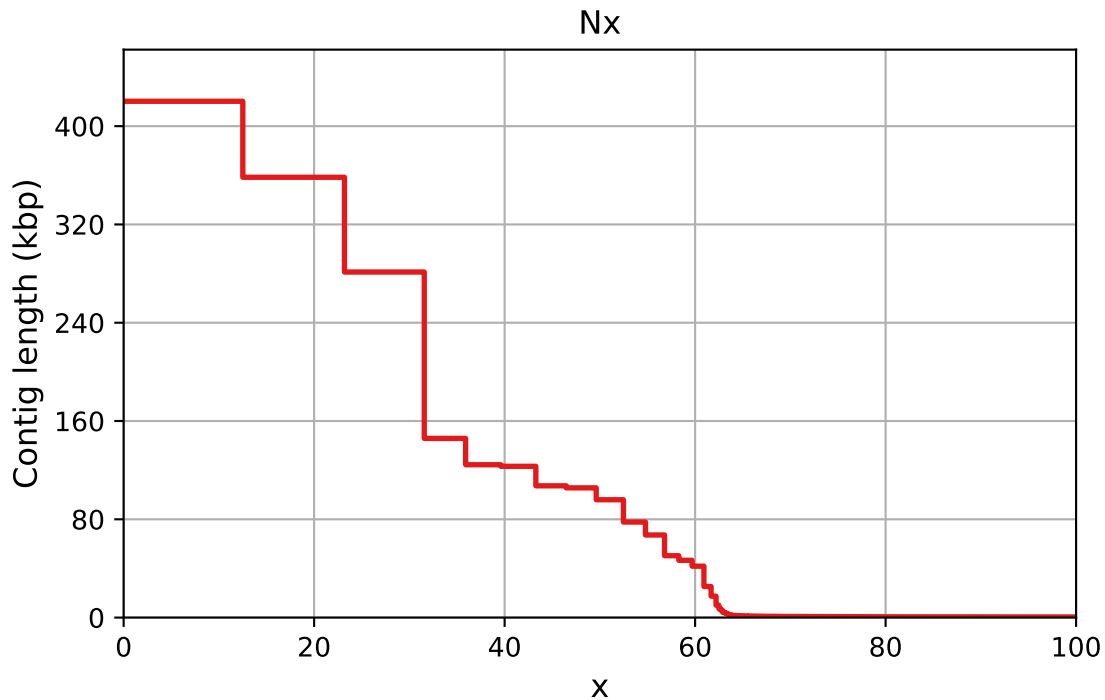


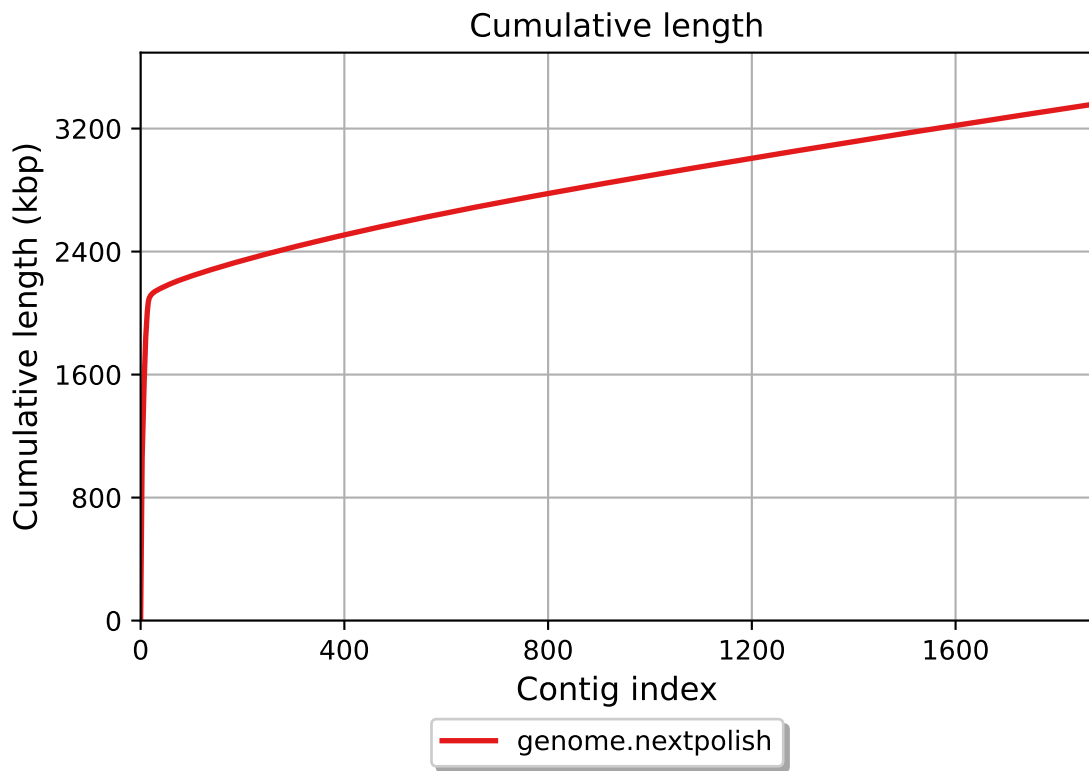
Report

	genome.nextpolish
# contigs (>= 0 bp)	4886
# contigs (>= 1000 bp)	152
# contigs (>= 5000 bp)	19
# contigs (>= 10000 bp)	17
# contigs (>= 25000 bp)	15
# contigs (>= 50000 bp)	12
Total length (>= 0 bp)	4537165
Total length (>= 1000 bp)	2296039
Total length (>= 5000 bp)	2111776
Total length (>= 10000 bp)	2098698
