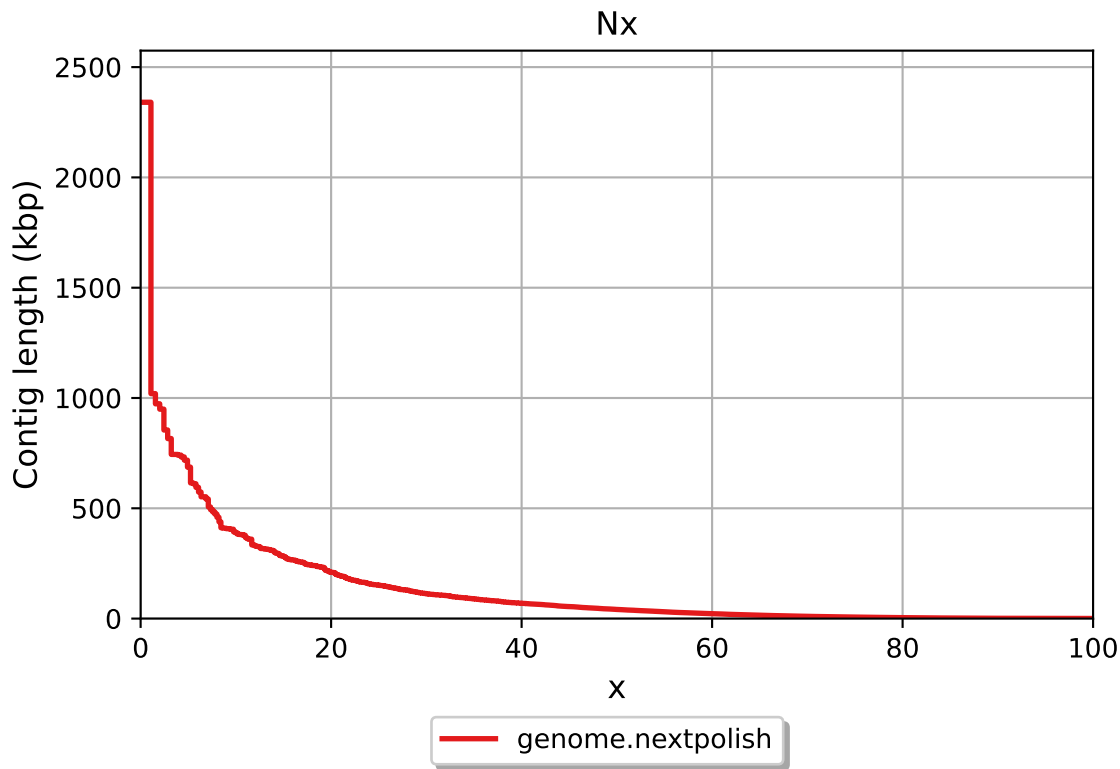


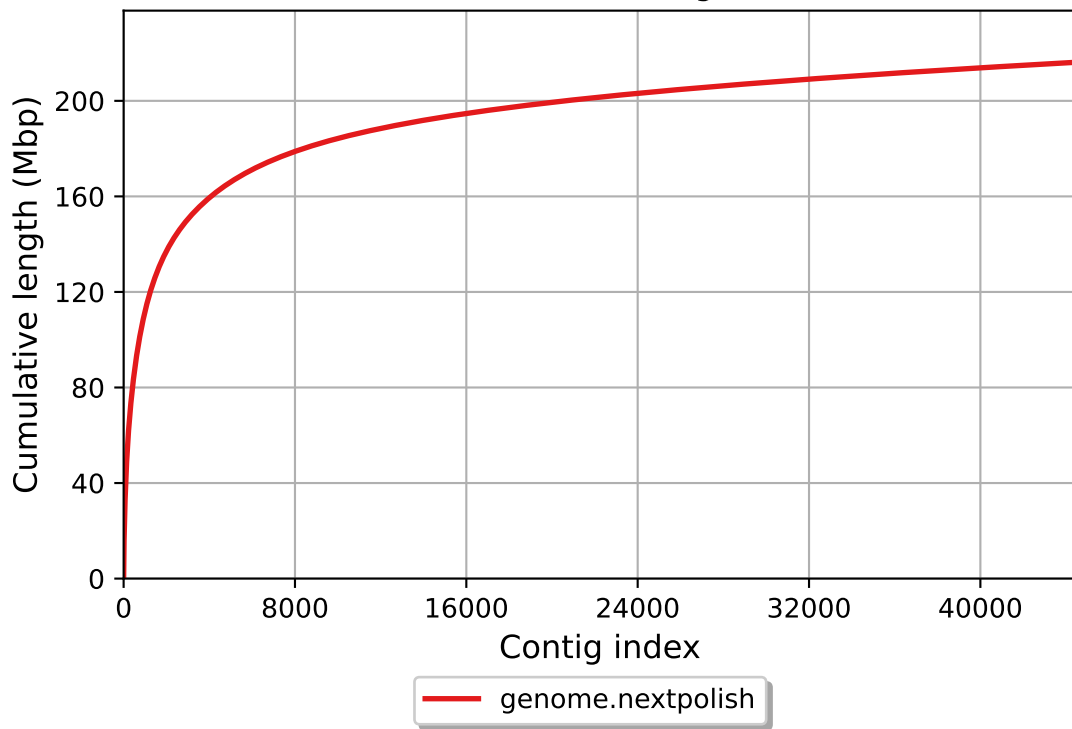
Report

	genome.nextpolish
# contigs (>= 0 bp)	183118
# contigs (>= 1000 bp)	20666
# contigs (>= 5000 bp)	5647
# contigs (>= 10000 bp)	3130
# contigs (>= 25000 bp)	1460
# contigs (>= 50000 bp)	741
Total length (>= 0 bp)	255315328
Total length (>= 1000 bp)	200066826
Total length (>= 5000 bp)	169524782
