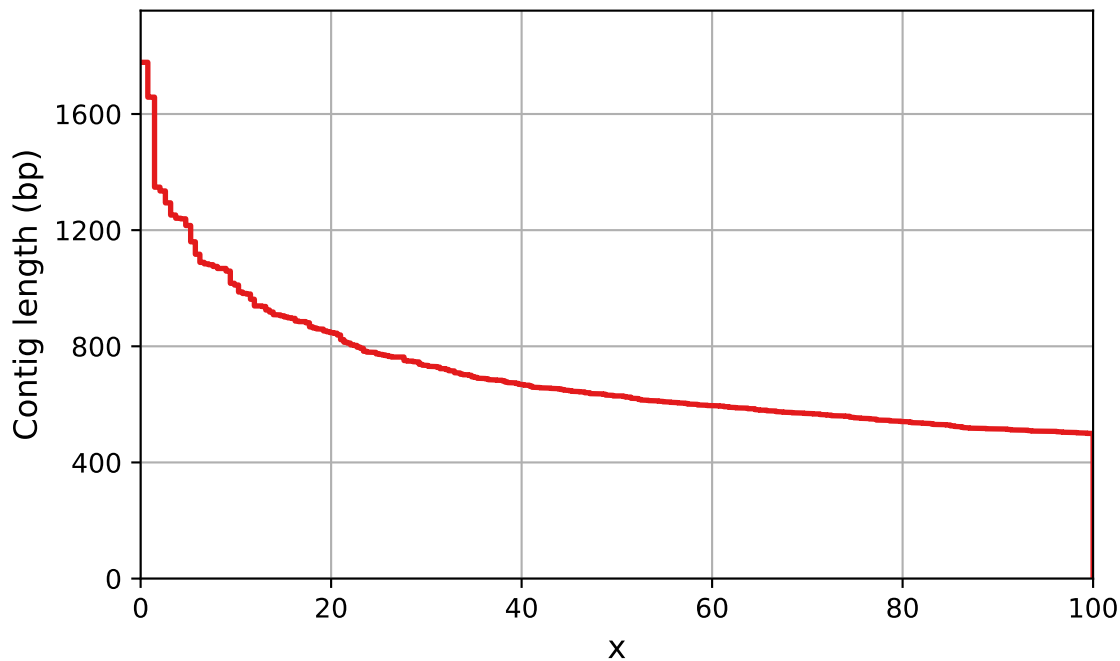


Report

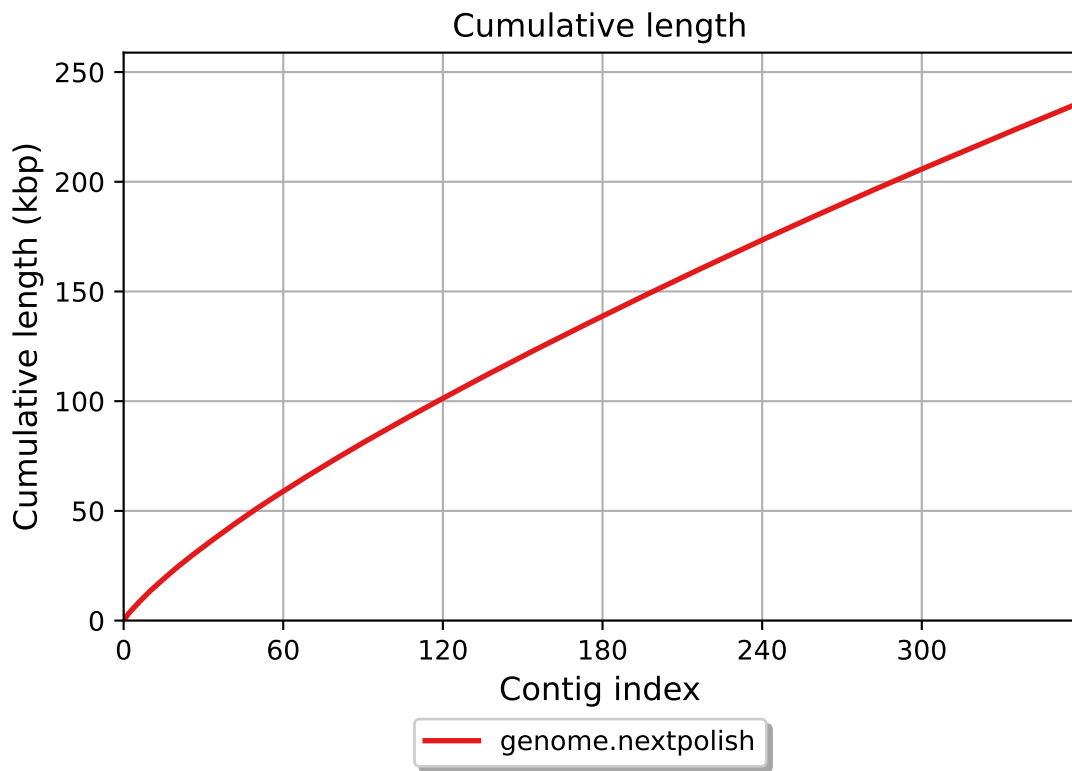
	genome.nextpolish
# contigs (>= 0 bp)	7958
# contigs (>= 1000 bp)	20
# contigs (>= 5000 bp)	0
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1663051
Total length (>= 1000 bp)	24190
Total length (>= 5000 bp)	0
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	358
Largest contig	1778
Total length	235317
GC (%)	54.23
N50	629
N90	515
auN	706.7
L50	146
L90	312
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

