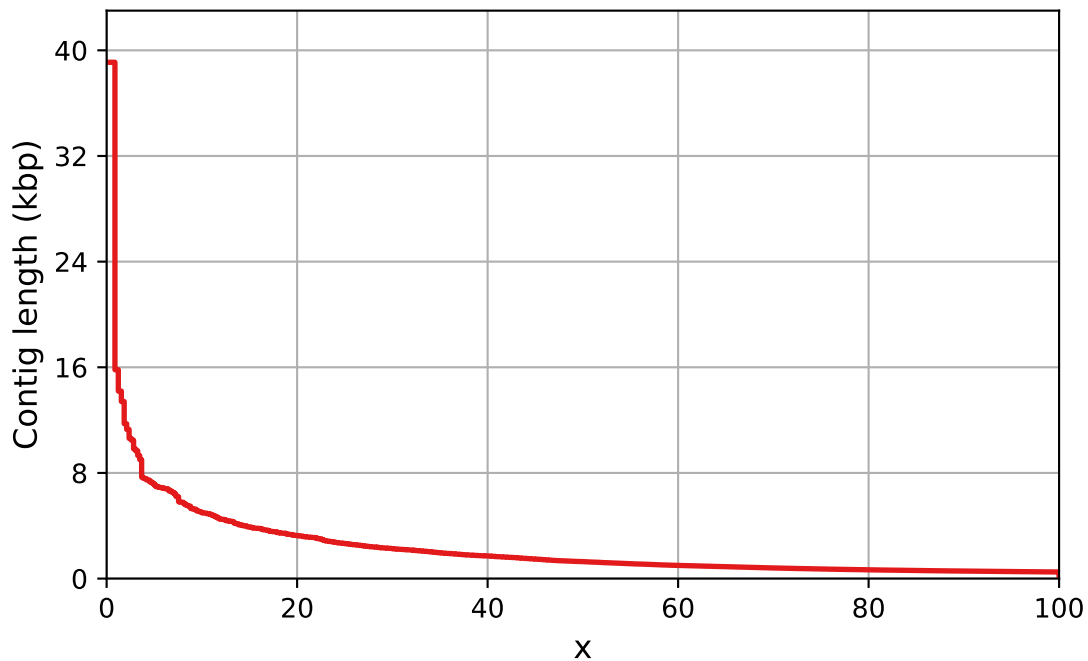


Report

	genome.nextpolish
# contigs (>= 0 bp)	76925
# contigs (>= 1000 bp)	1267
# contigs (>= 5000 bp)	57
# contigs (>= 10000 bp)	8
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	23270143
Total length (>= 1000 bp)	2683069
Total length (>= 5000 bp)	446351
Total length (>= 10000 bp)	126633
Total length (>= 25000 bp)	39095
Total length (>= 50000 bp)	0
# contigs	3950
Largest contig	39095
Total length	4474077
GC (%)	48.47
N50	1284
N90	569
auN	2482.4
L50	872
L90	3111
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 1 part

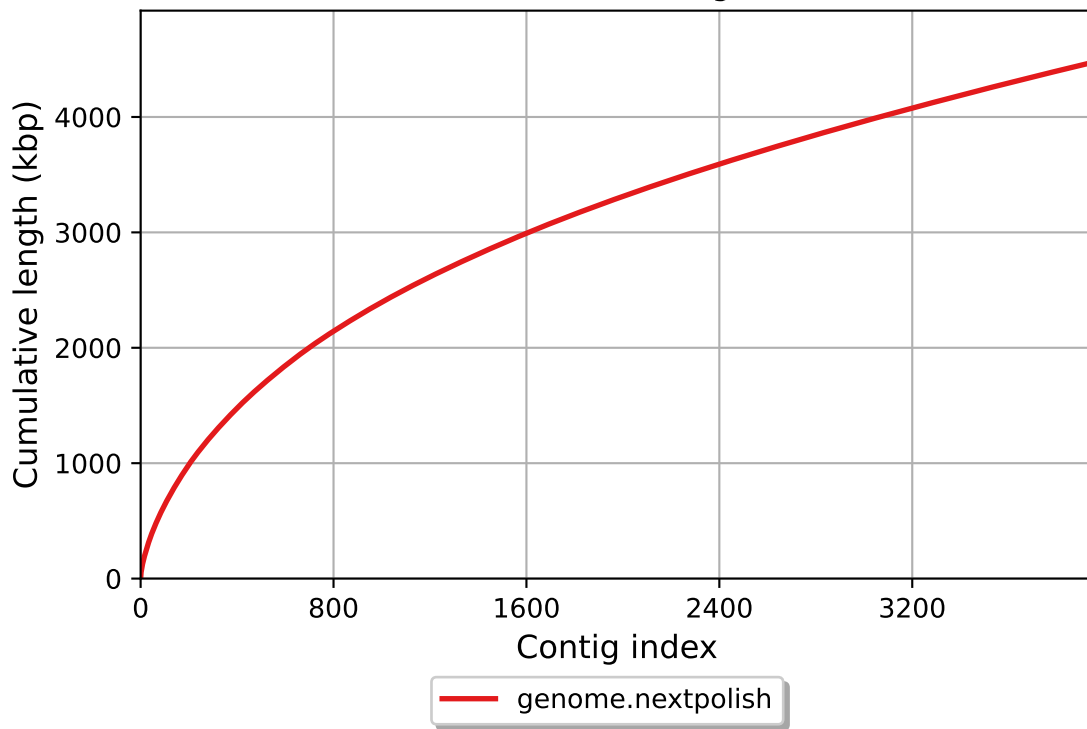
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

