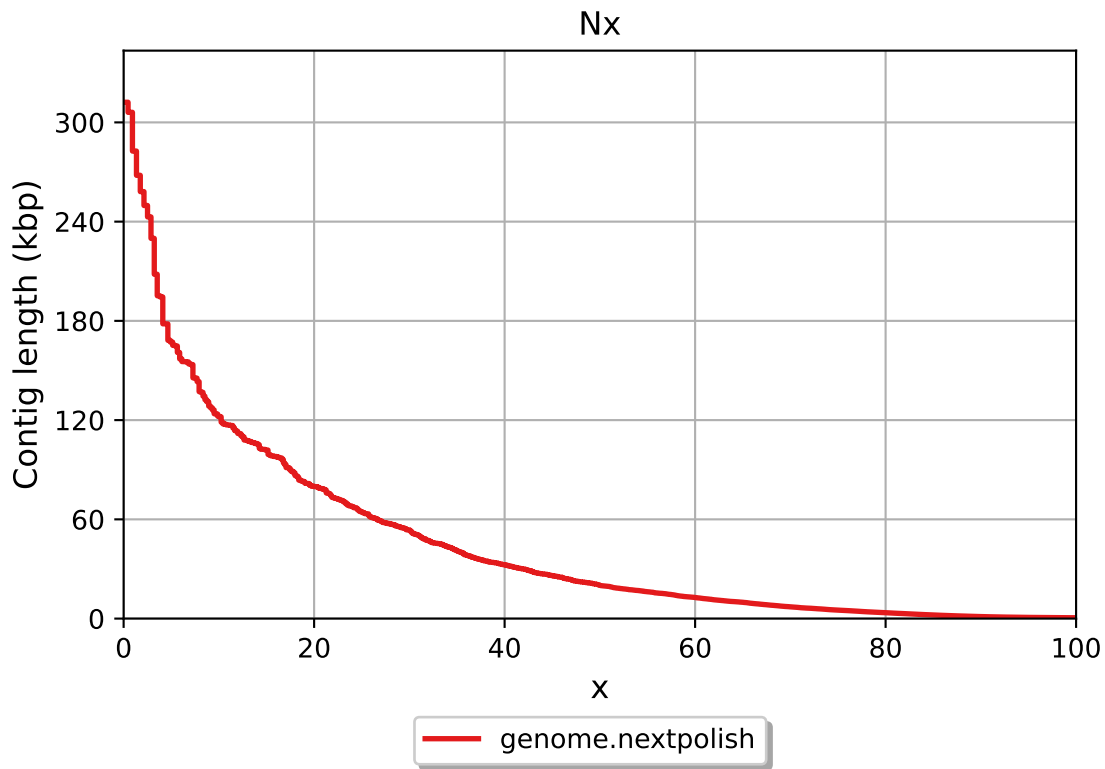


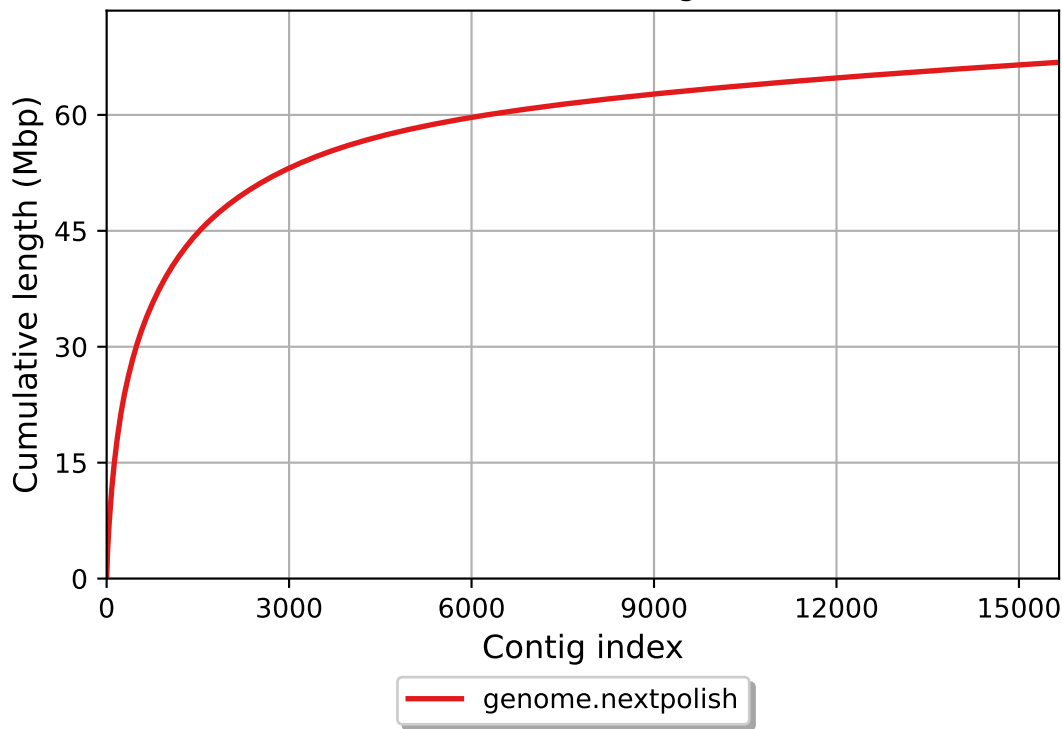
Report

	genome.nextpolish
# contigs (>= 0 bp)	72253
# contigs (>= 1000 bp)	7395
# contigs (>= 5000 bp)	2367
# contigs (>= 10000 bp)	1333
# contigs (>= 25000 bp)	515
# contigs (>= 50000 bp)	223
Total length (>= 0 bp)	83003339
