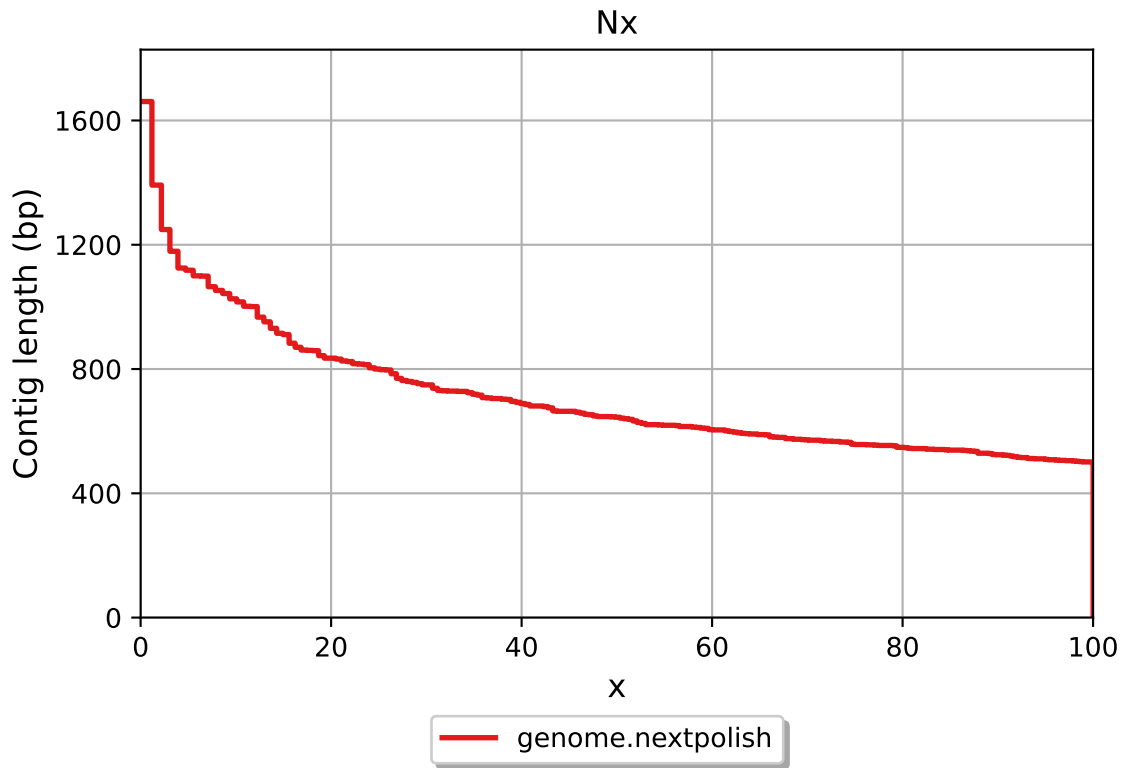
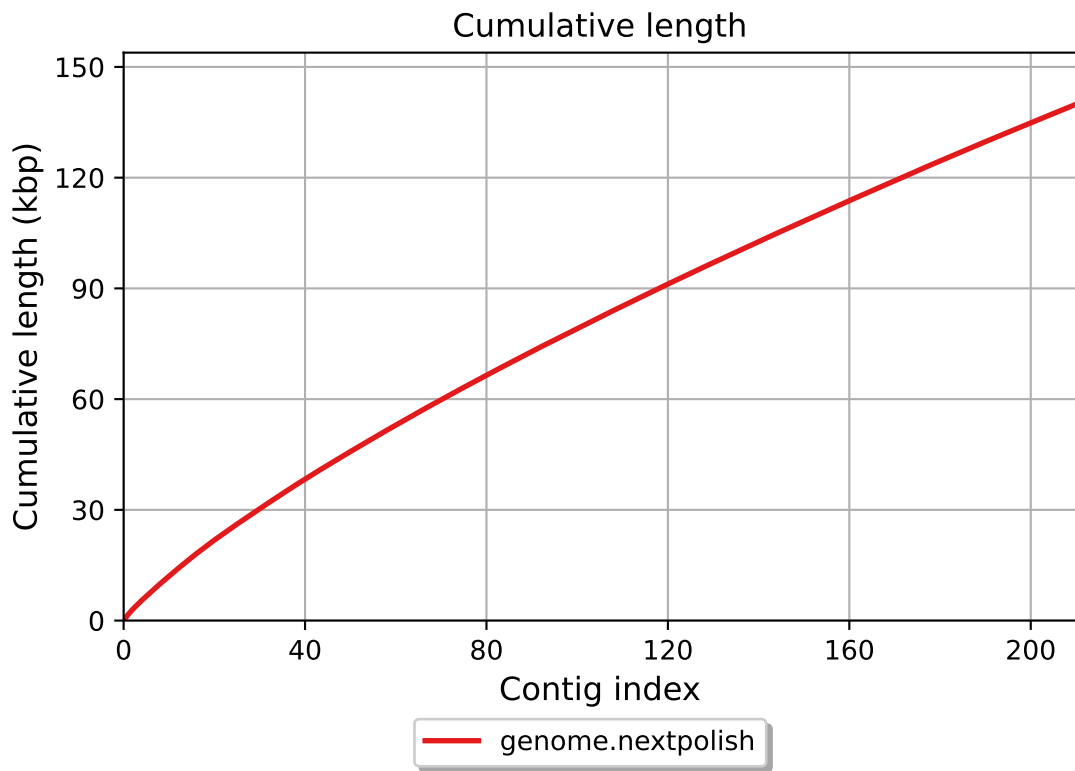


