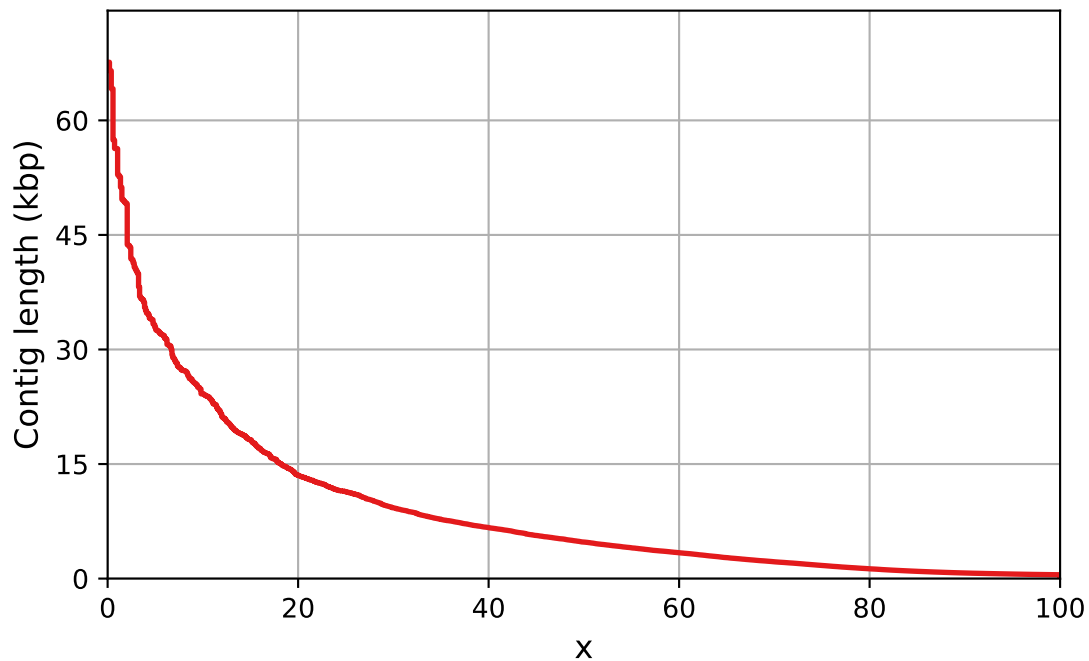


Report

	genome.nextpolish
# contigs (>= 0 bp)	27178
# contigs (>= 1000 bp)	7210
# contigs (>= 5000 bp)	1597
# contigs (>= 10000 bp)	551
# contigs (>= 25000 bp)	98
# contigs (>= 50000 bp)	9
Total length (>= 0 bp)	39628616
Total length (>= 1000 bp)	29511063
Total length (>= 5000 bp)	17112354
Total length (>= 10000 bp)	9939772
Total length (>= 25000 bp)	3390608
Total length (>= 50000 bp)	525014
# contigs	15558
Largest contig	67609
Total length	35087841
GC (%)	46.20
N50	4780
N90	732
auN	9025.5
L50	1686
L90	9649
# N's per 100 kbp	1.55
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	19 + 12 part

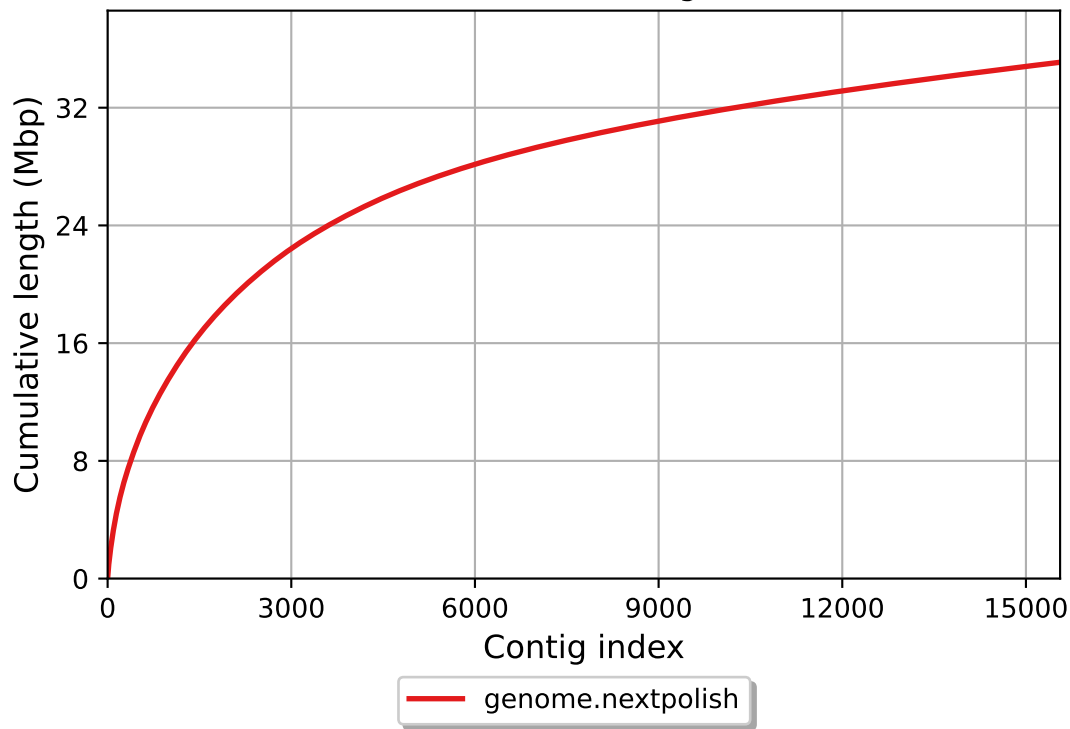
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

