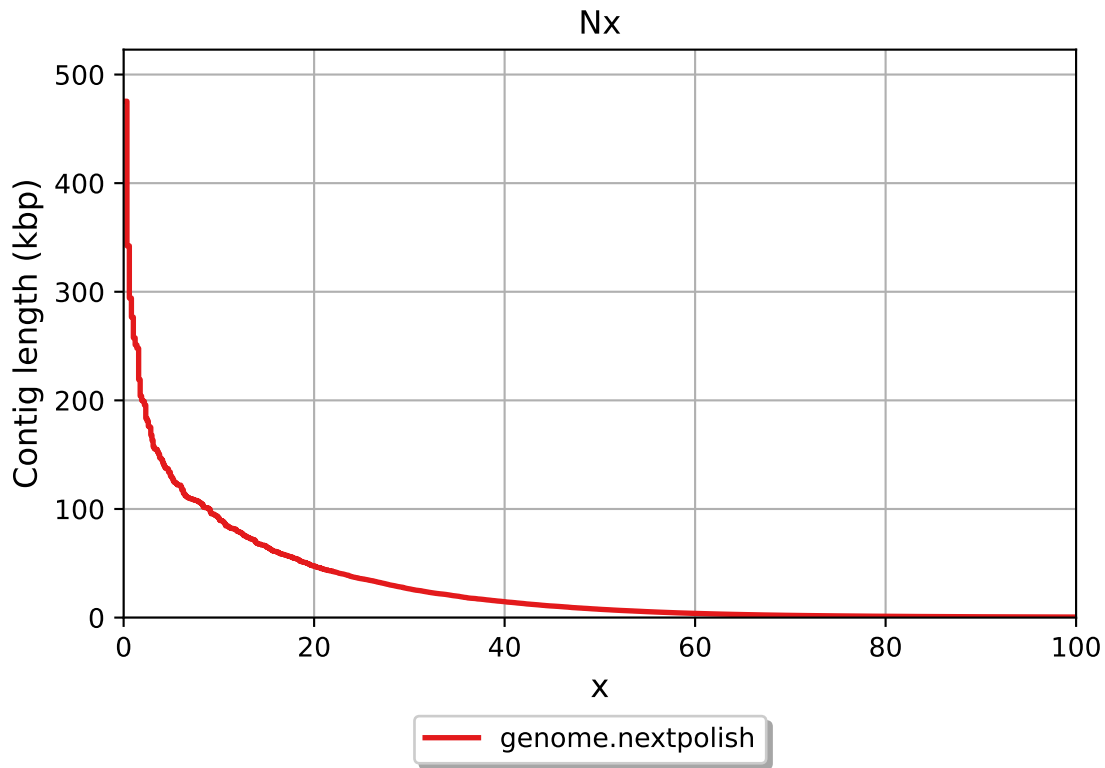


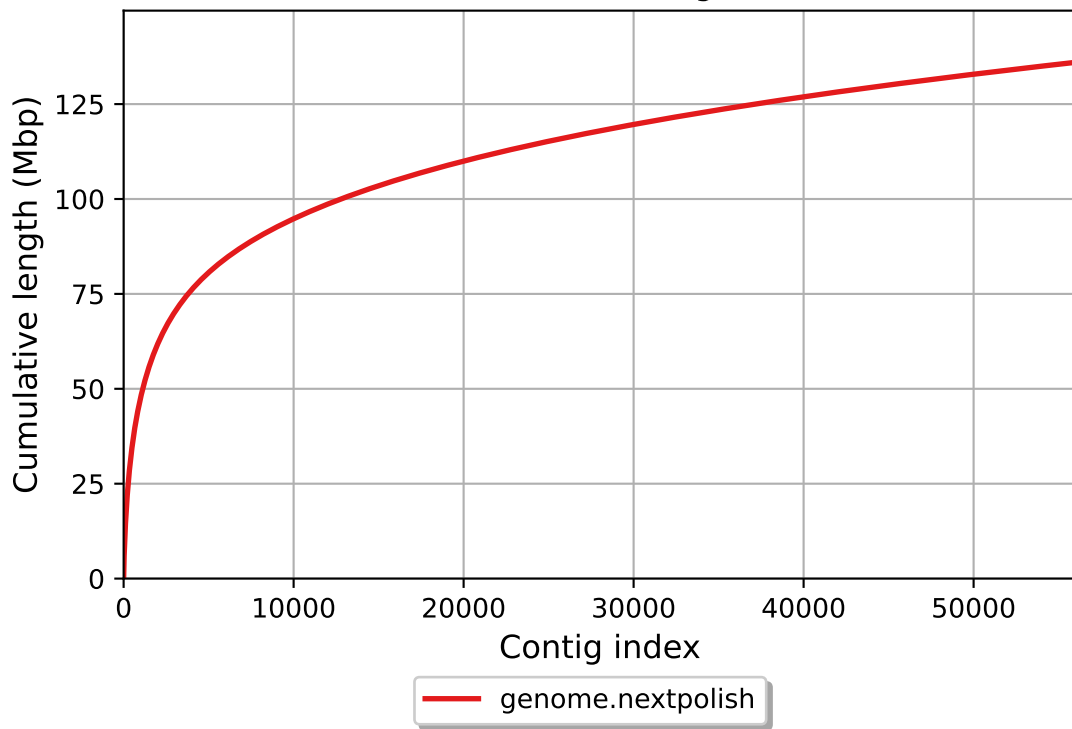
Report

	genome.nextpolish
# contigs (>= 0 bp)	205243
# contigs (>= 1000 bp)	23652
# contigs (>= 5000 bp)	4081
# contigs (>= 10000 bp)	2091
# contigs (>= 25000 bp)	747
# contigs (>= 50000 bp)	290
Total length (>= 0 bp)	179181461
Total length (>= 1000 bp)	113895889
Total length (>= 5000 bp)	76501164
