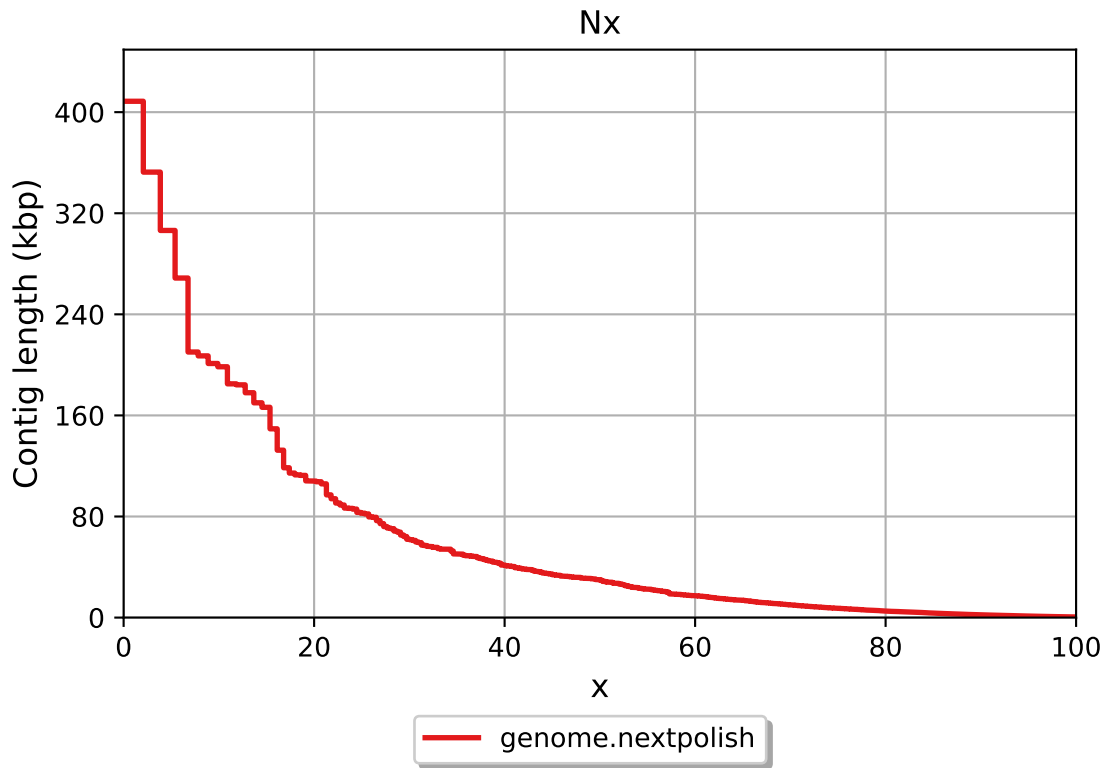


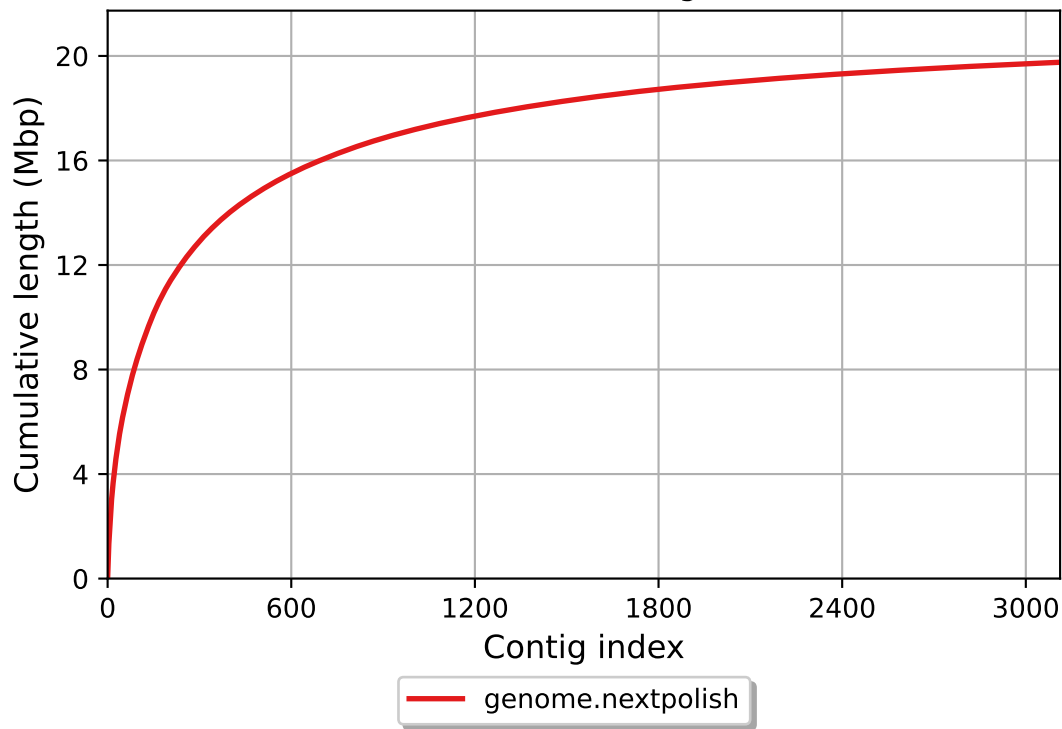
Report

	genome.nextpolish
# contigs (>= 0 bp)	10806
# contigs (>= 1000 bp)	2070
# contigs (>= 5000 bp)	671
# contigs (>= 10000 bp)	385
# contigs (>= 25000 bp)	162
# contigs (>= 50000 bp)	65
Total length (>= 0 bp)	21408140
Total length (>= 1000 bp)	19024944
Total length (>= 5000 bp)	15887552
