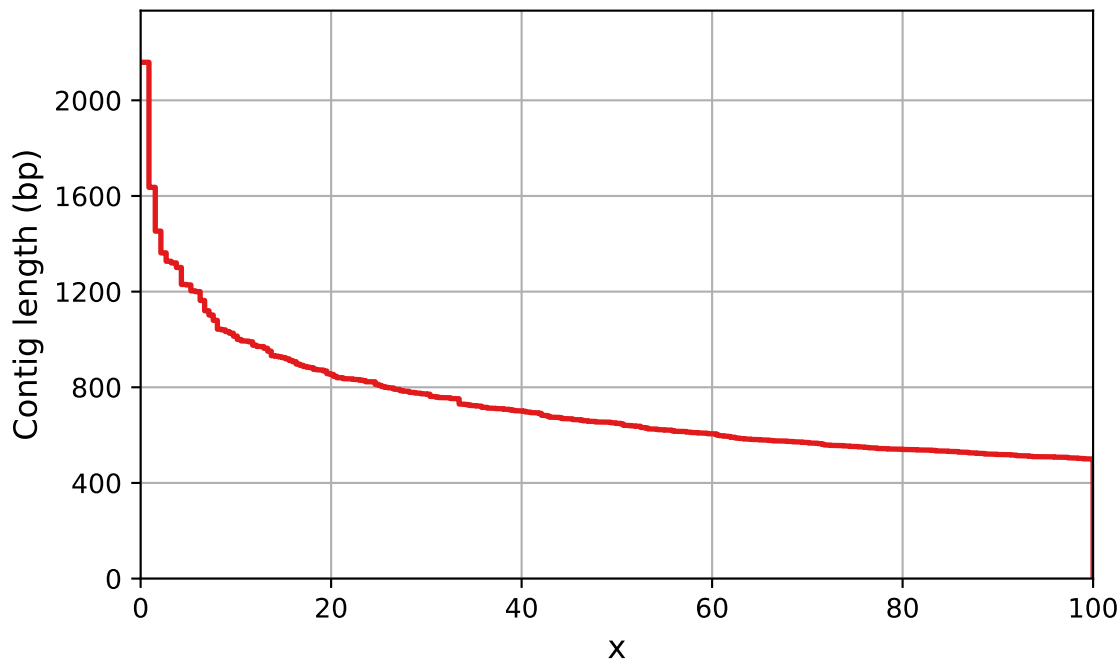


Report

	genome.nextpolish
# contigs (>= 0 bp)	7202
# contigs (>= 1000 bp)	21
# contigs (>= 5000 bp)	0
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1594138
Total length (>= 1000 bp)	26040
Total length (>= 5000 bp)	0
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	369
Largest contig	2159
Total length	246741
GC (%)	55.20
N50	649
N90	519
auN	724.7
L50	148
L90	321
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

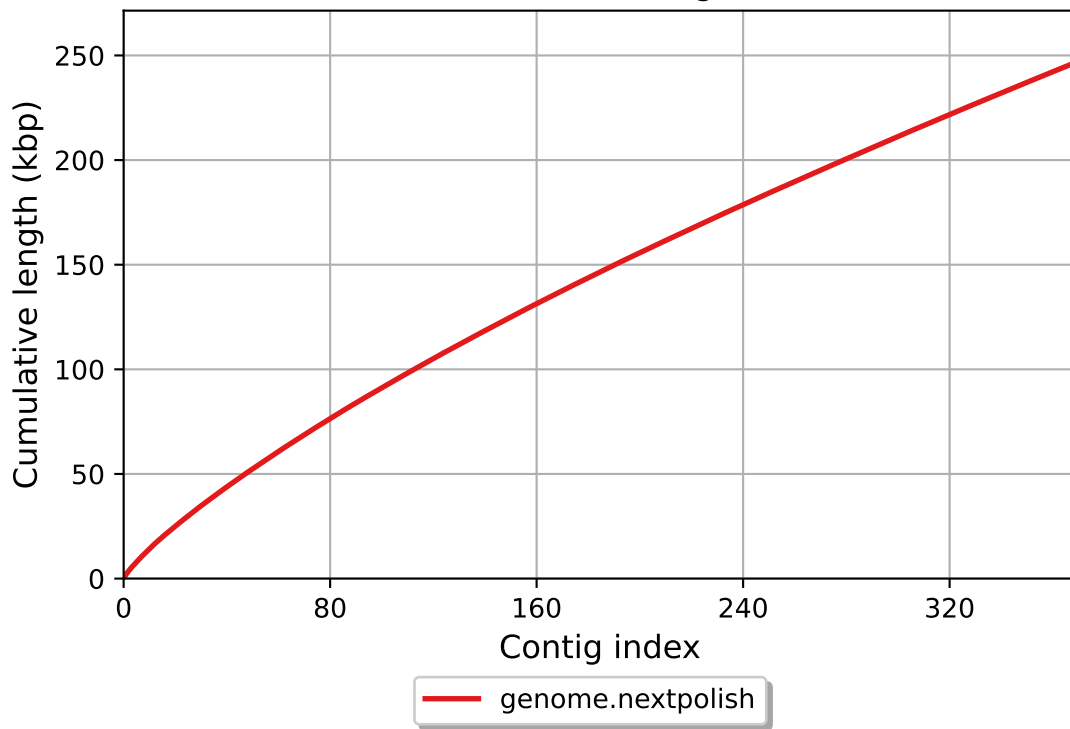
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