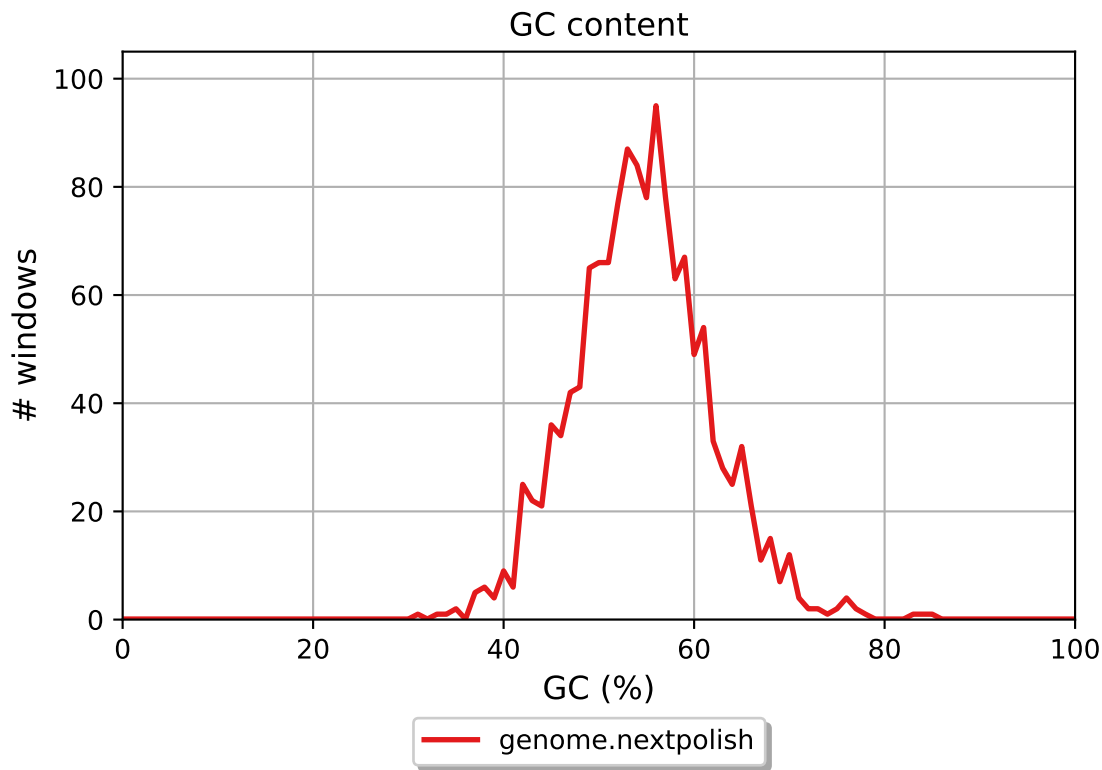
Report

	genome.nextpolish
# contigs (>= 0 bp)	4082
# contigs (>= 1000 bp)	15
# contigs (>= 5000 bp)	0
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	901656
Total length (>= 1000 bp)	17129
Total length (>= 5000 bp)	0
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	210
Largest contig	1661
Total length	139883
GC (%)	54.61
N50	645
N90	524
auN	711.9
L50	86
L90	183
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 1 part

