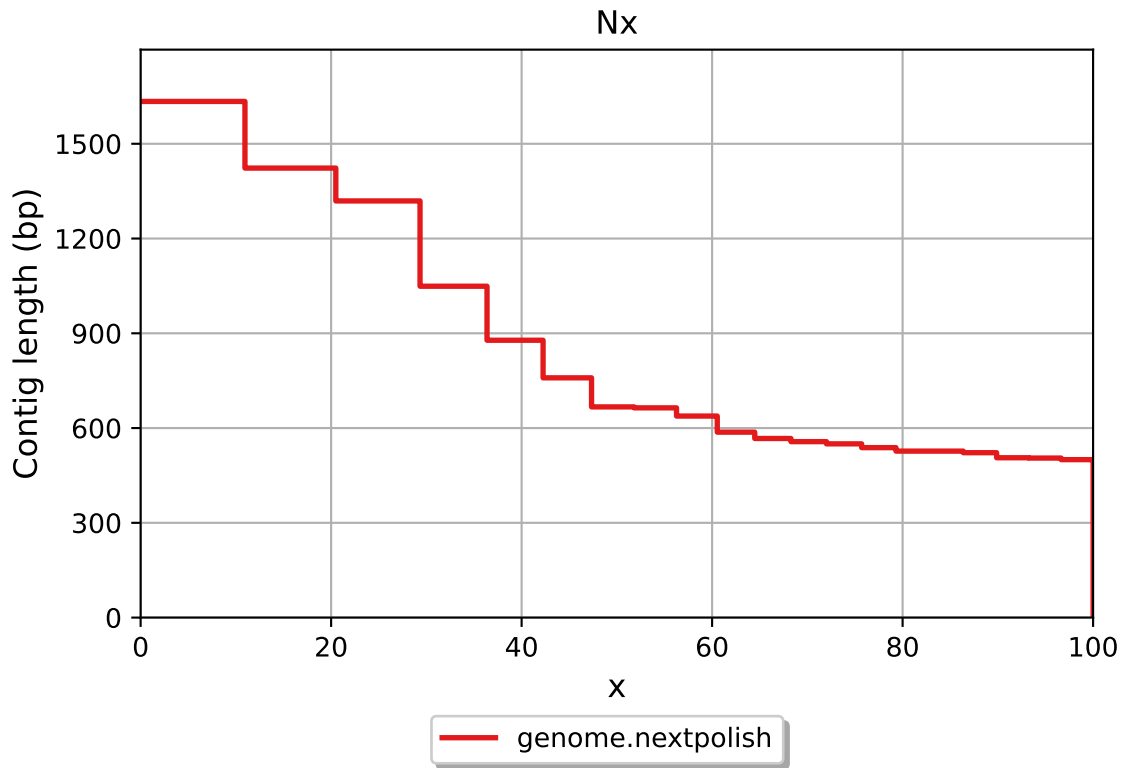


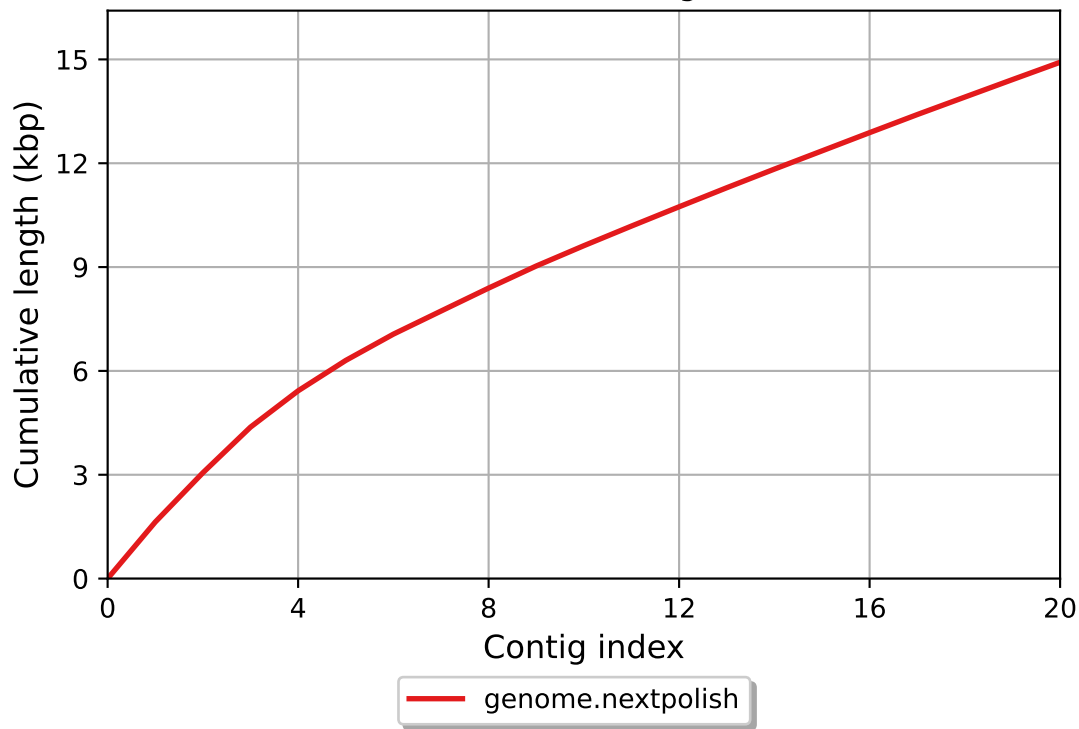
Report

	genome.nextpolish
# contigs (>= 0 bp)	795
# contigs (>= 1000 bp)	4
# contigs (>= 5000 bp)	0
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	167164
Total length (>= 1000 bp)	5425
Total length (>= 5000 bp)	0
