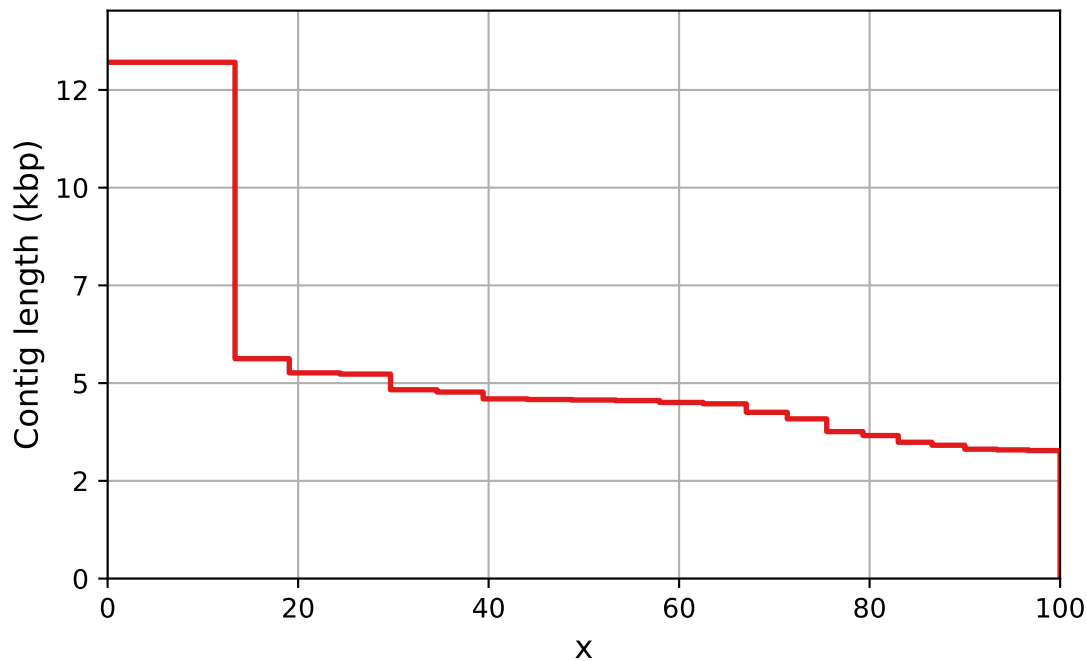


# Report

	genome.nextpolish
# contigs (>= 0 bp)	21
# contigs (>= 1000 bp)	21
# contigs (>= 5000 bp)	4
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	98735
Total length (>= 1000 bp)	98735
Total length (>= 5000 bp)	29325
Total length (>= 10000 bp)	13207
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	21
Largest contig	13207
Total length	98735
GC (%)	49.43
N50	4568
N90	3310
auN	5568.8
L50	9
L90	19
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

