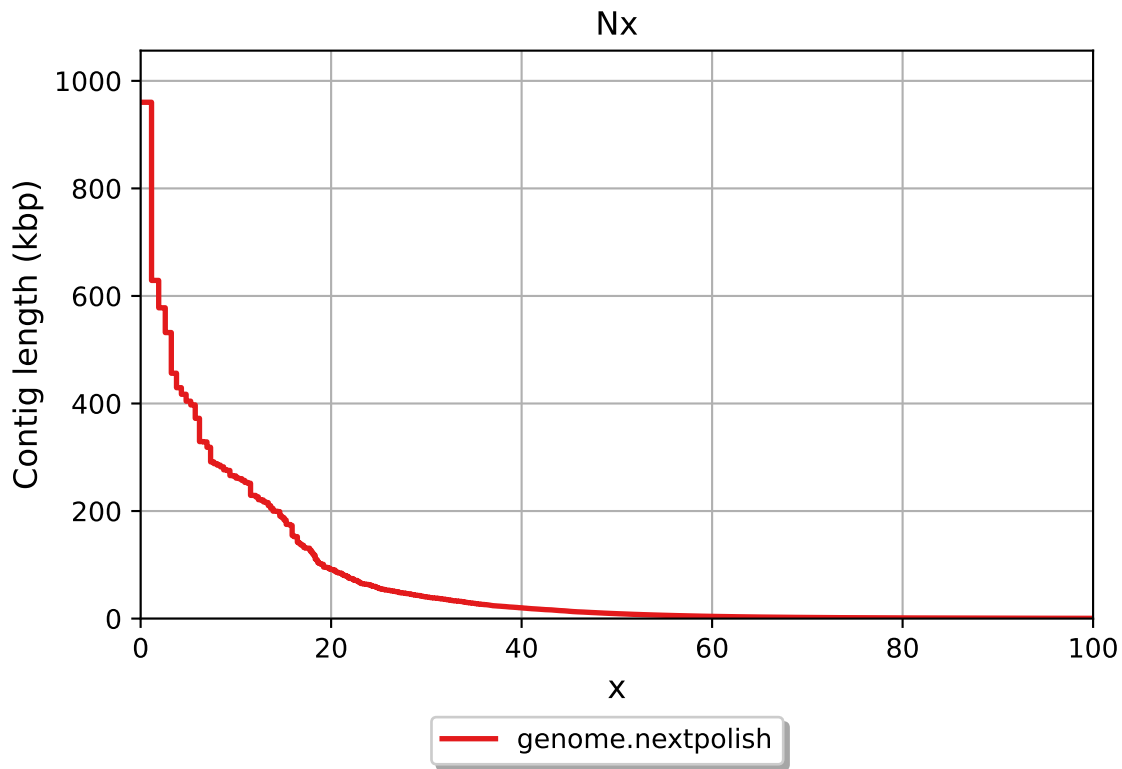


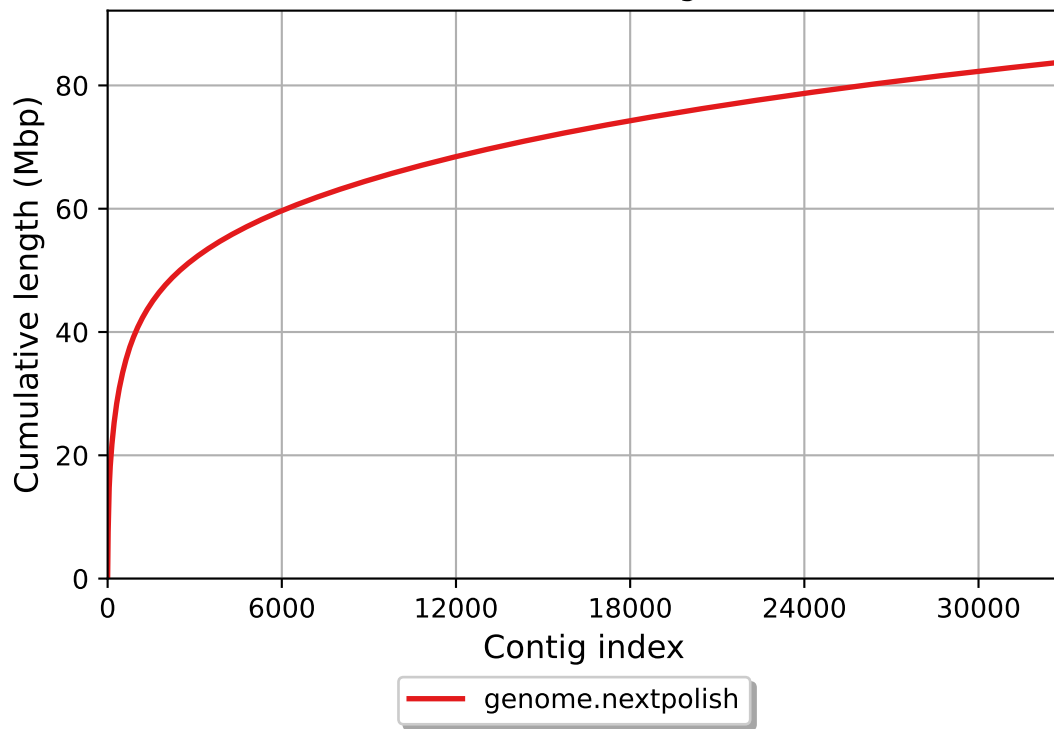
Report

	genome.nextpolish
# contigs (>= 0 bp)	89644
# contigs (>= 1000 bp)	14338
# contigs (>= 5000 bp)	2112
# contigs (>= 10000 bp)	1055
# contigs (>= 25000 bp)	394
# contigs (>= 50000 bp)	161
Total length (>= 0 bp)	100362971
Total length (>= 1000 bp)	70947123
Total length (>= 5000 bp)	48264985