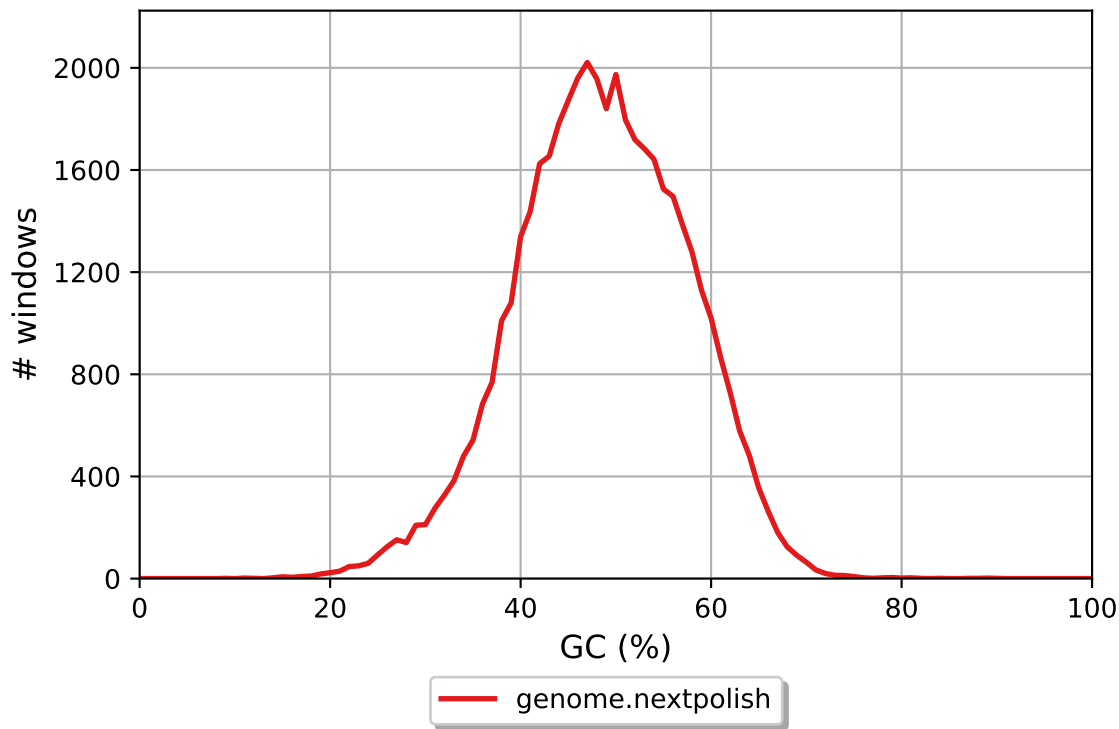


— genome.nextpolish

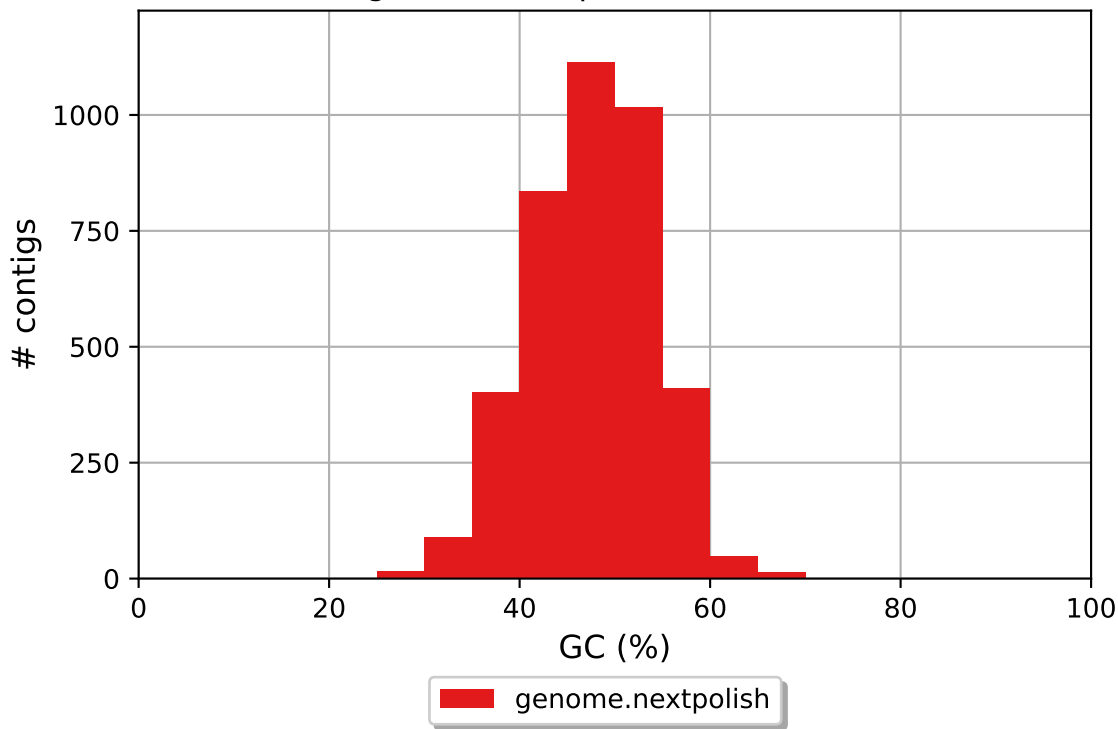
Cumulative length



