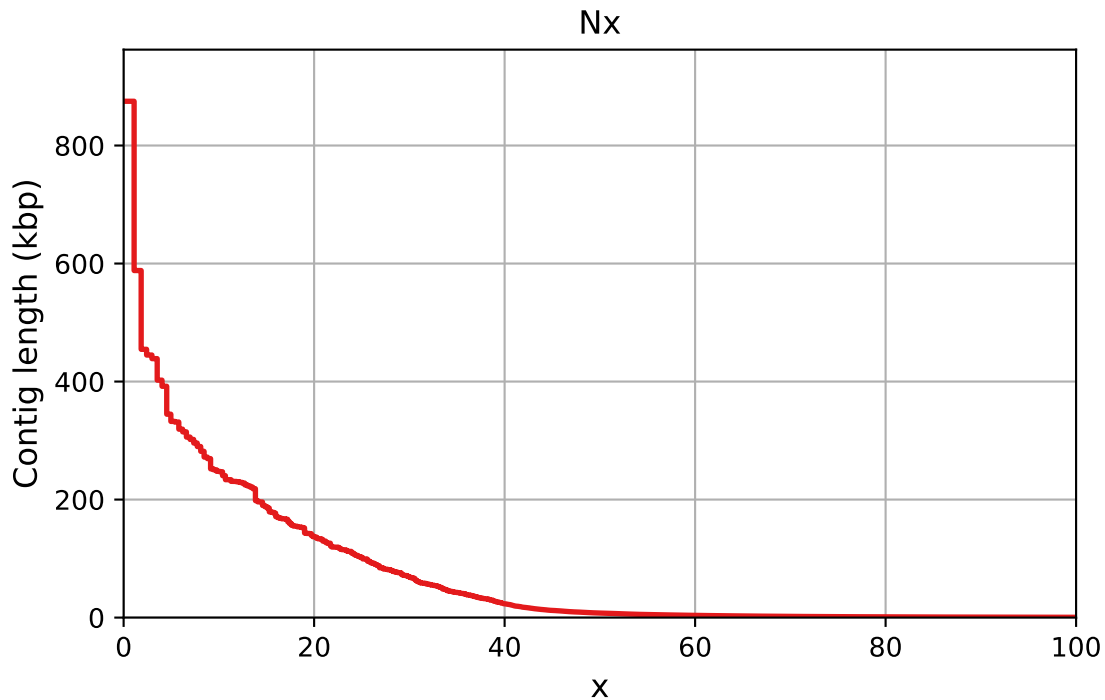


Report

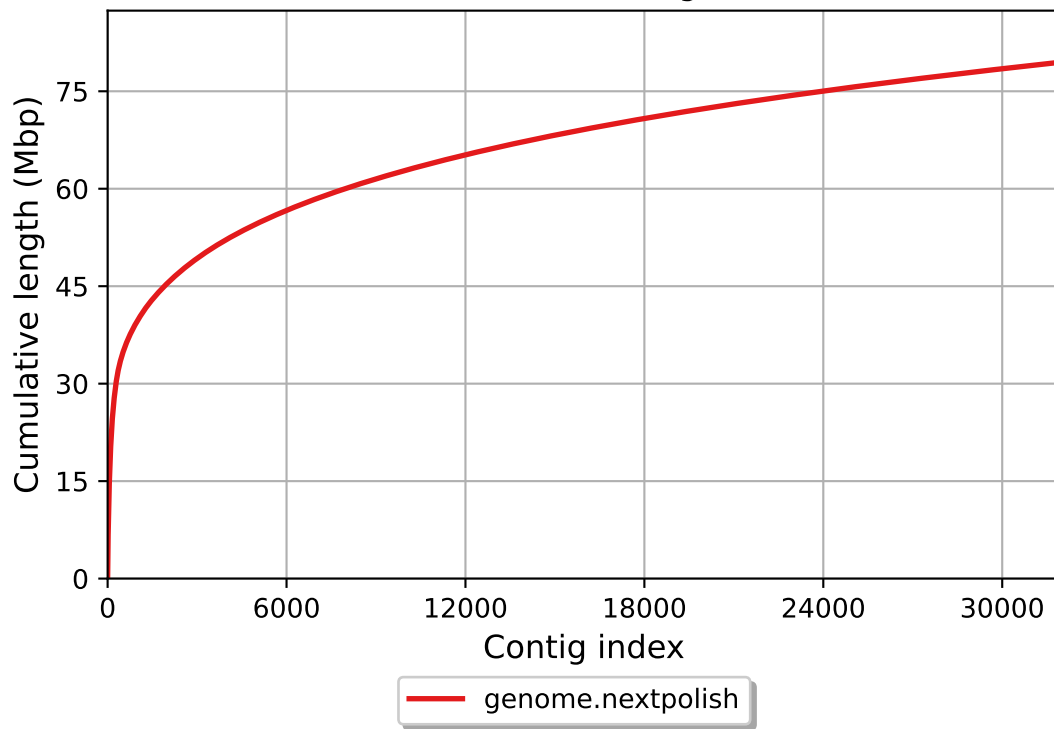
	genome.nextpolish
# contigs (>= 0 bp)	124124
# contigs (>= 1000 bp)	13617
# contigs (>= 5000 bp)	1722
# contigs (>= 10000 bp)	727
# contigs (>= 25000 bp)	332
# contigs (>= 50000 bp)	194
Total length (>= 0 bp)	105977221
Total length (>= 1000 bp)	66918779
Total length (>= 5000 bp)	44125282
Total length (>= 10000 bp)	37303969
Total length (>= 25000 bp)	31523689
Total length (>= 50000 bp)	26609696
# contigs	31941
Largest contig	874989
Total length	79463316
GC (%)	45.05
N50	7644
N90	765
auN	77934.3
L50	1006
L90	18904
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	34 + 11 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).

