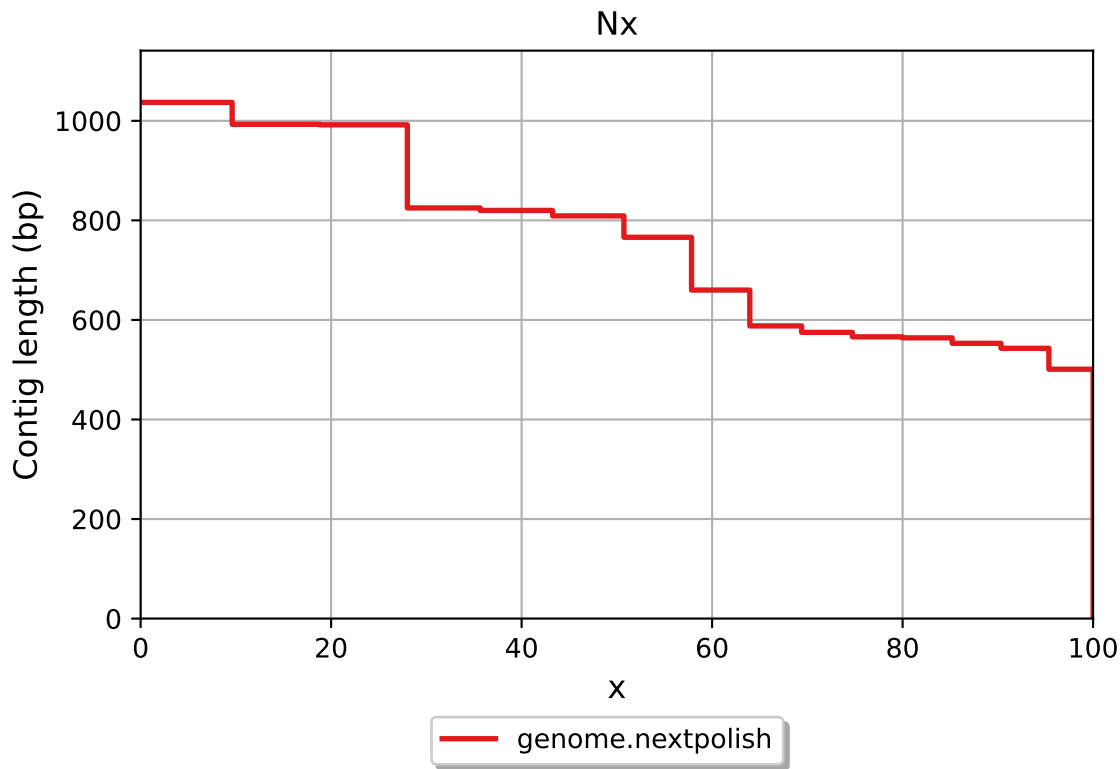


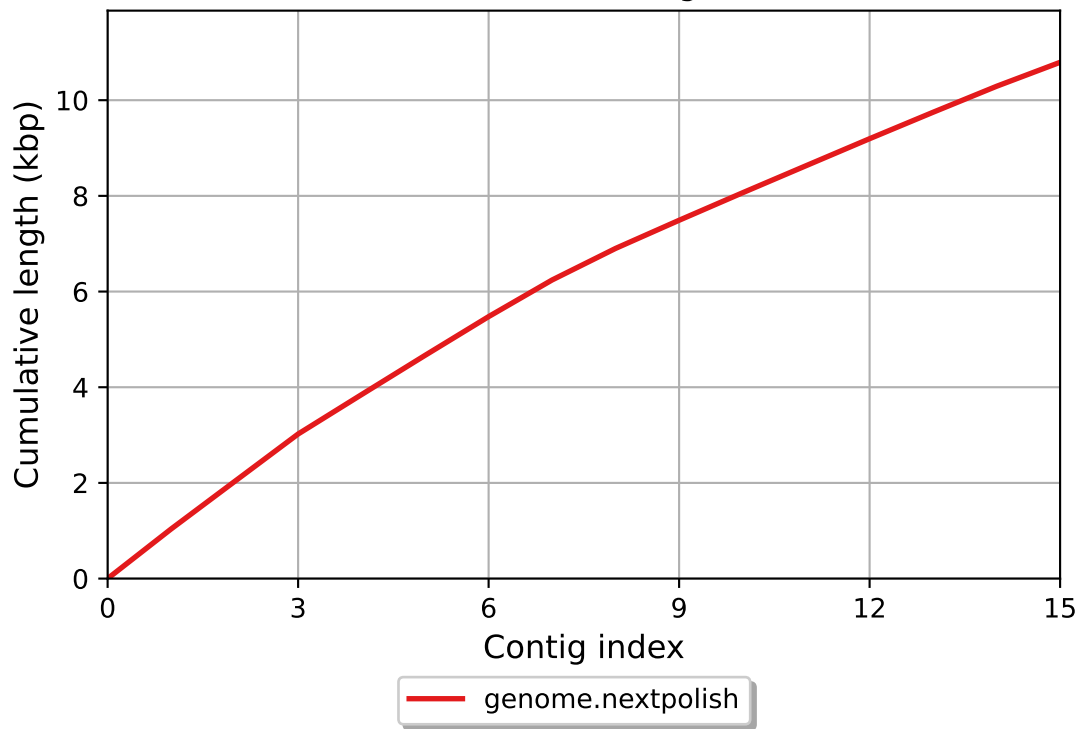
Report

	genome.nextpolish
# contigs (≥ 0 bp)	692
# contigs (≥ 1000 bp)	1
# contigs (≥ 5000 bp)	0
# contigs (≥ 10000 bp)	0
# contigs (≥ 25000 bp)	0
# contigs (≥ 50000 bp)	0
Total length (≥ 0 bp)	126593
Total length (≥ 1000 bp)	1037
Total length (≥ 5000 bp)	0