Total length (>= 10000 bp)	62698452
Total length (>= 25000 bp)	42085479
Total length (>= 50000 bp)	26188253
# contigs	56029
Largest contig	475392
Total length	136020333
GC (%)	46.95
N50	7627
N90	750
auN	30679.7
L50	2702
L90	33542
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	45 + 22 part

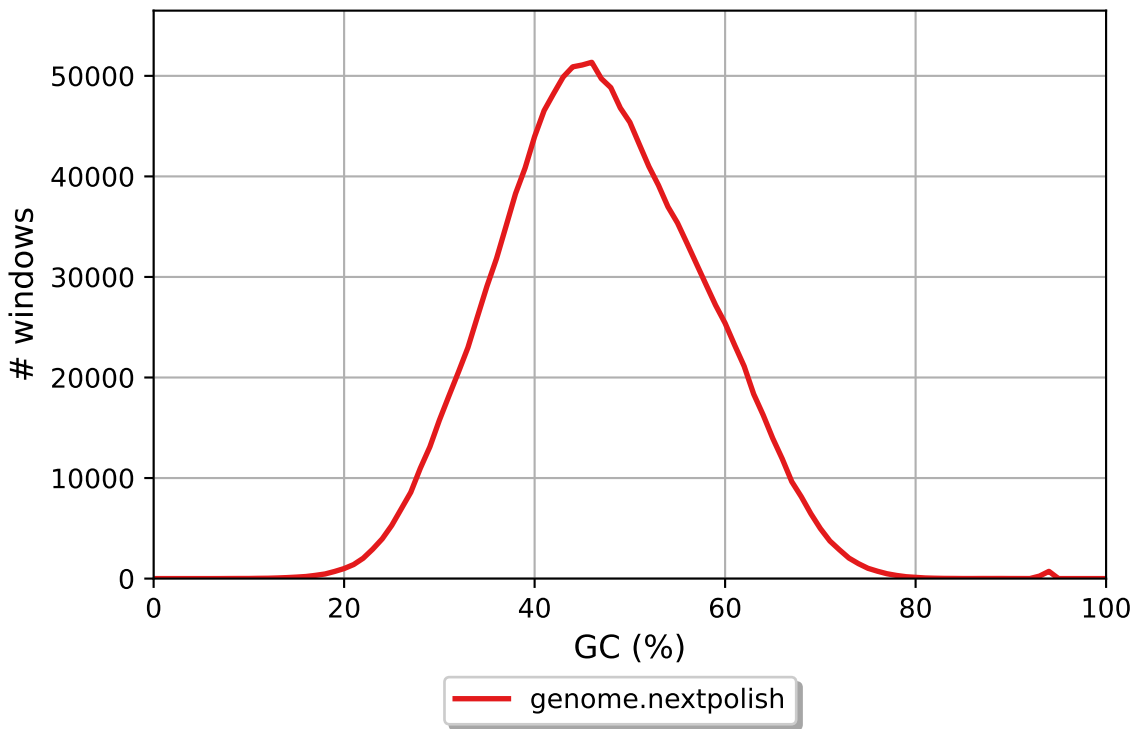
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



