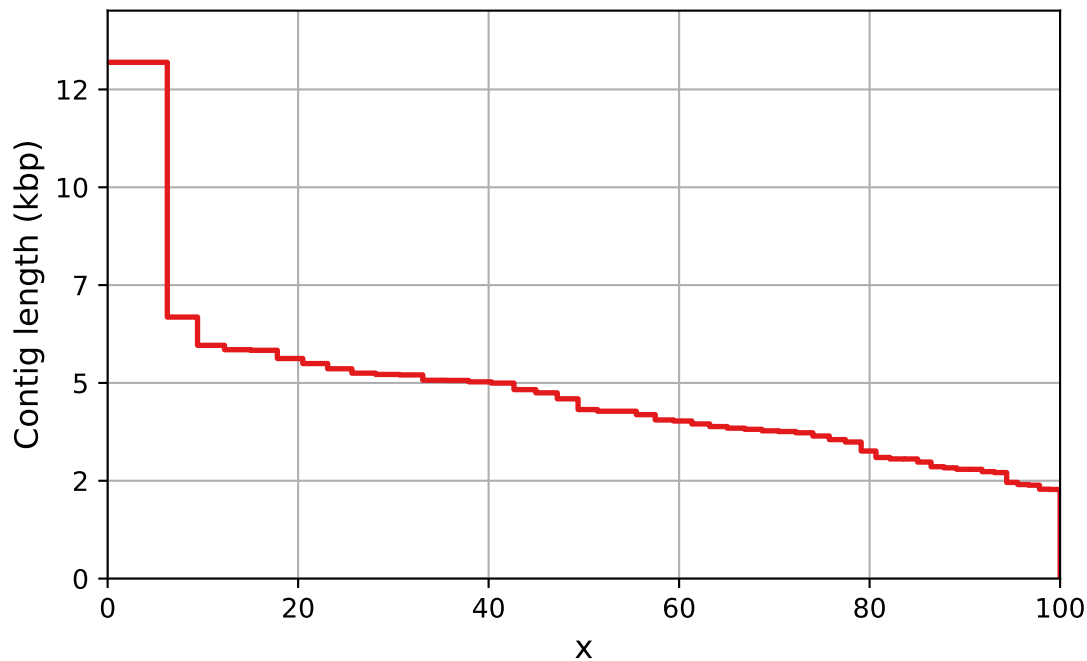


Report

	genome.nextpolish
# contigs (>= 0 bp)	50
# contigs (>= 1000 bp)	50
# contigs (>= 5000 bp)	14
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	210560
Total length (>= 1000 bp)	210560
Total length (>= 5000 bp)	84834
Total length (>= 10000 bp)	13194
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	50
Largest contig	13194
Total length	210560
GC (%)	51.06
N50	4319
N90	2793
auN	4899.3
L50	19
L90	42
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