Total length (>= 25000 bp)	2071188
Total length (>= 50000 bp)	1957434
# contigs	1870
Largest contig	420257
Total length	3358021
GC (%)	40.75
N50	95922
N90	548
auN	145049.3
L50	9
L90	1228
# N's per 100 kbp	0.03
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 1 part

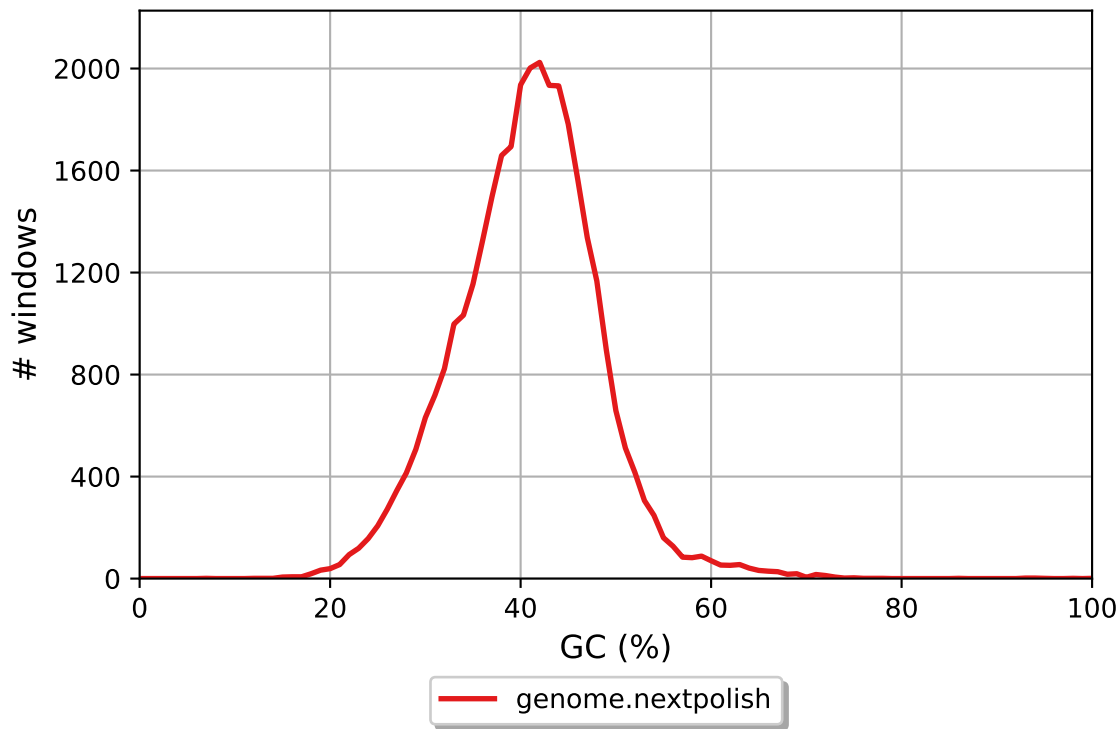
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