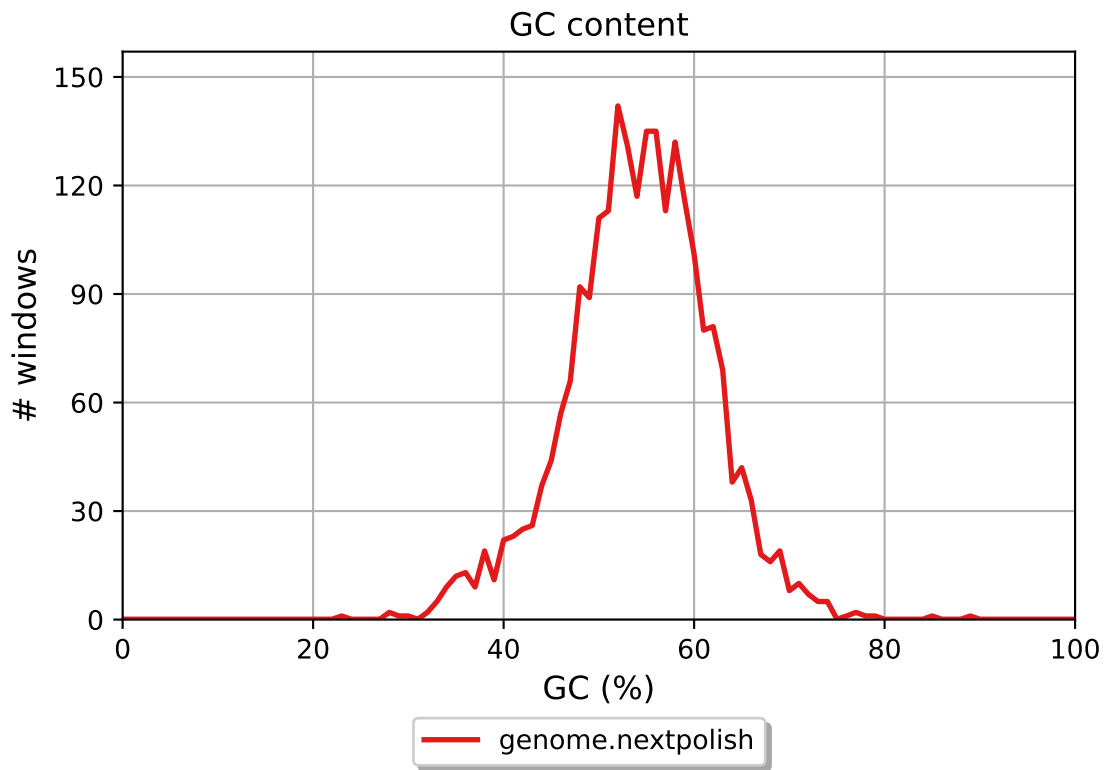
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

