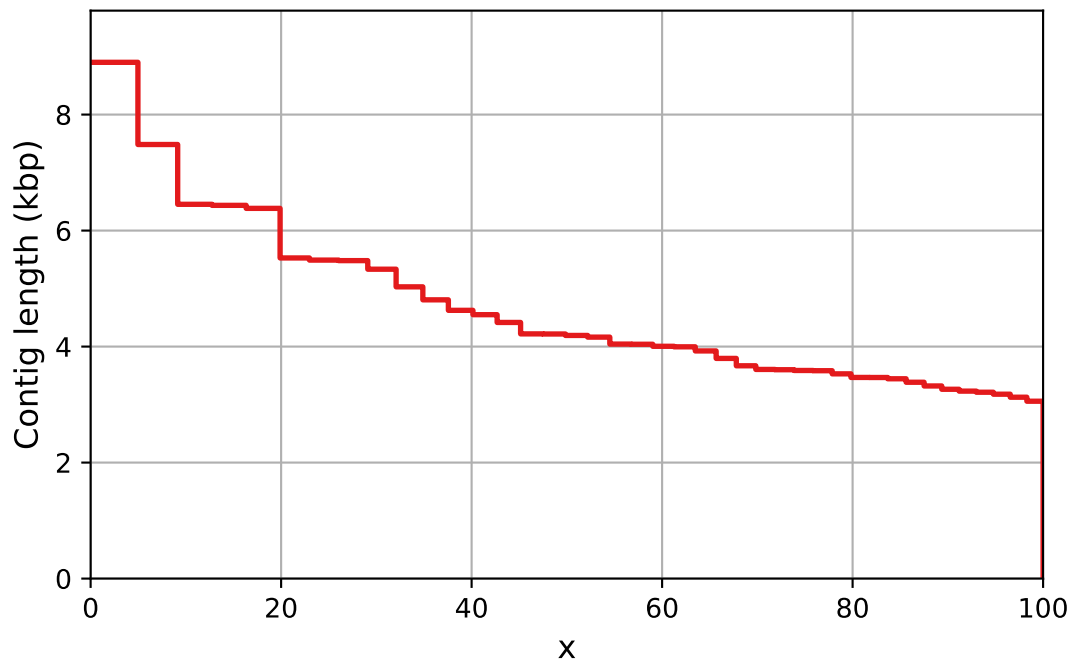


Report

	genome.nextpolish
# contigs (>= 0 bp)	41
# contigs (>= 1000 bp)	41
# contigs (>= 5000 bp)	10
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	179251
Total length (>= 1000 bp)	179251
Total length (>= 5000 bp)	62521
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	41
Largest contig	8901
Total length	179251
GC (%)	50.44
N50	4192
N90	3264
auN	4741.3
L50	17
L90	36
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