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	20
Largest contig	1634
Total length	14917
GC (%)	54.40
N50	667
N90	506
auN	893.8
L50	7
L90	18
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

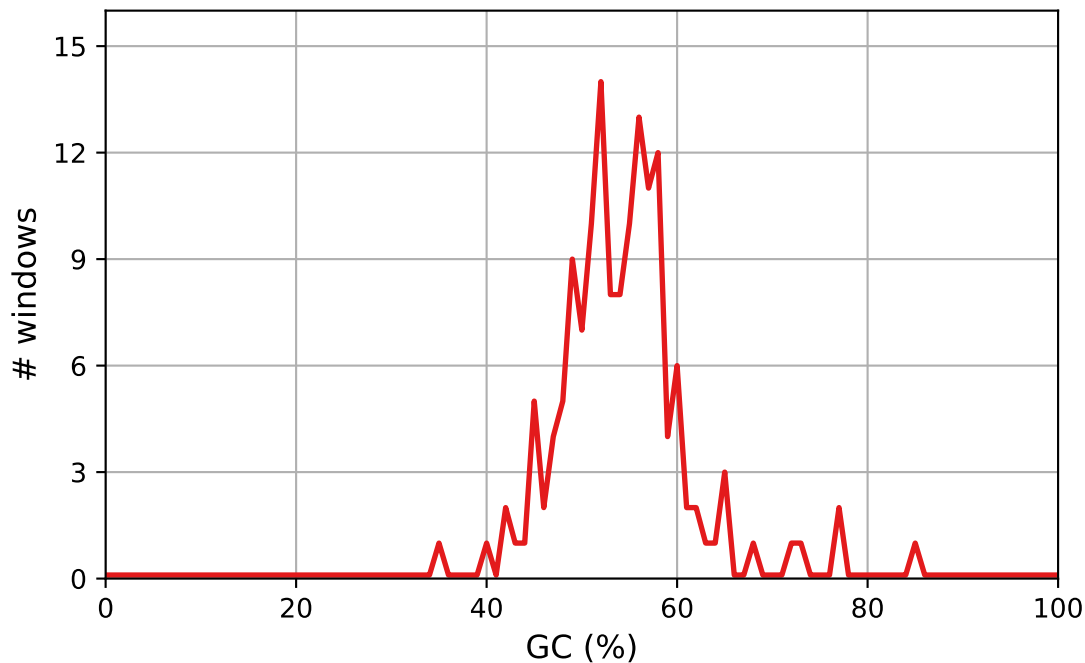
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length

