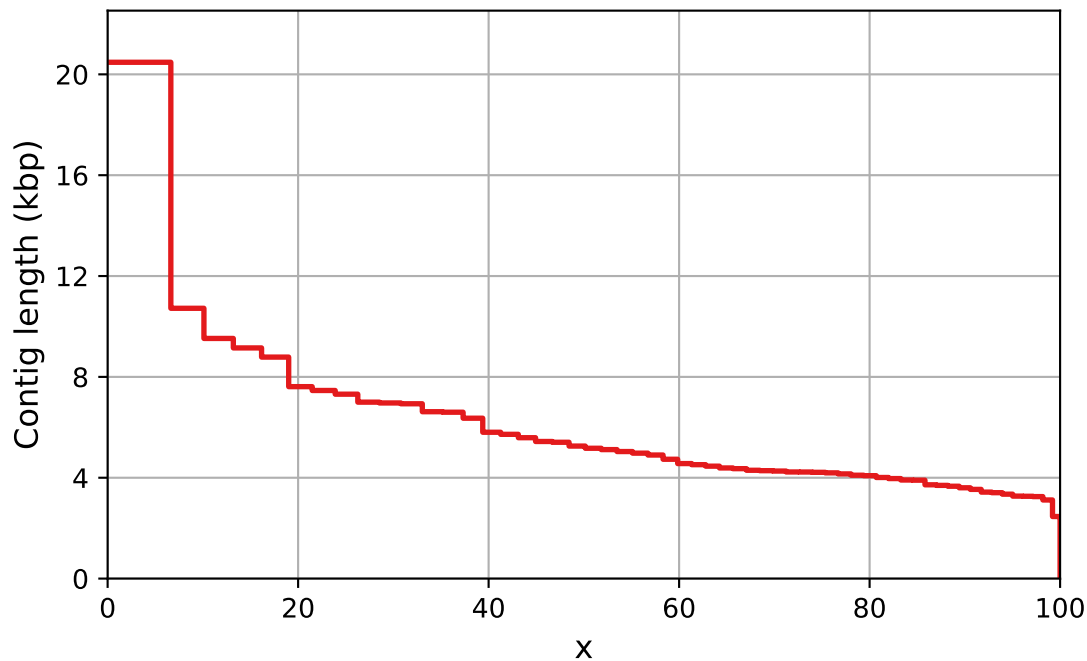


Report

	genome.nextpolish
# contigs (>= 0 bp)	58
# contigs (>= 1000 bp)	58
# contigs (>= 5000 bp)	23
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	308562
Total length (>= 1000 bp)	308562
Total length (>= 5000 bp)	170063
Total length (>= 10000 bp)	31200
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	58
Largest contig	20479
Total length	308562
GC (%)	34.57
N50	5256
N90	3602
auN	6634.7
L50	20
L90	49
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

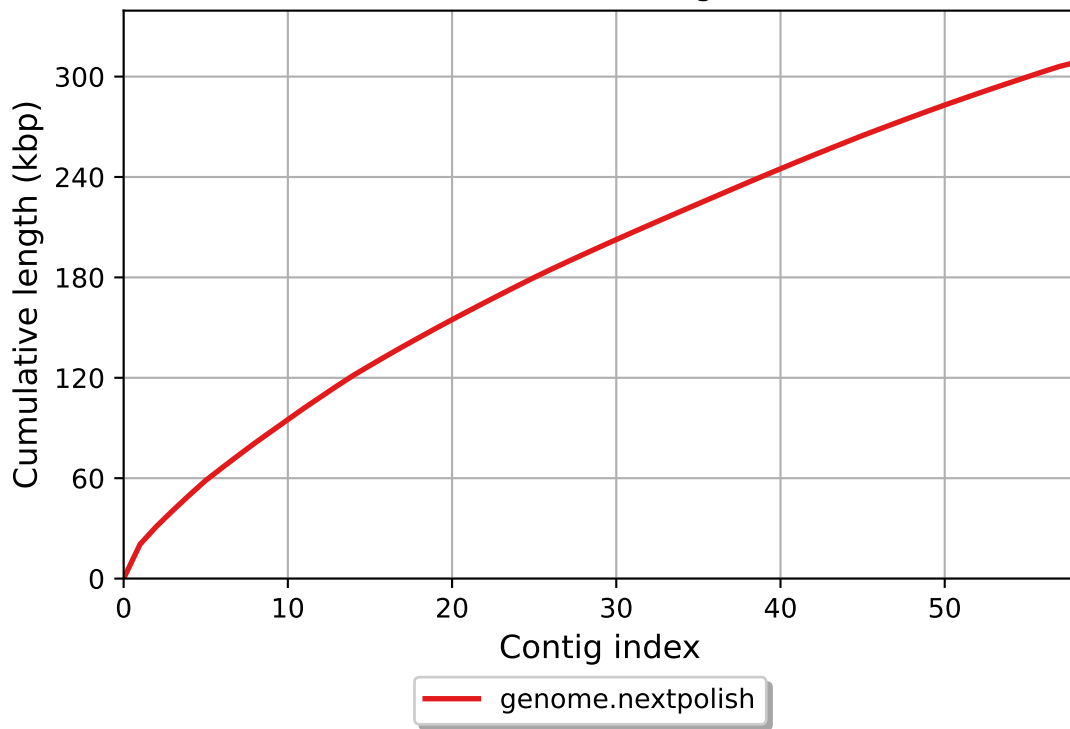
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