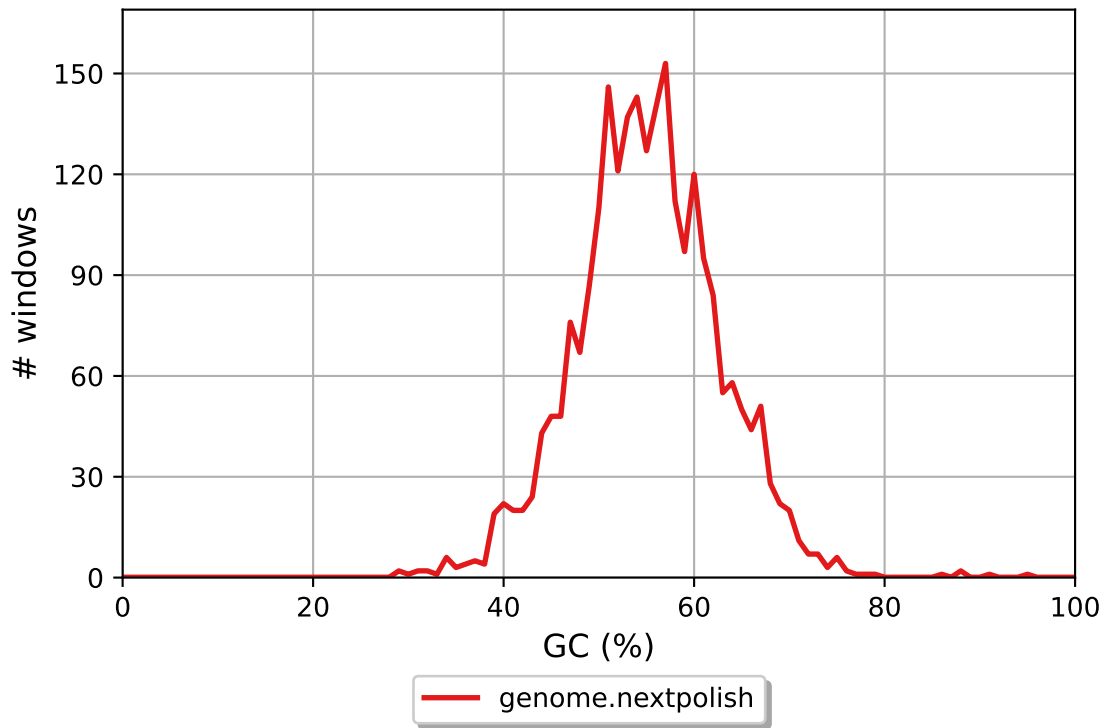


— genome.nextpolish

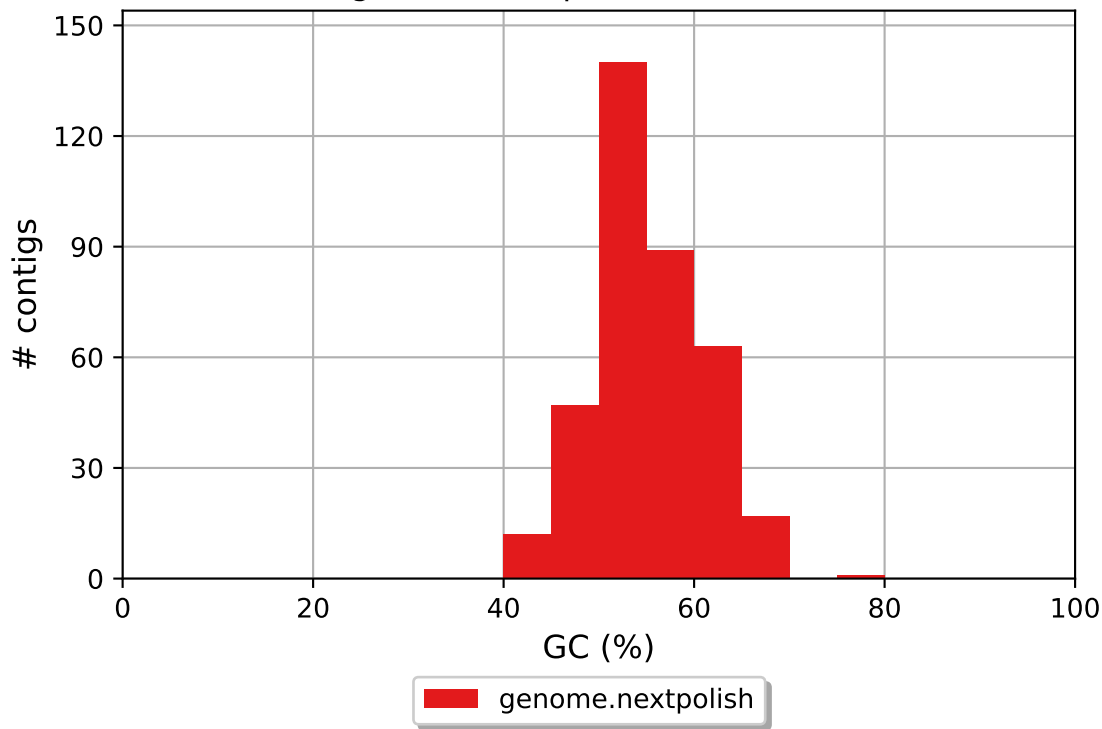
Cumulative length