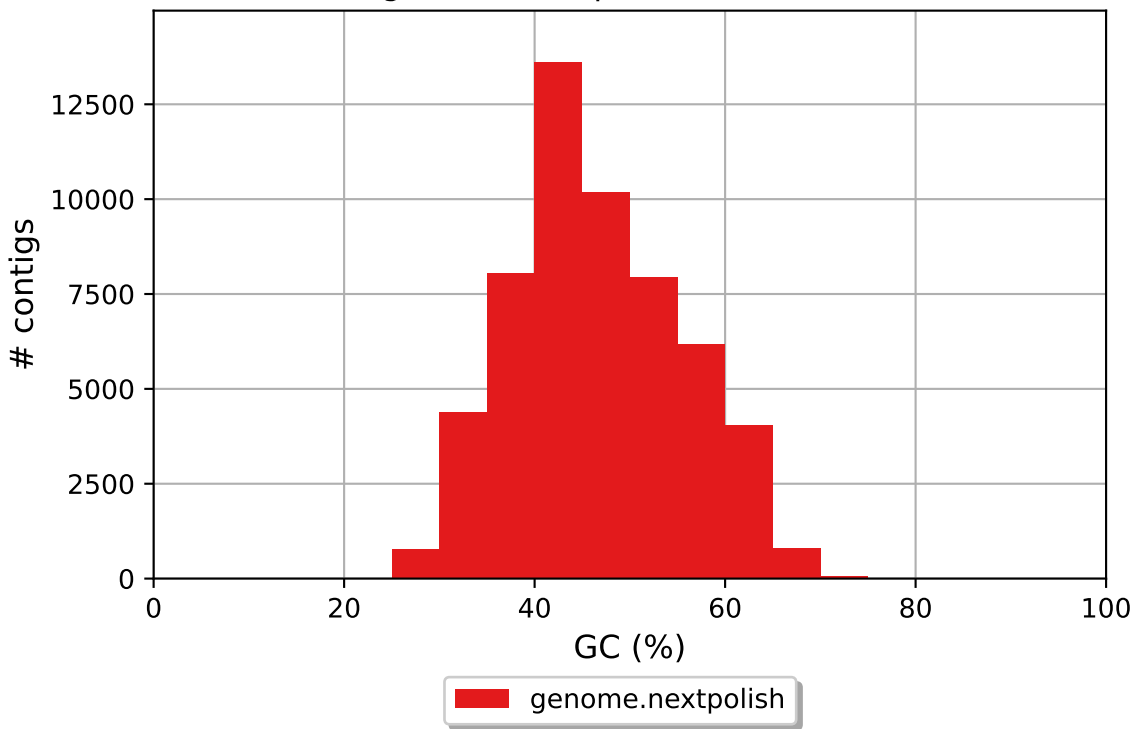
Cumulative length



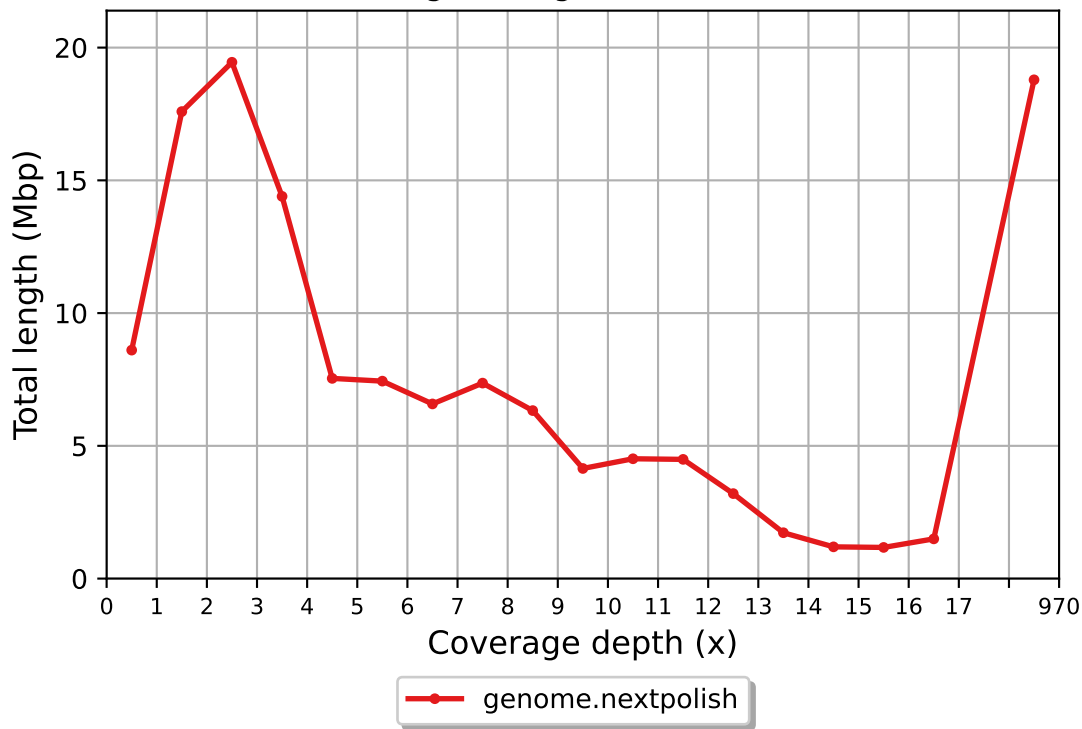
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