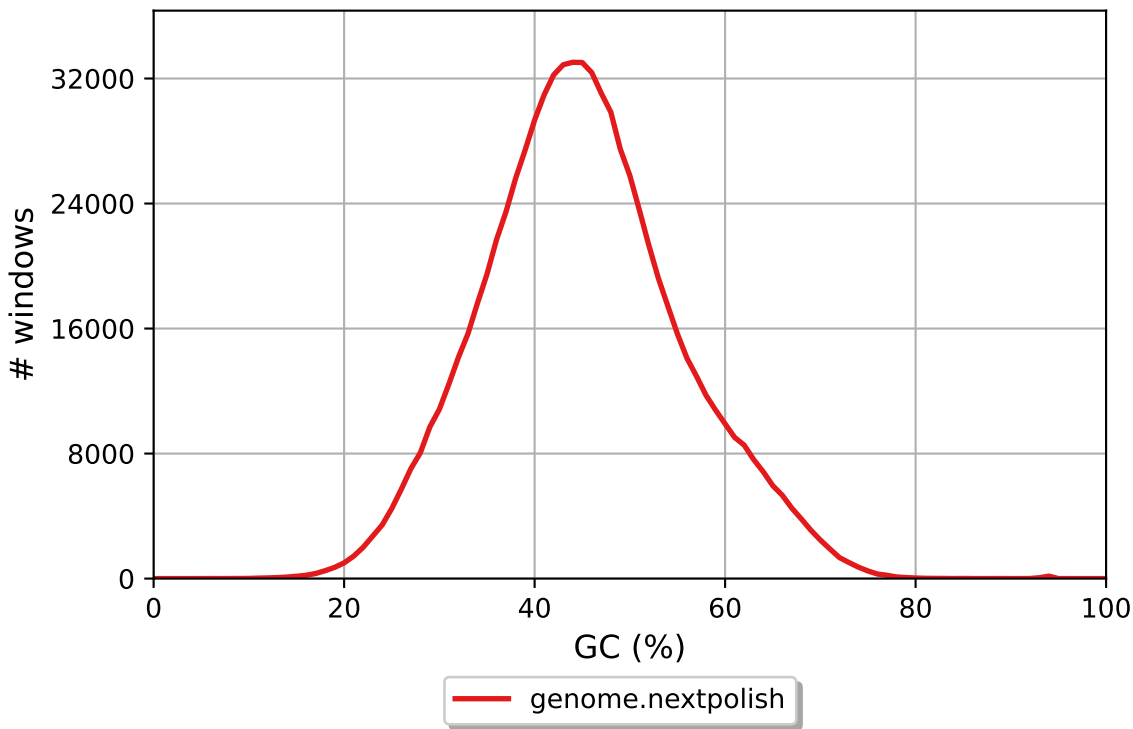


— genome.nextpolish

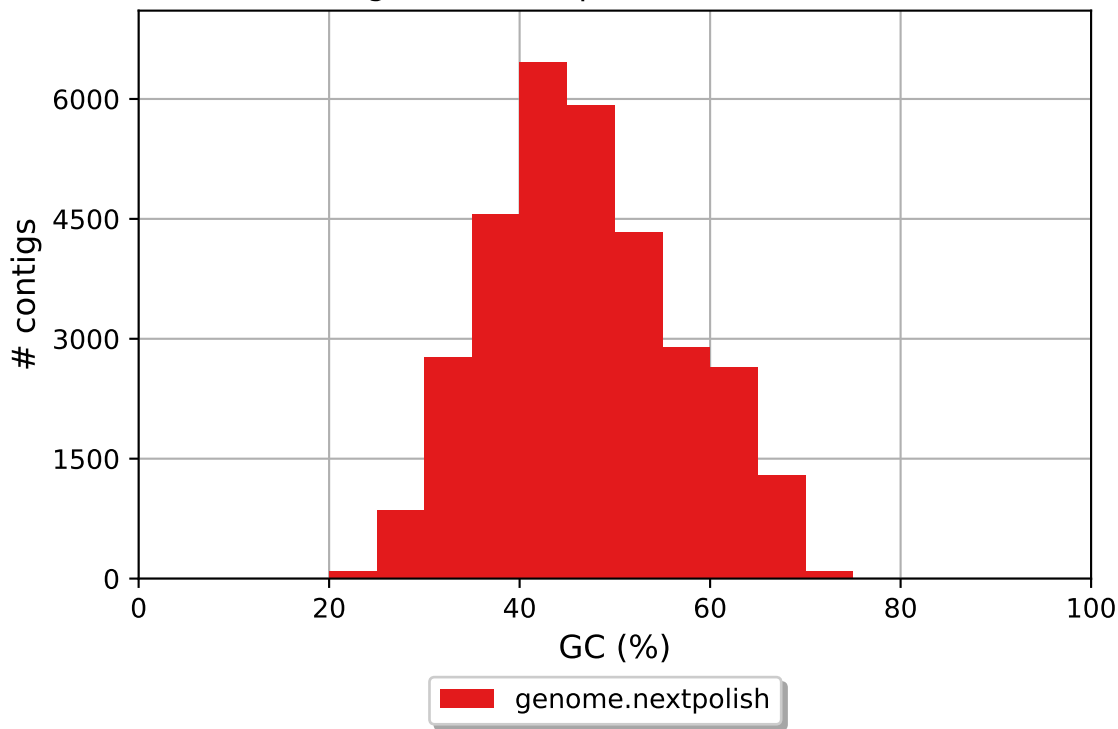