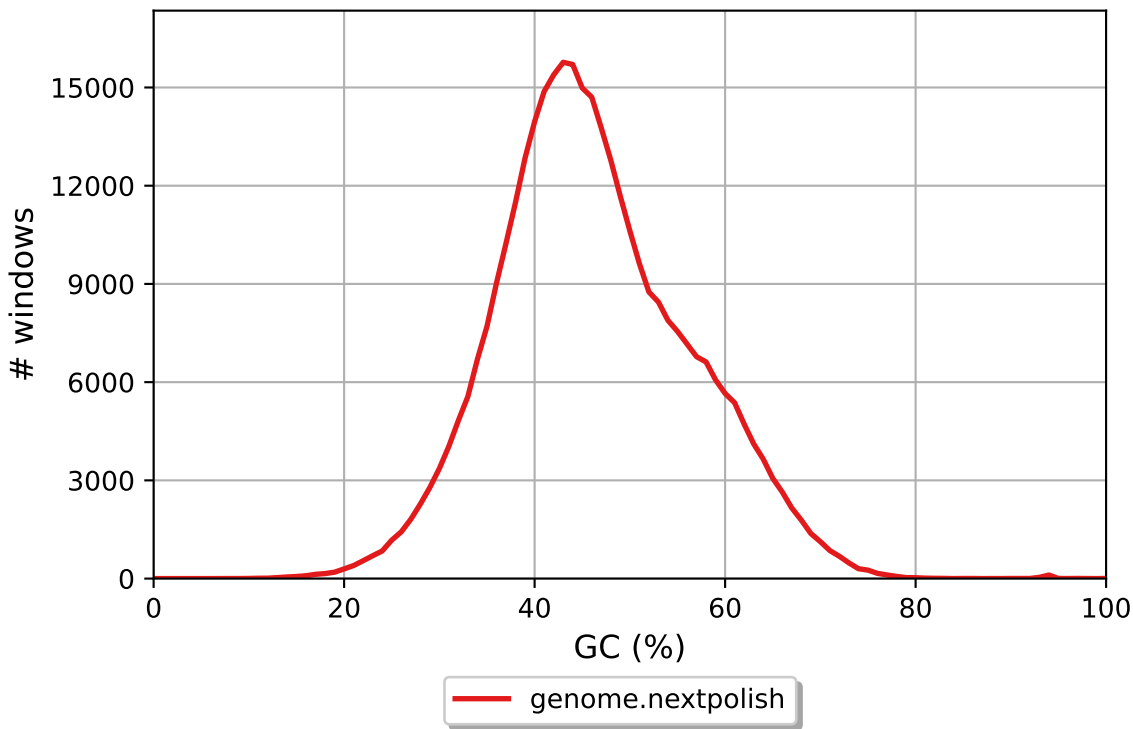


— genome.nextpolish

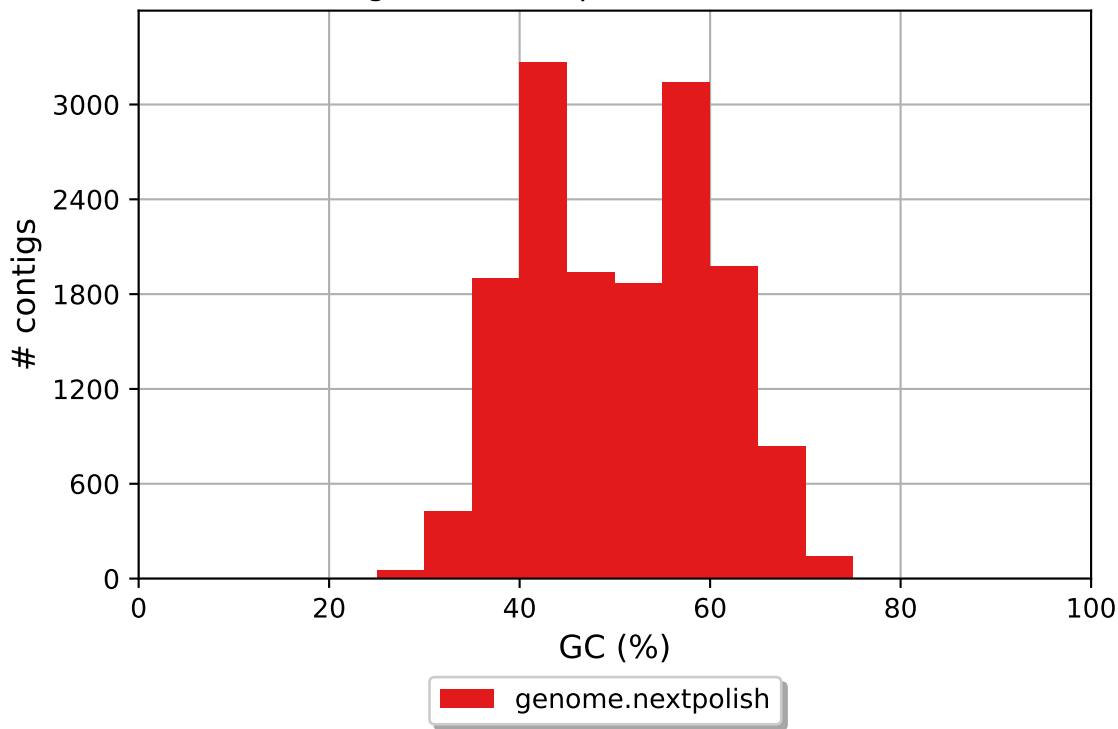