Total length (>= 10000 bp)	151855890
Total length (>= 25000 bp)	125934827
Total length (>= 50000 bp)	100704637
# contigs	44469
Largest contig	2340634
Total length	216151634
GC (%)	45.37
N50	41545
N90	1338
auN	147962.8
L50	903
L90	15862
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	176 + 26 part

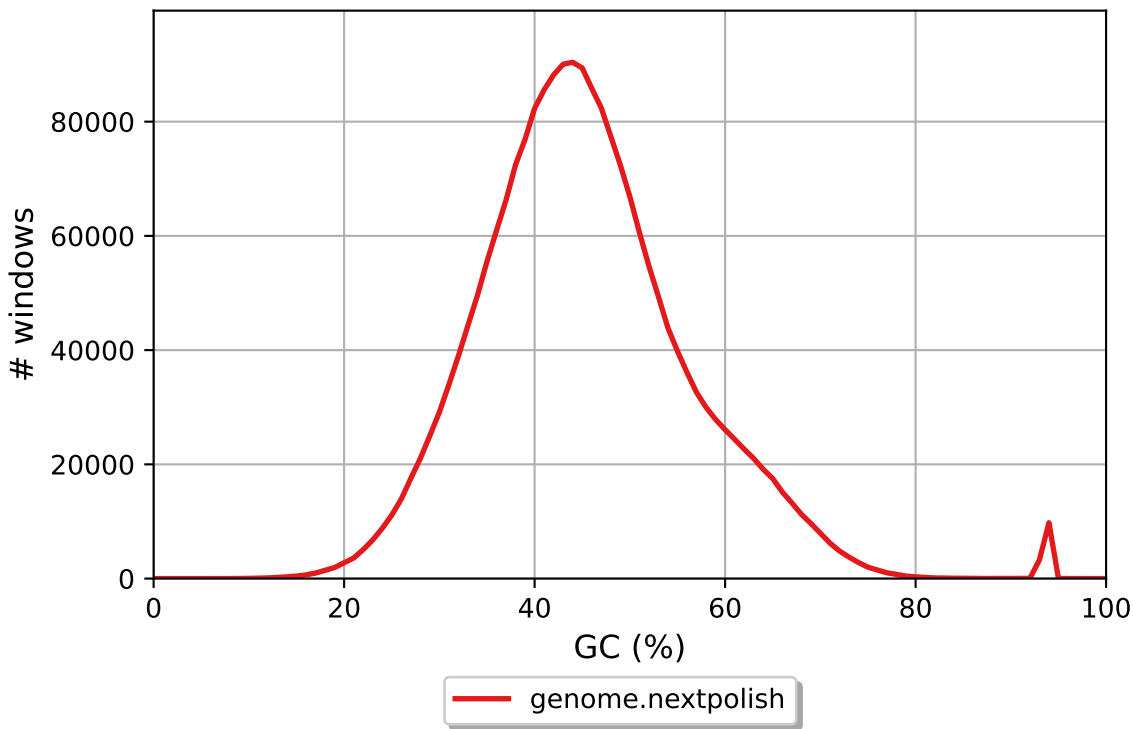
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