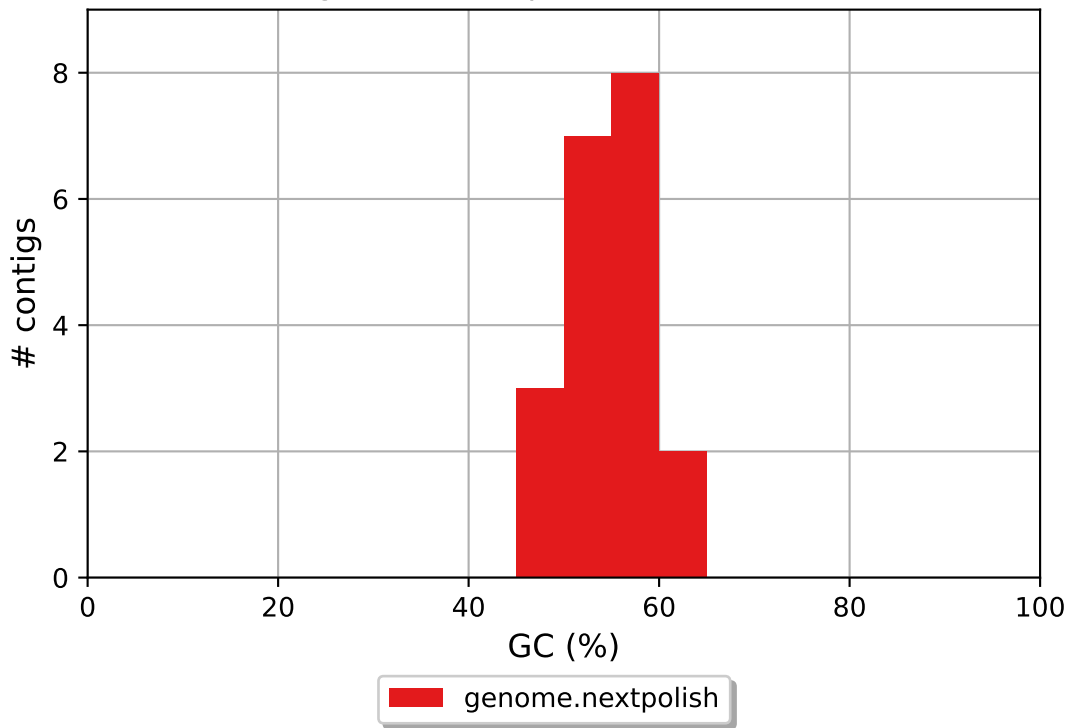


GC content

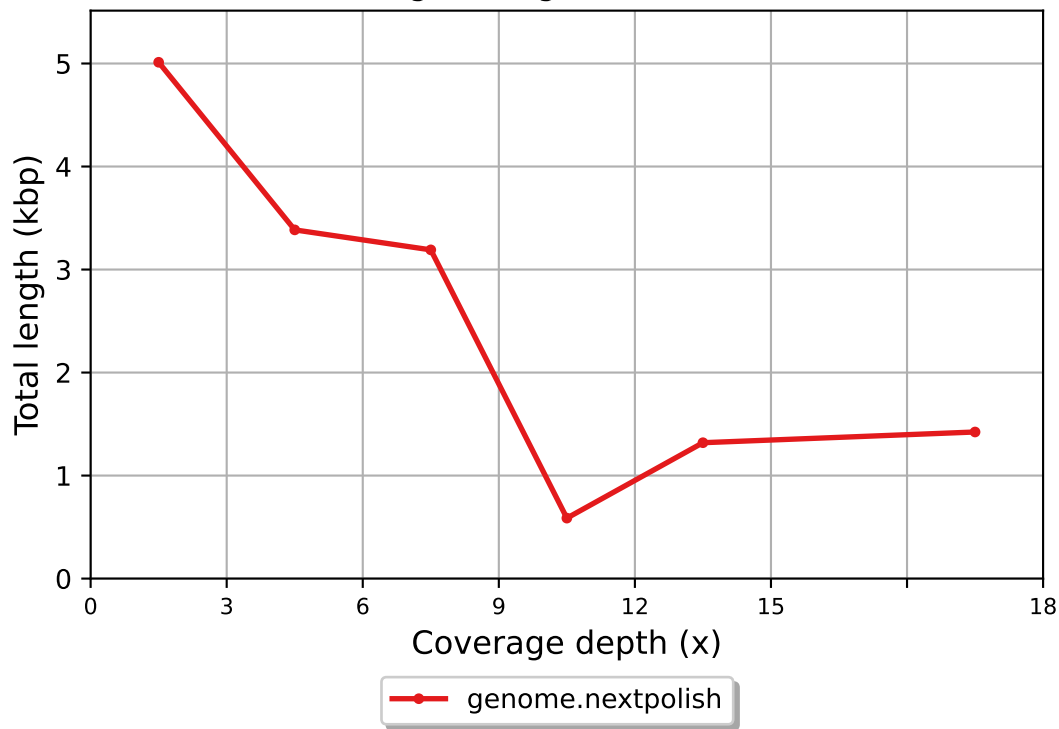


— genome.nextpolish

genome.nextpolish GC content



Coverage histogram (bin size: 3x)



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