Cumulative length



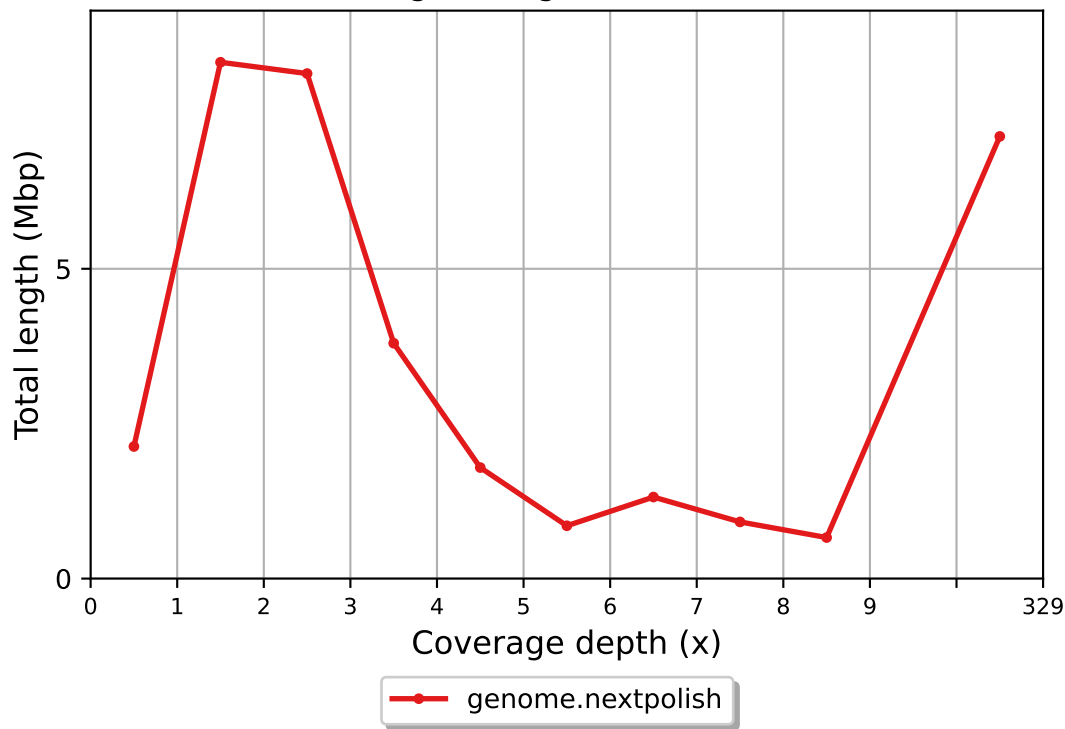
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