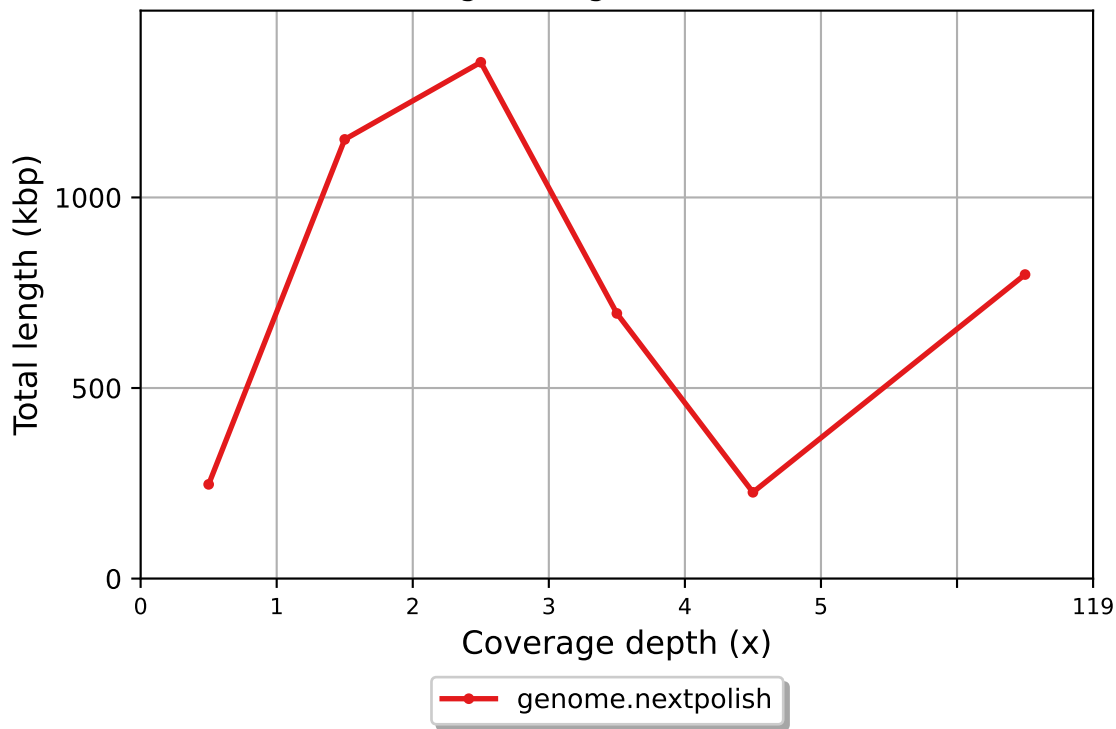
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



genome.nextpolish coverage histogram (bin size: 1x)