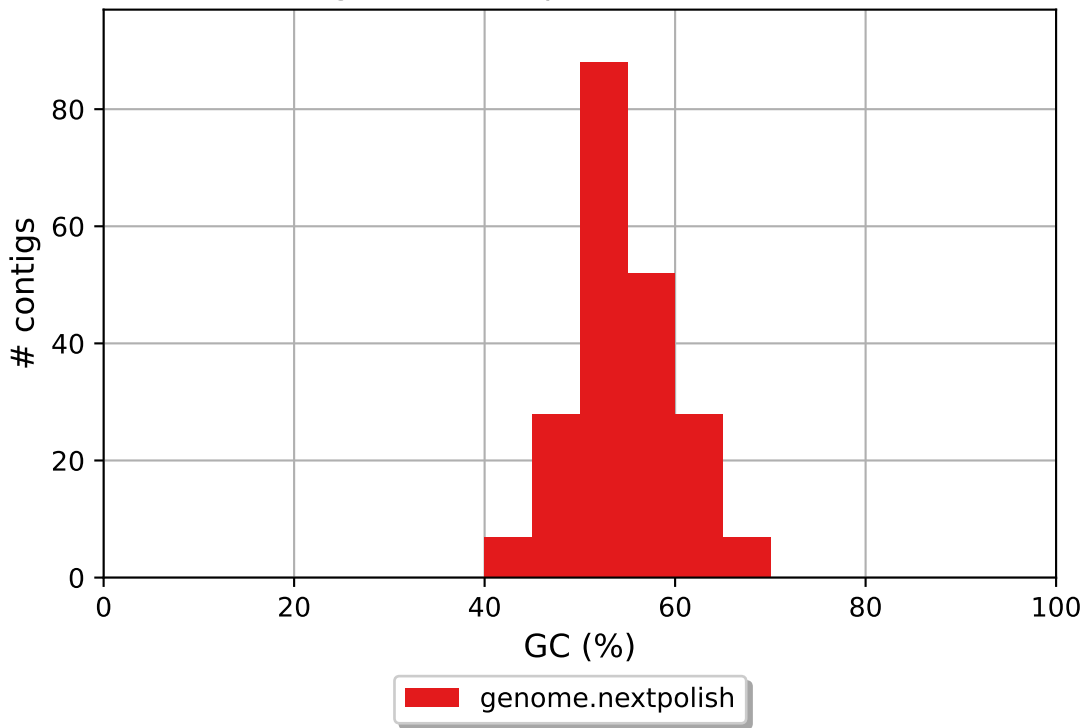
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



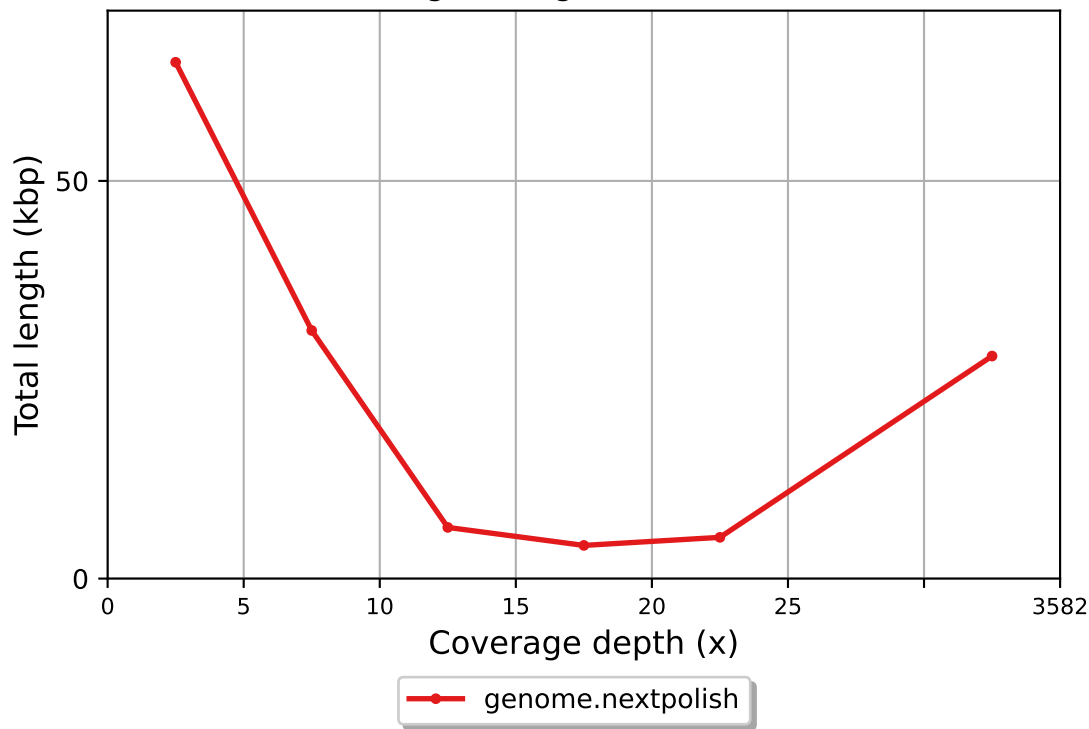




genome.nextpolish GC content



Coverage histogram (bin size: 5x)



genome.nextpolish coverage histogram (bin size: 5x)