Total length (>= 10000 bp)	13881811
Total length (>= 25000 bp)	10453966
Total length (>= 50000 bp)	7040061
# contigs	3112
Largest contig	408613
Total length	19758112
GC (%)	40.42
N50	29952
N90	2208
auN	66960.7
L50	141
L90	1238
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 3 part

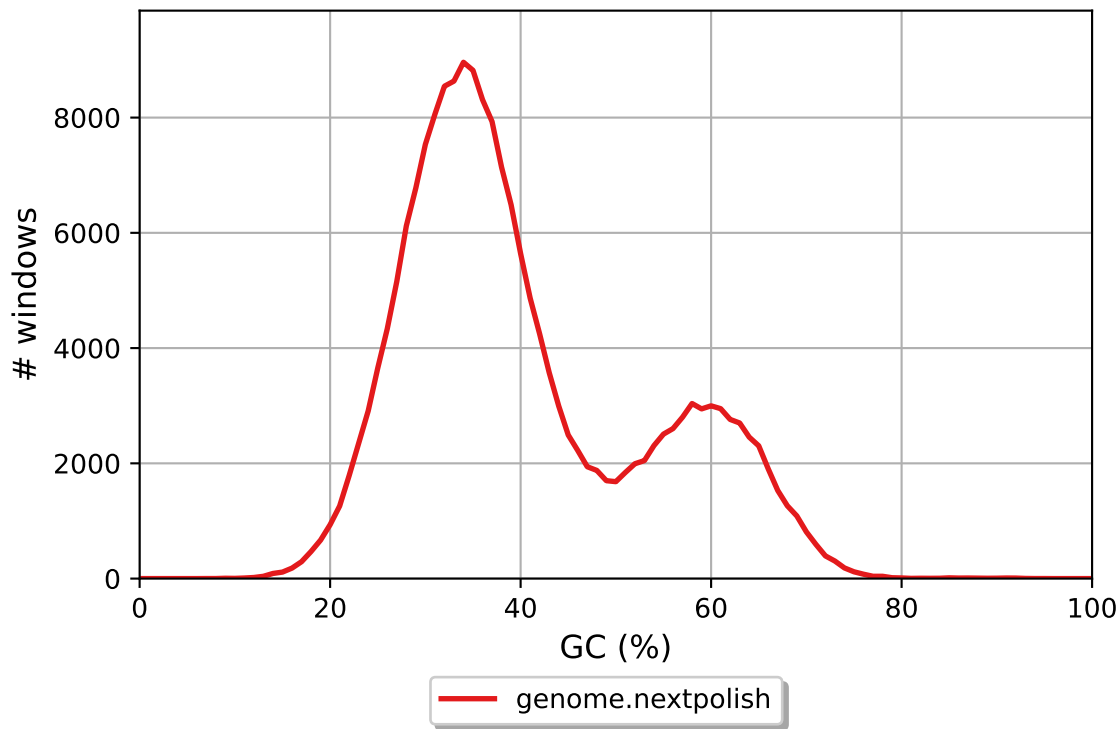
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