Total length (>= 10000 bp)	40953888
Total length (>= 25000 bp)	30672515
Total length (>= 50000 bp)	22482075
# contigs	32810
Largest contig	960198
Total length	83756838
GC (%)	44.83
N50	9130
N90	783
auN	76012.3
L50	1152
L90	19362
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	24 + 15 part

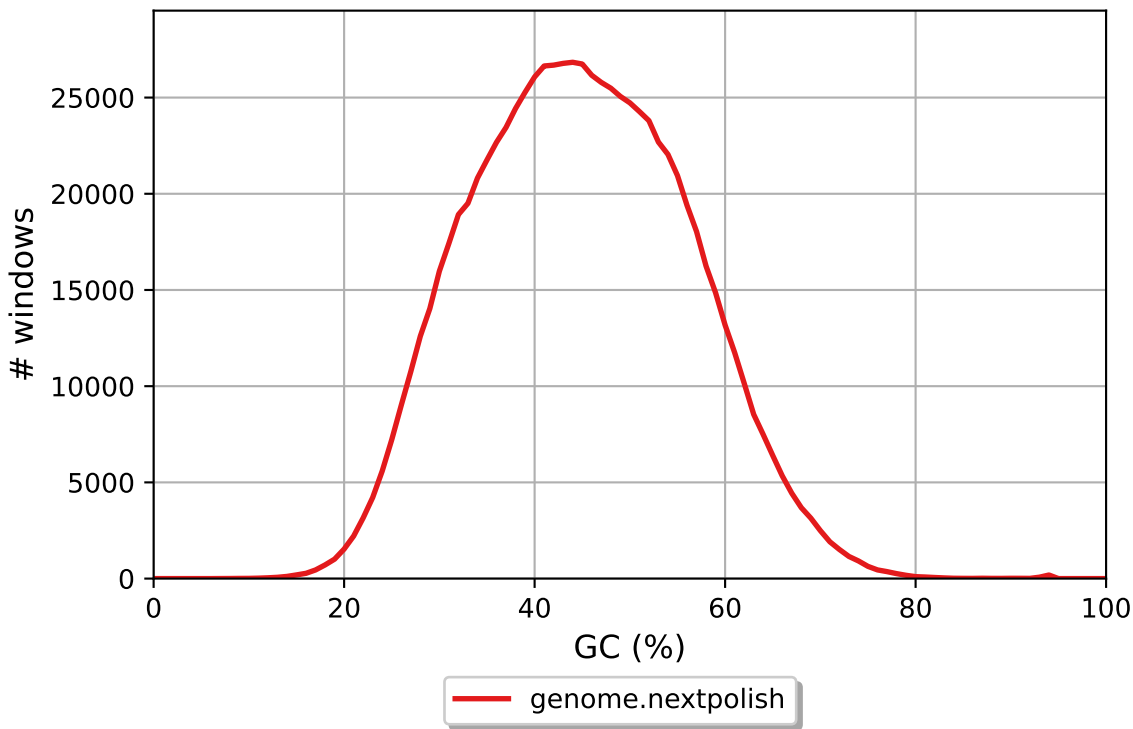
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