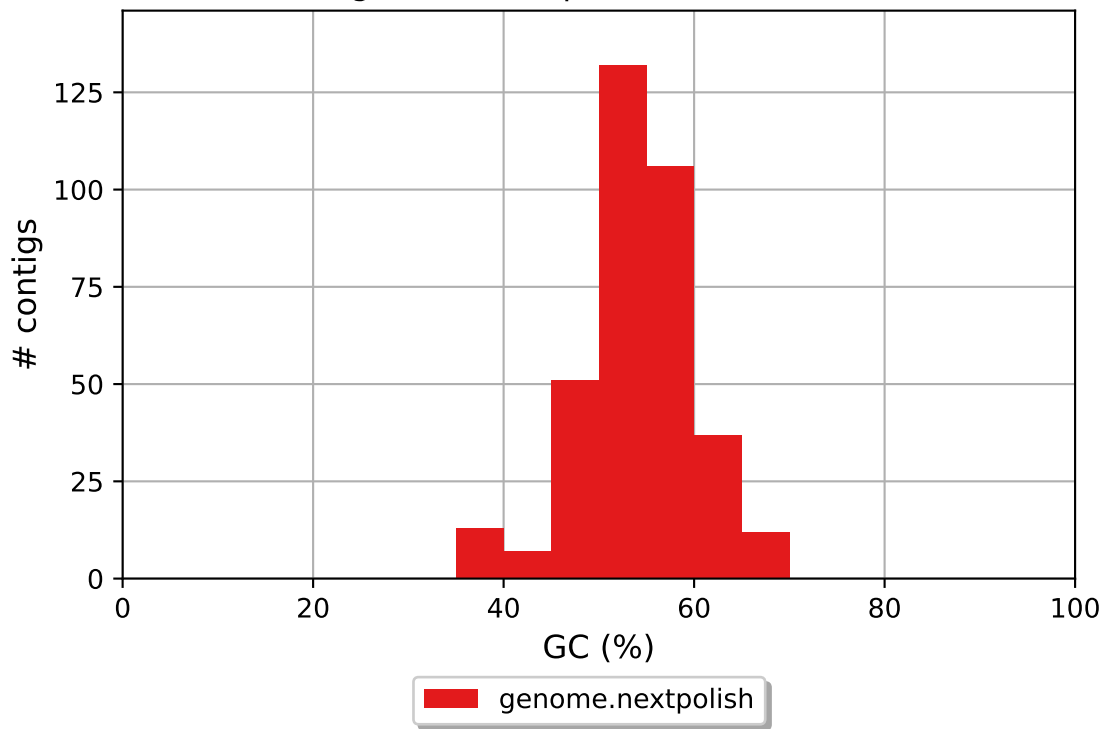


— genome.nextpolish

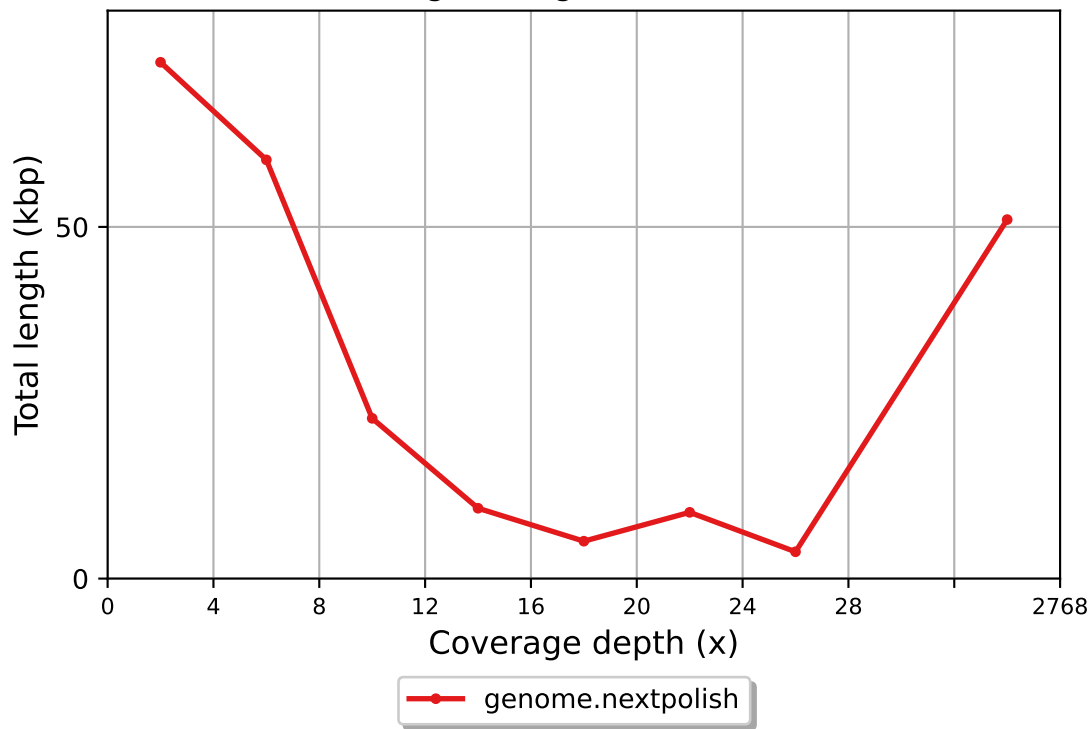




genome.nextpolish GC content



Coverage histogram (bin size: 4x)



genome.nextpolish coverage histogram (bin size: 4x)