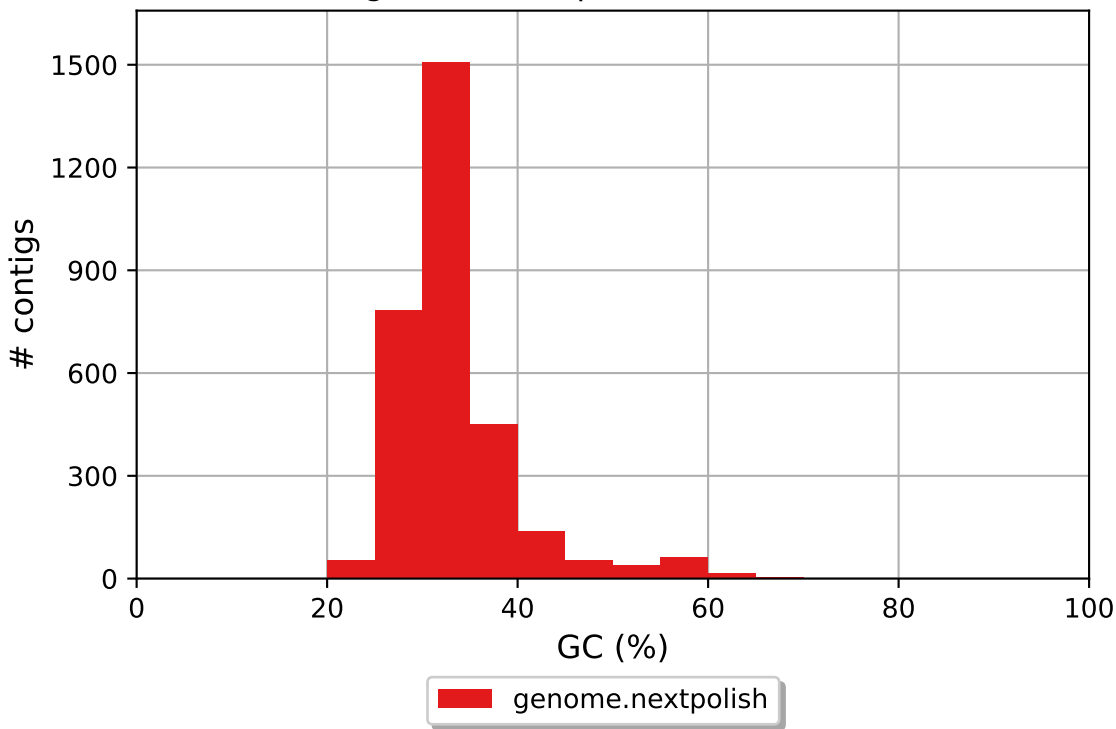
Cumulative length



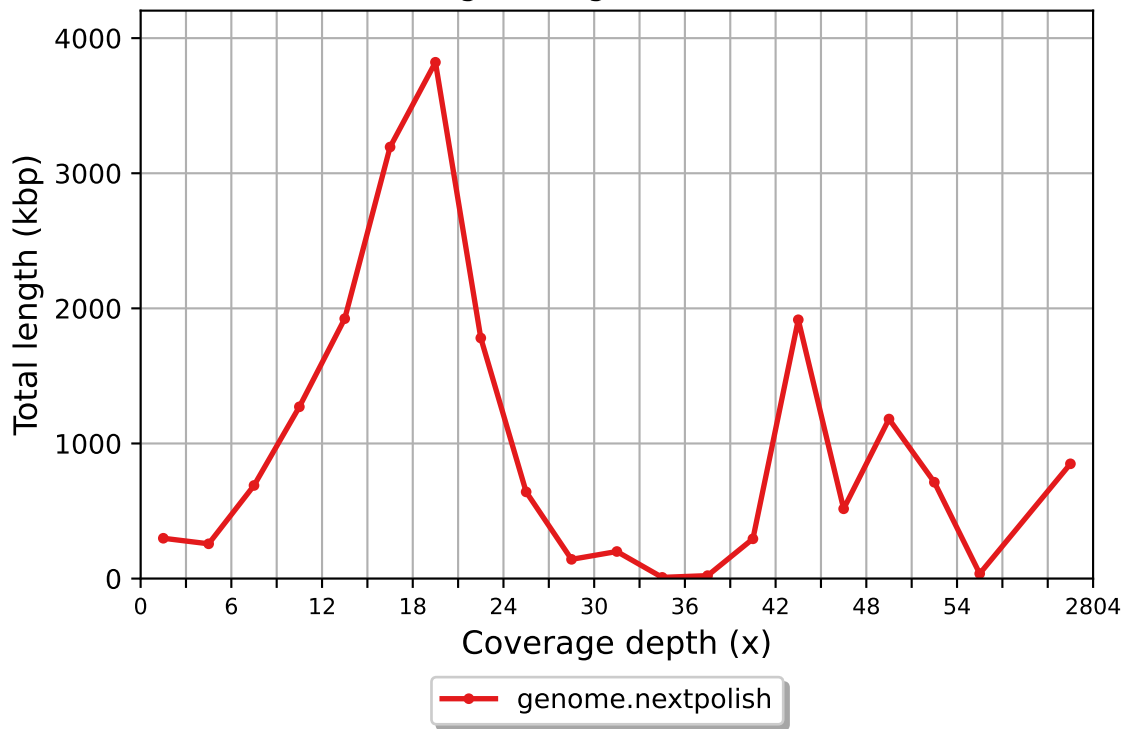
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 3x)



genome.nextpolish coverage histogram (bin size: 3x)