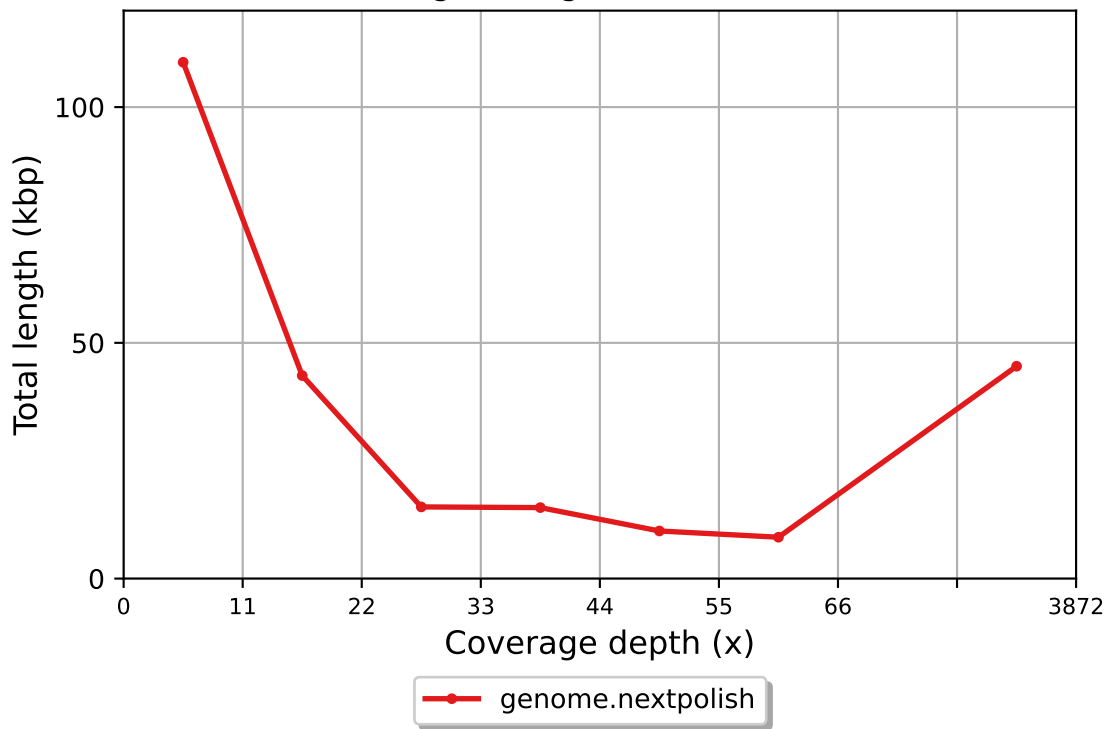
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 11x)



genome.nextpolish coverage histogram (bin size: 11x)