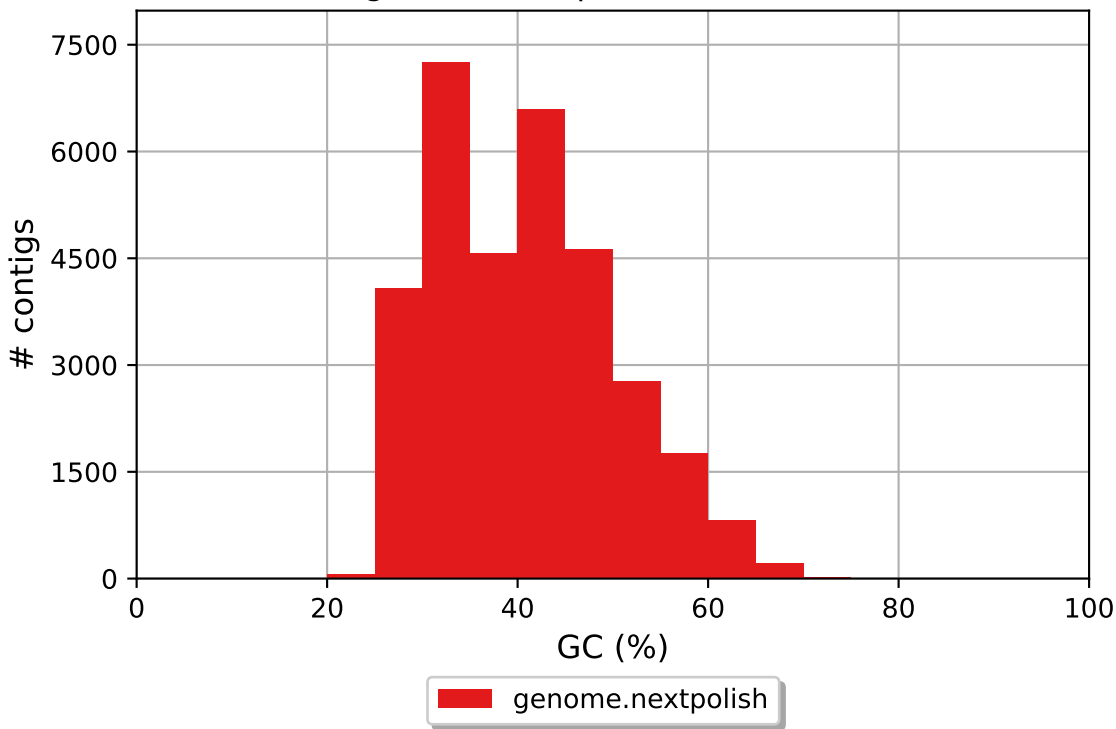
Cumulative length



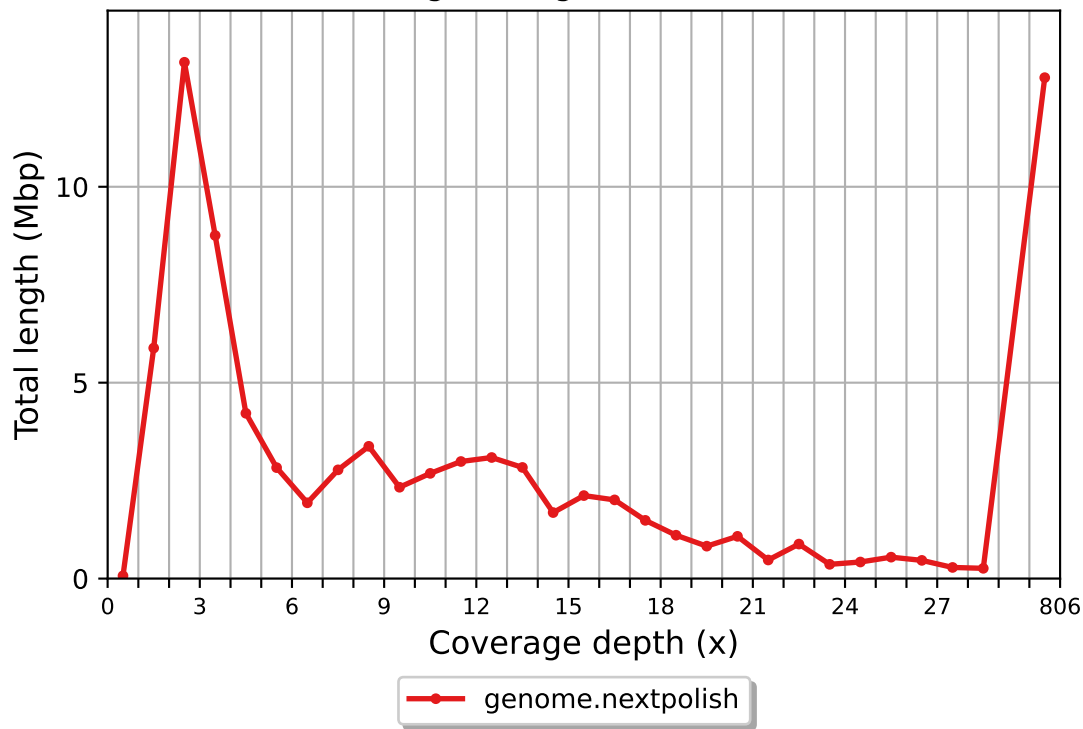
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