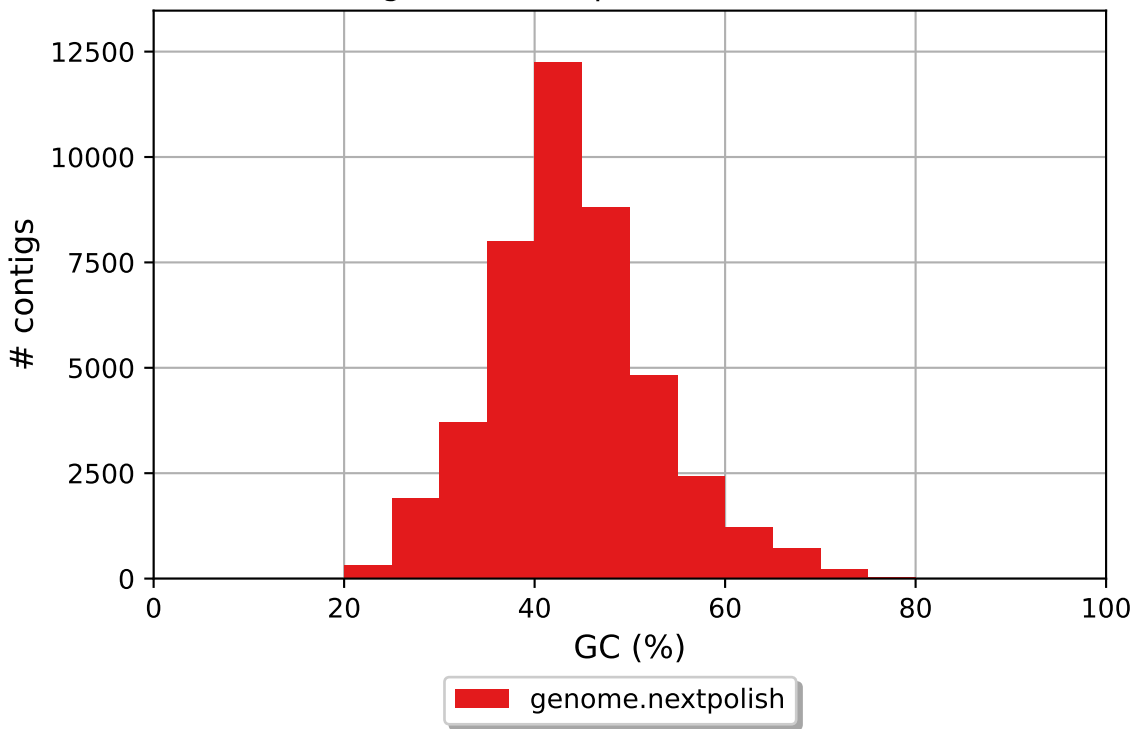
Cumulative length



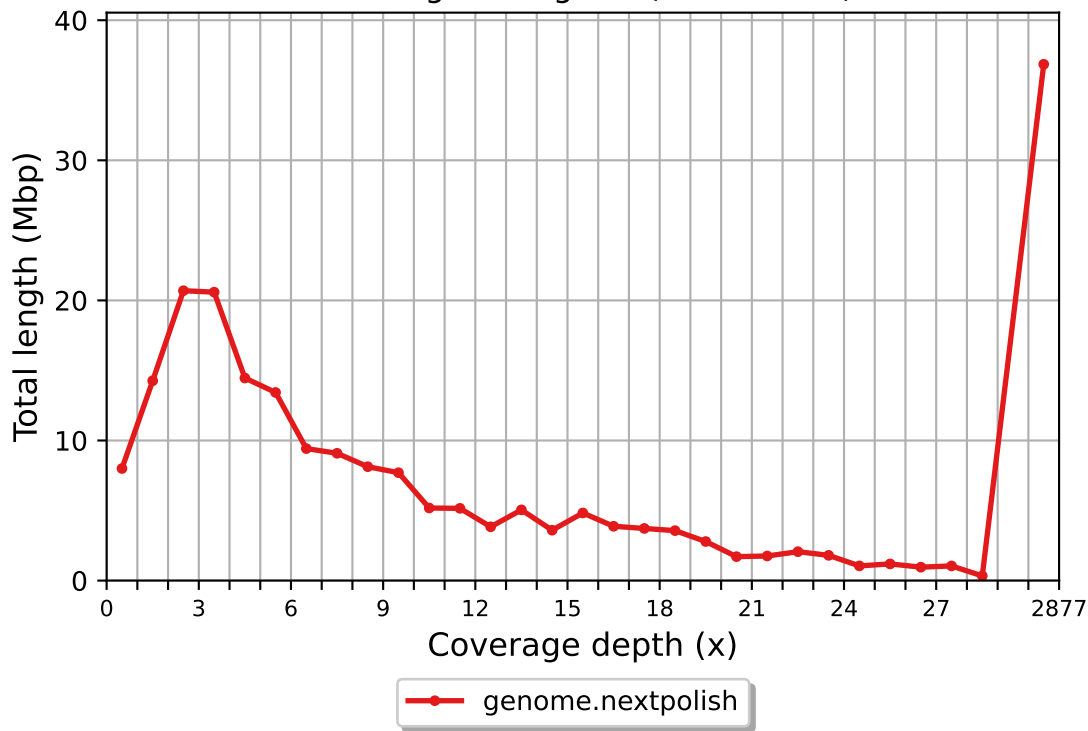
GC content



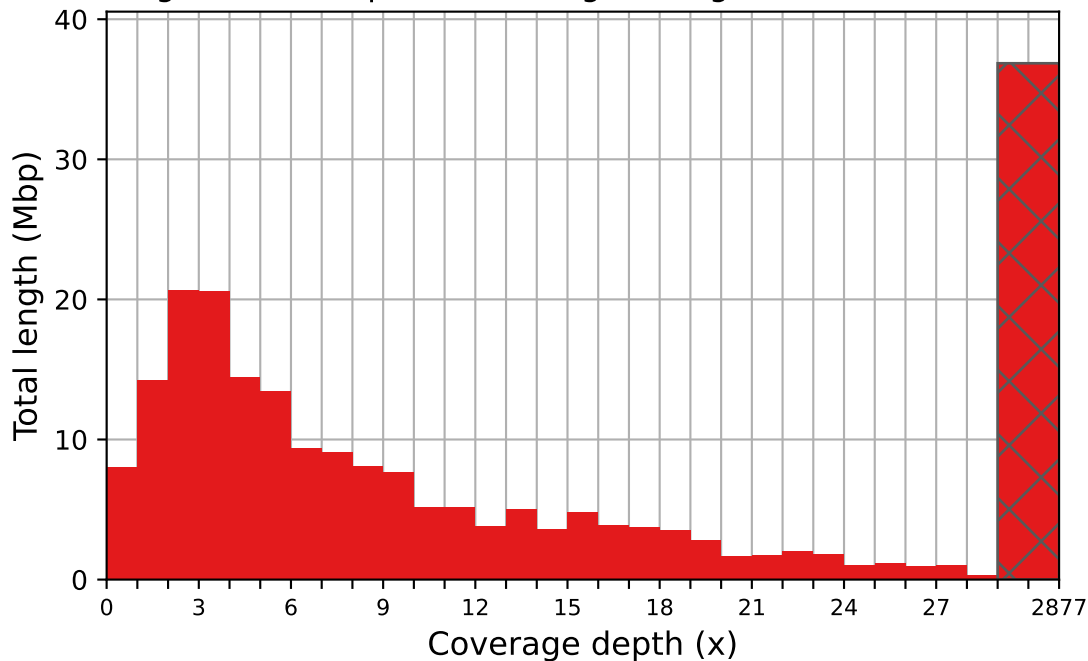
genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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



genome.nextpolish