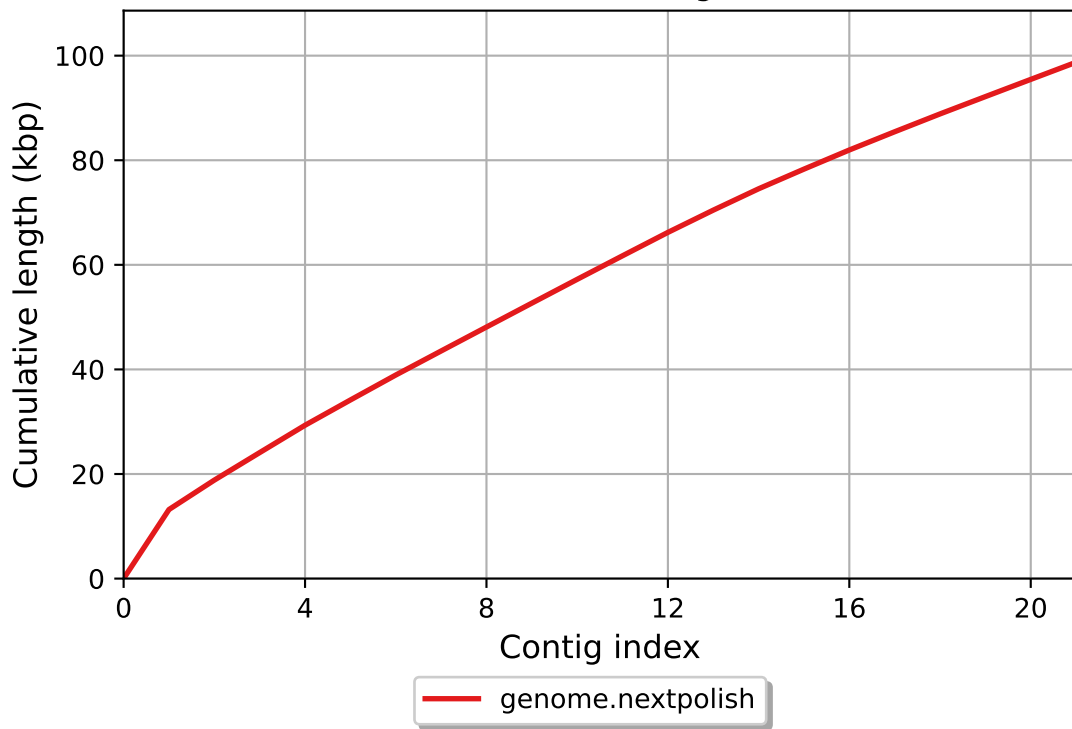
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

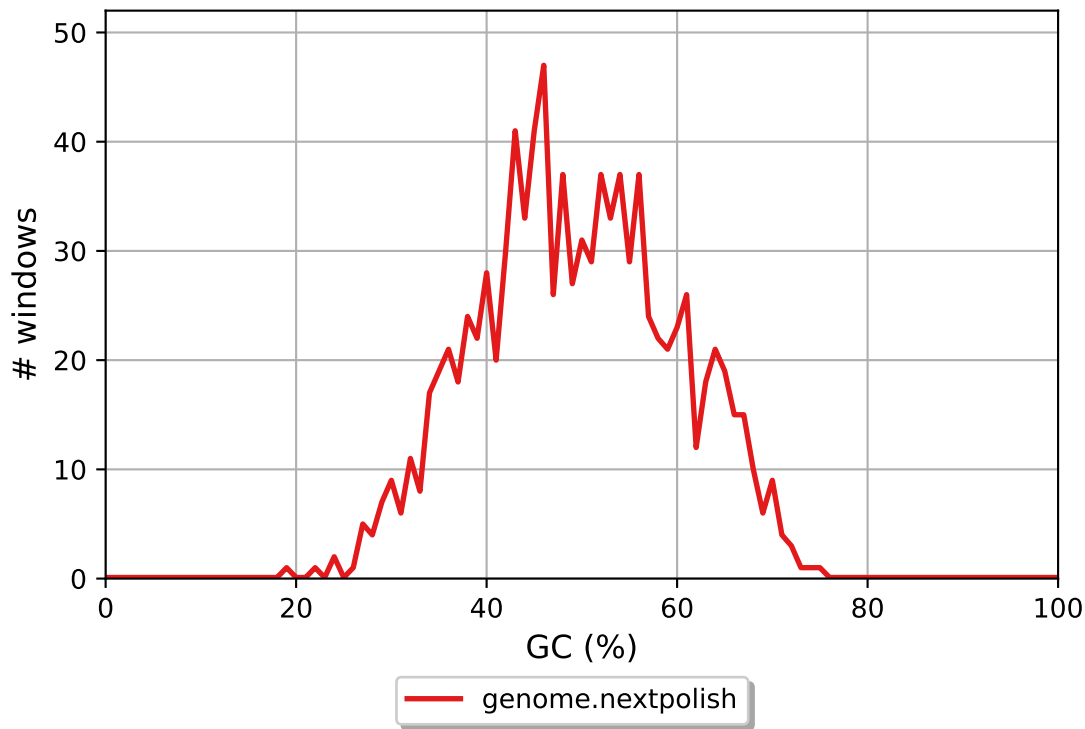


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