Cumulative length



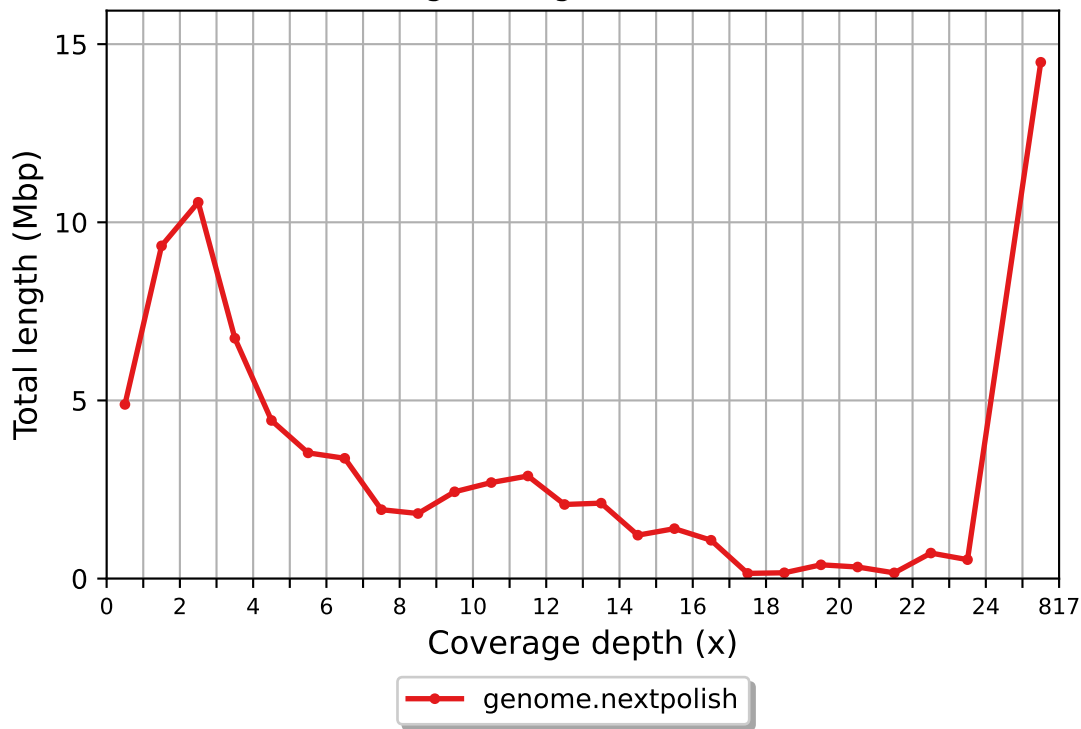
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