Total length (≥ 10000 bp)	0
Total length (≥ 25000 bp)	0
Total length (≥ 50000 bp)	0
# contigs	15
Largest contig	1037
Total length	10792
GC (%)	53.82
N50	809
N90	553
auN	763.7
L50	6
L90	13
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

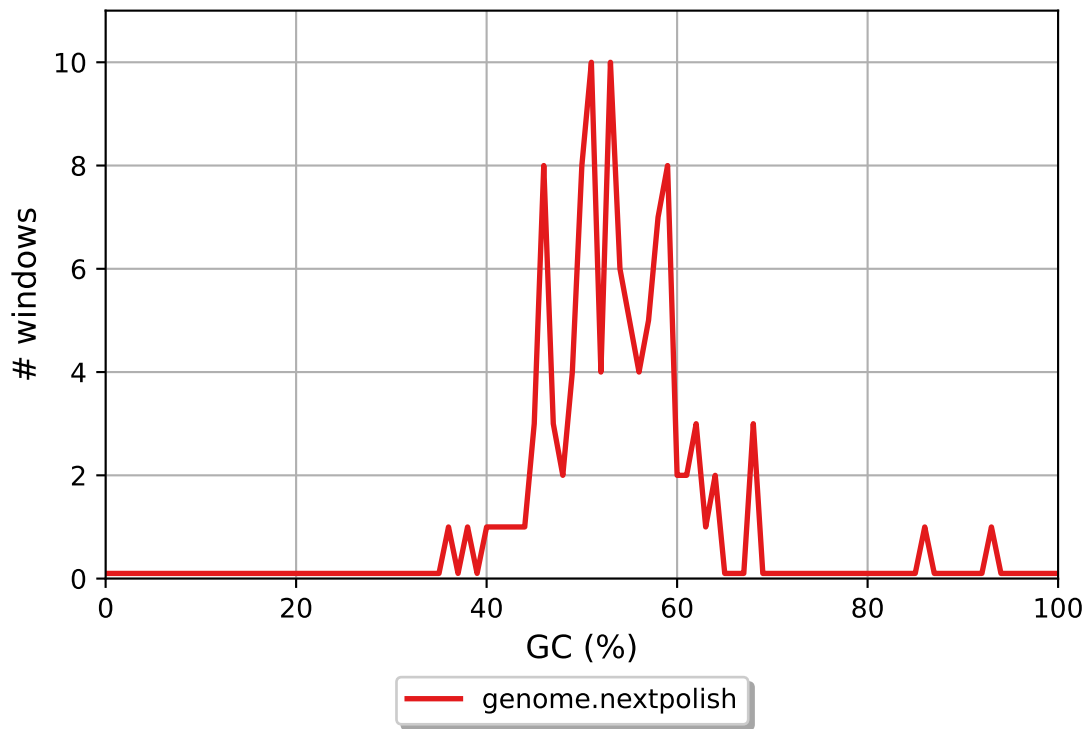
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).