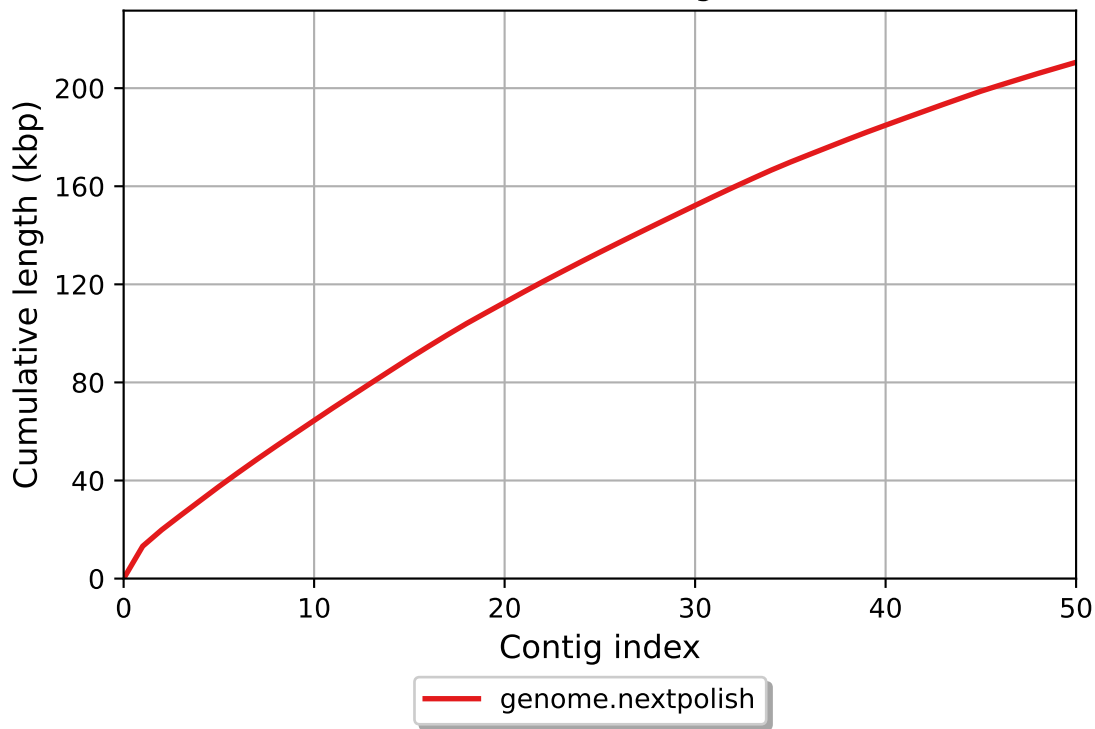
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

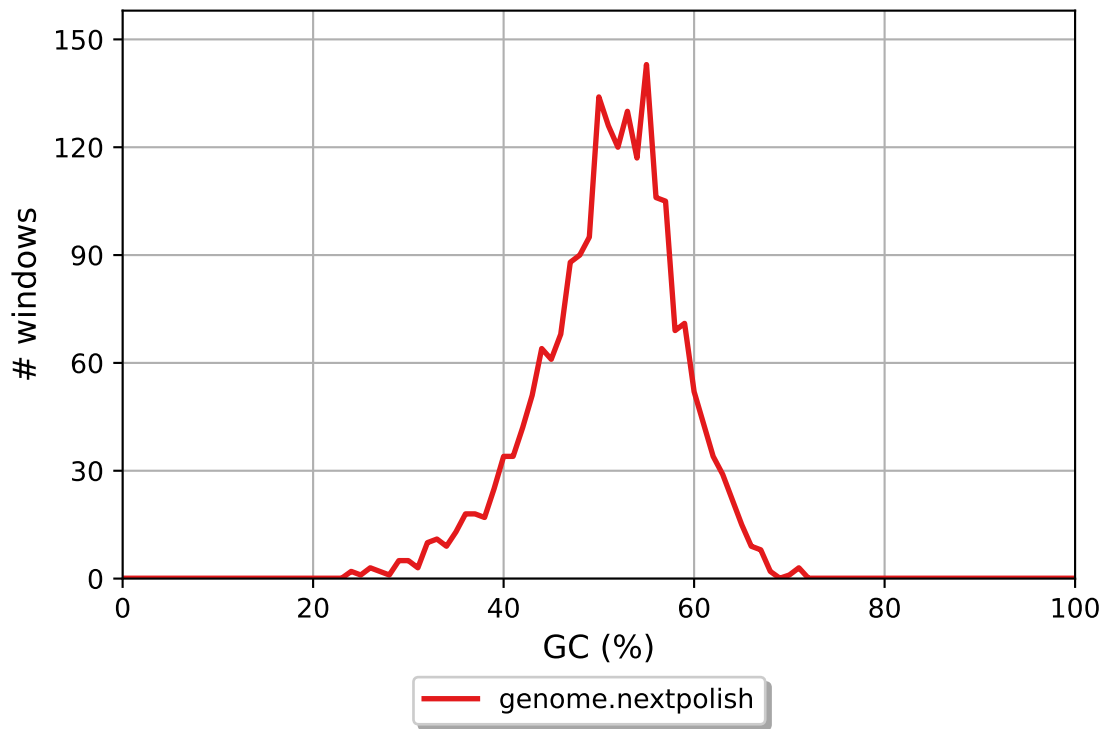


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