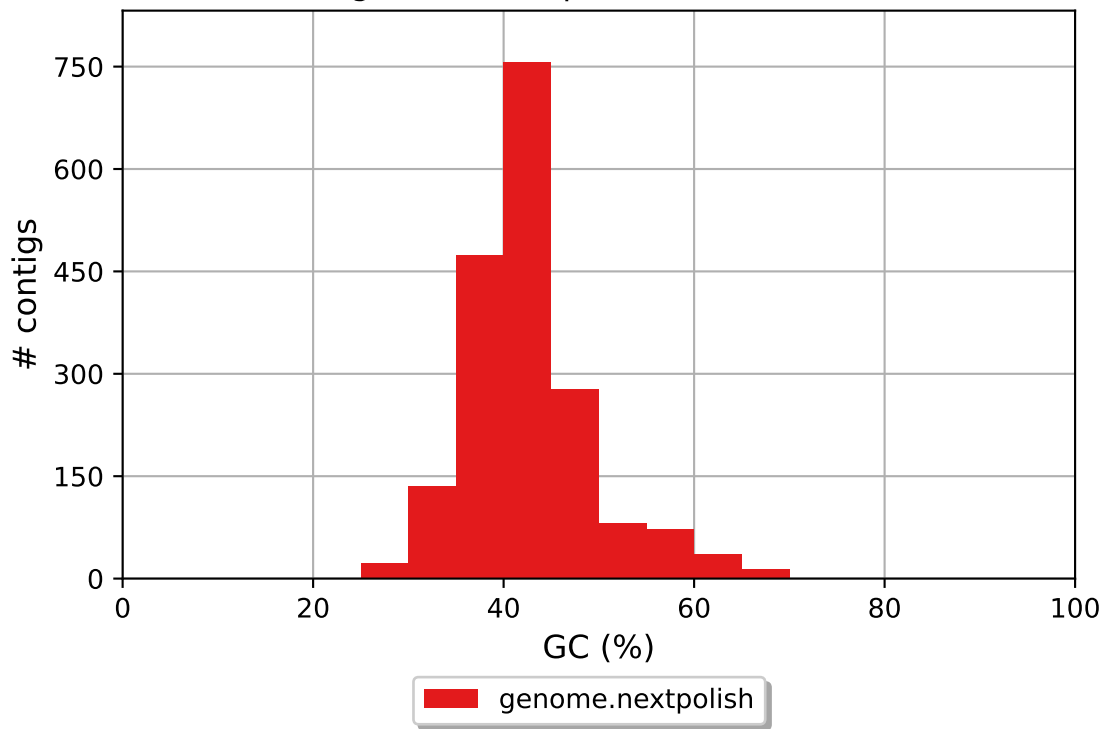
— genome.nextpolish



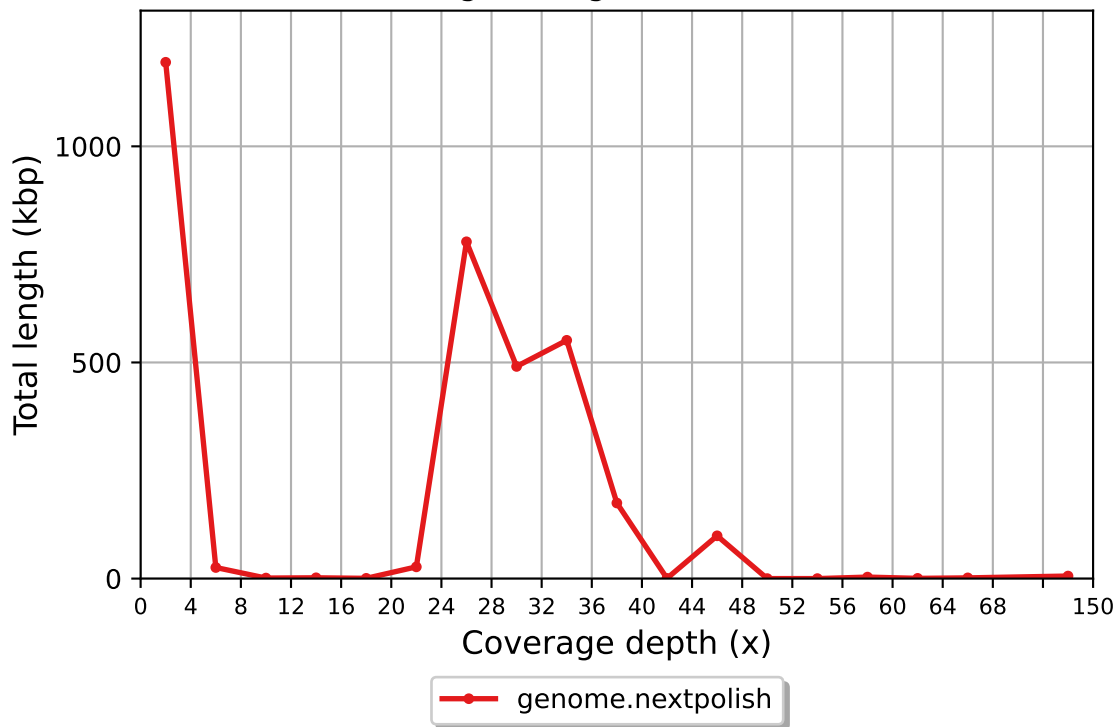
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 4x)



genome.nextpolish coverage histogram (bin size: 4x)