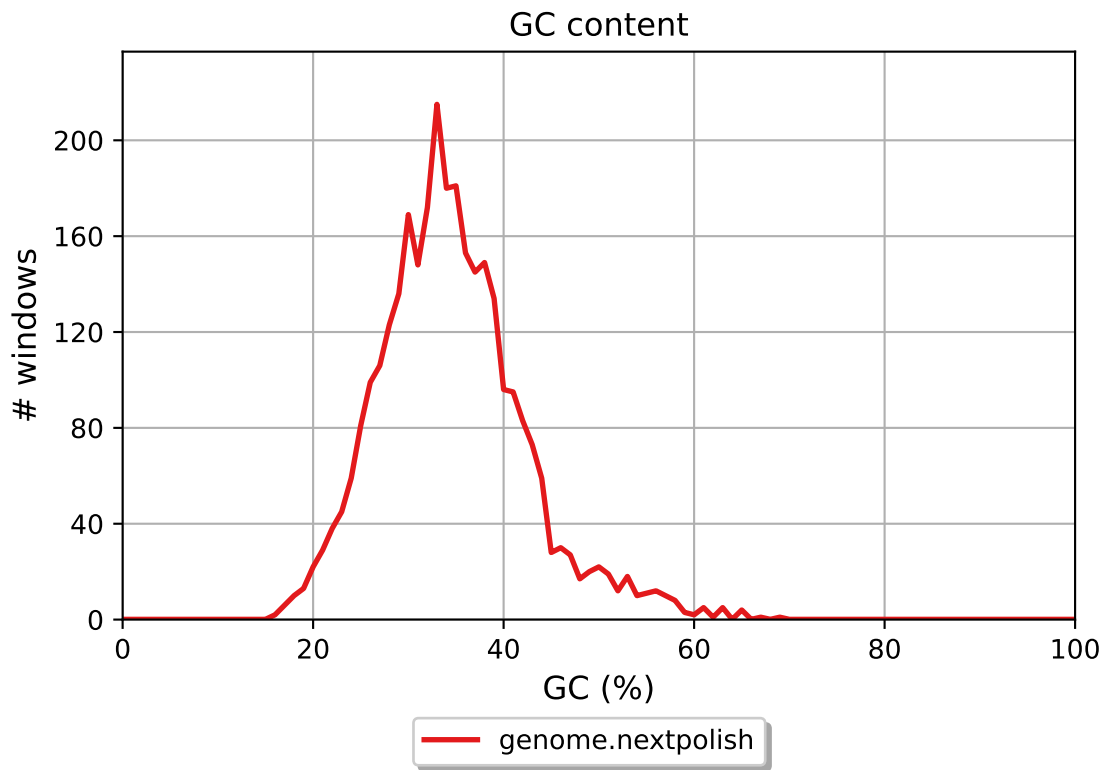
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