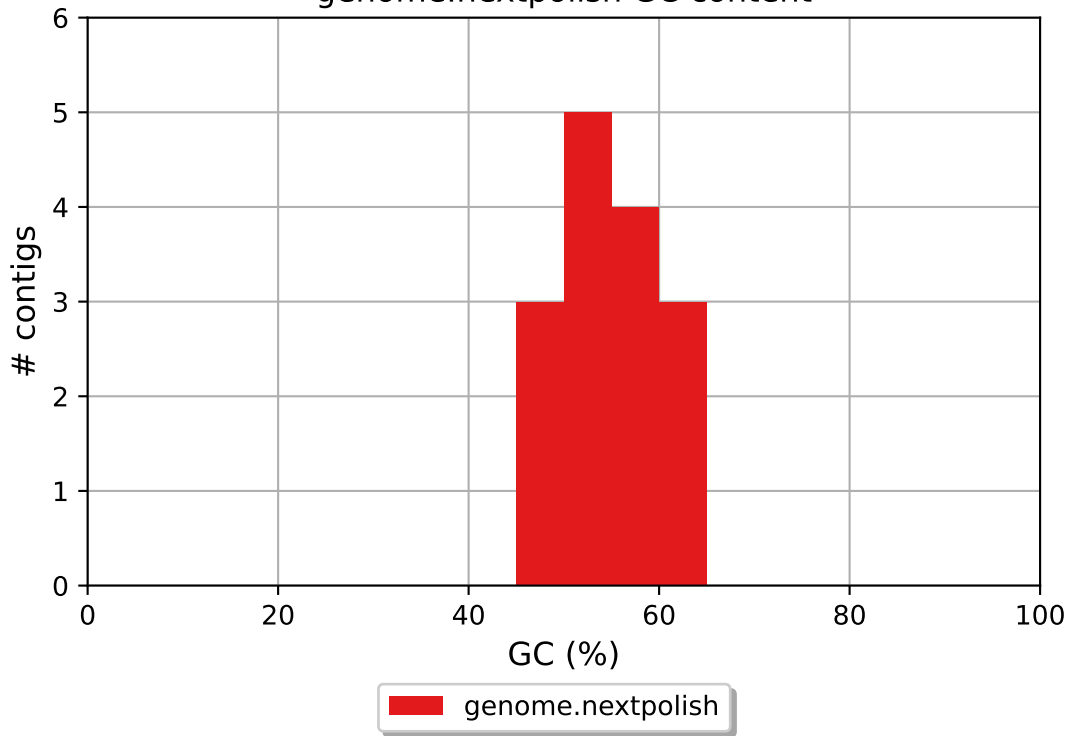
Cumulative length



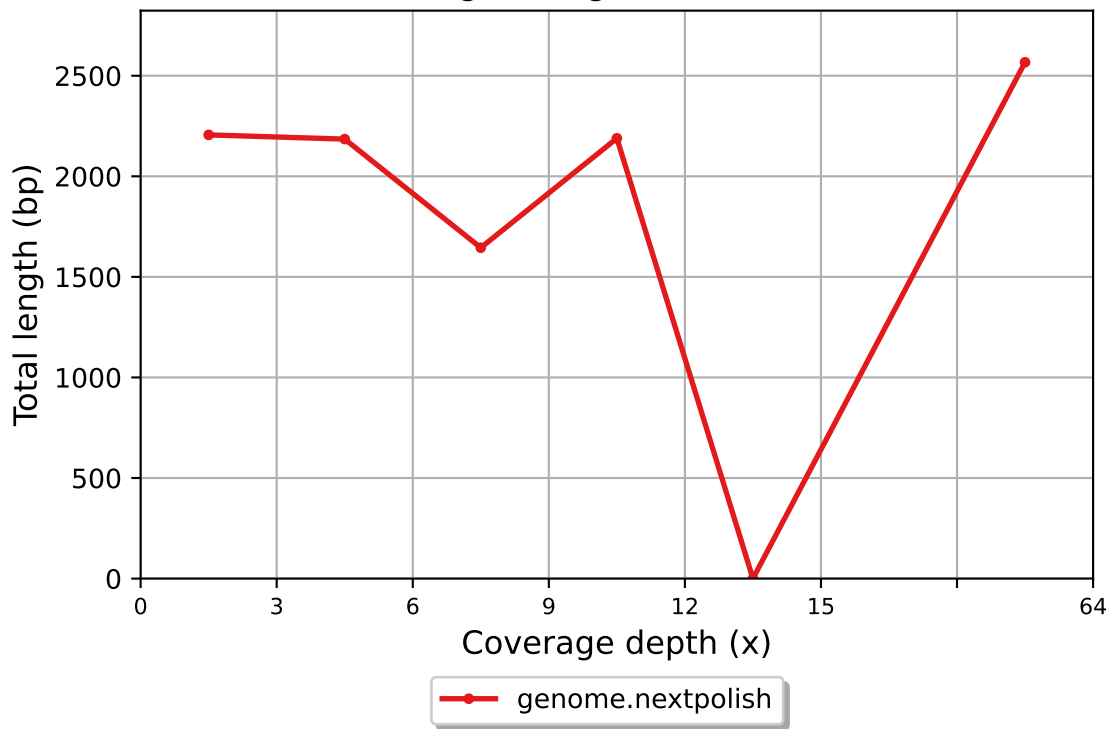
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 3x)



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