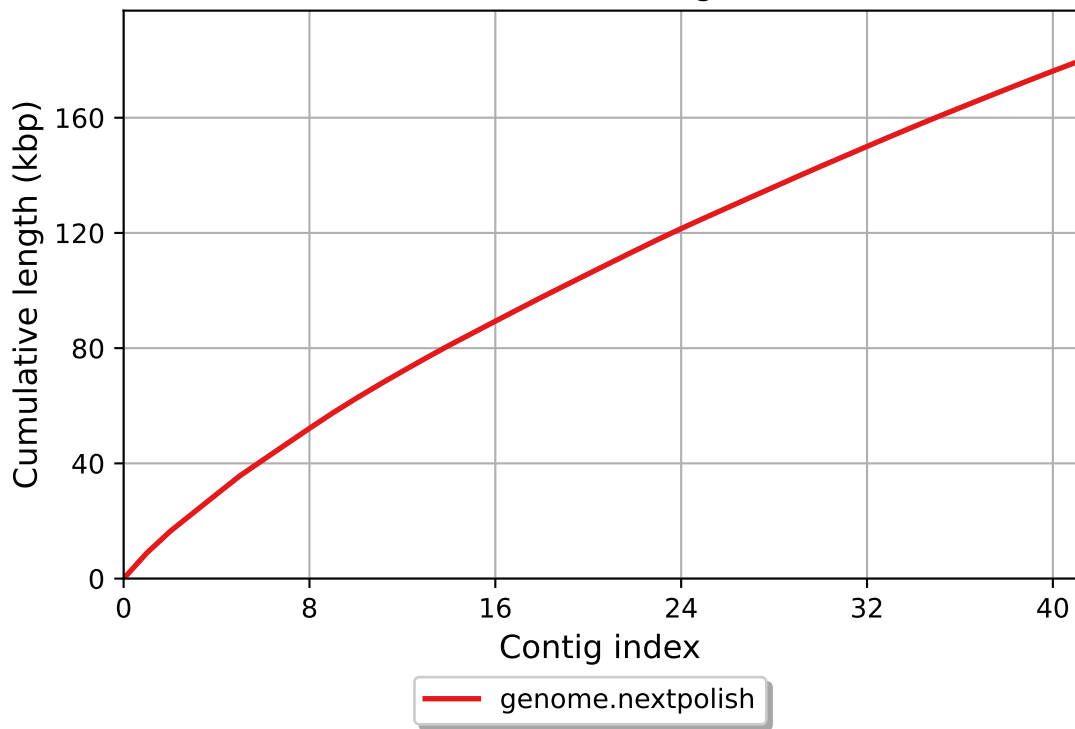
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

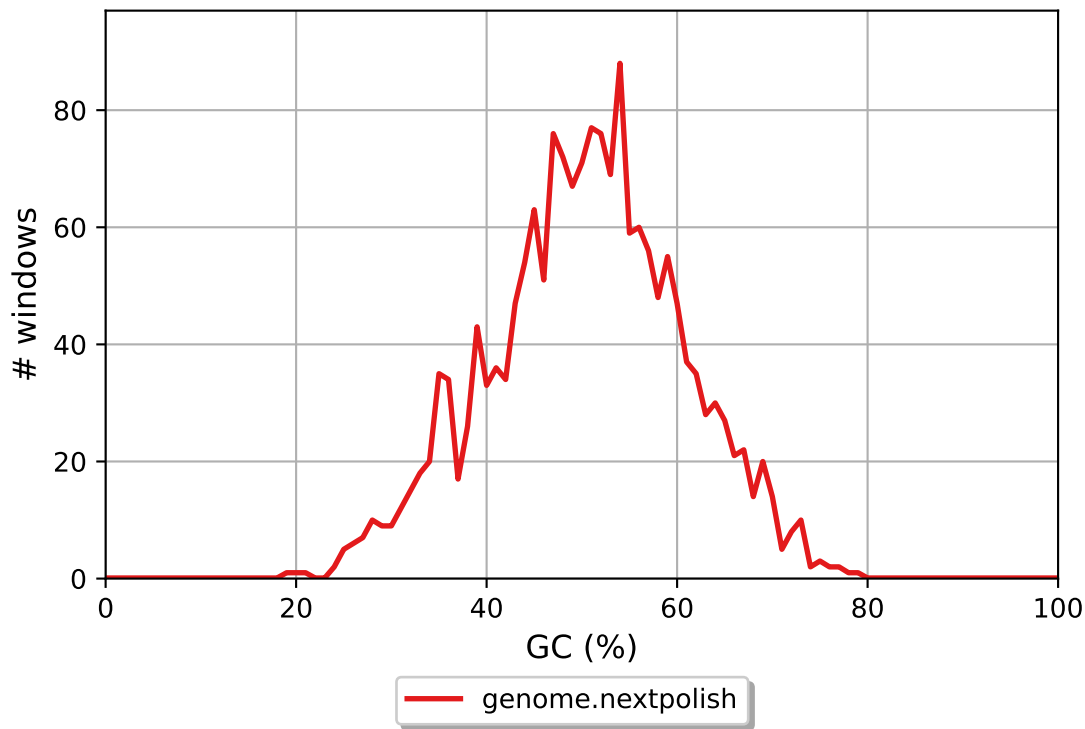


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