Total length (>= 1000 bp)	61271527
Total length (>= 5000 bp)	50392426
Total length (>= 10000 bp)	43215815
Total length (>= 25000 bp)	30720980
Total length (>= 50000 bp)	20773219
# contigs	15660
Largest contig	312166
Total length	66810656
GC (%)	46.18
N50	20045
N90	1213
auN	46066.3
L50	636
L90	6355
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	25 + 12 part

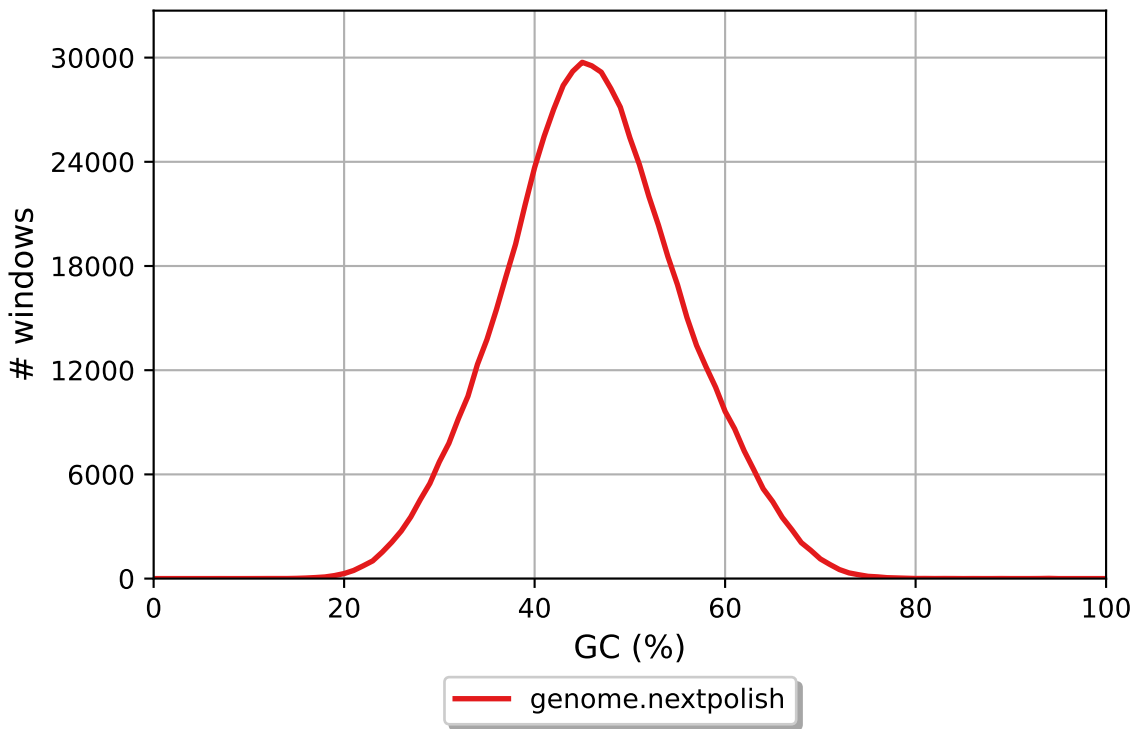
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



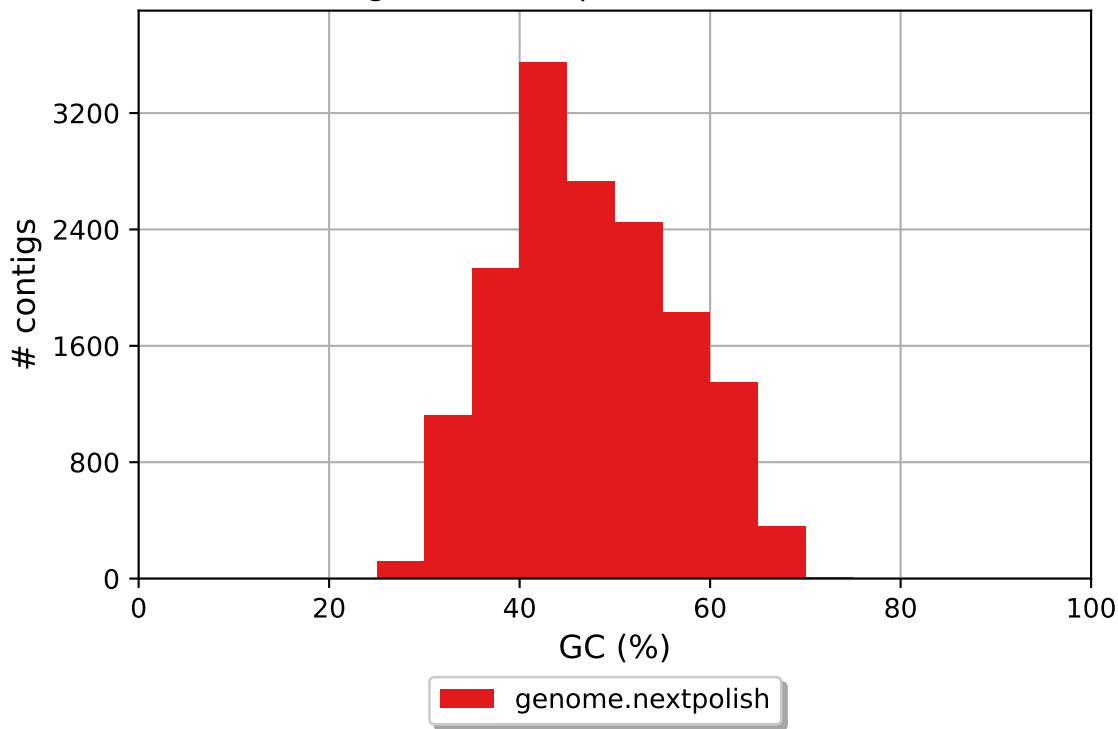
Cumulative length



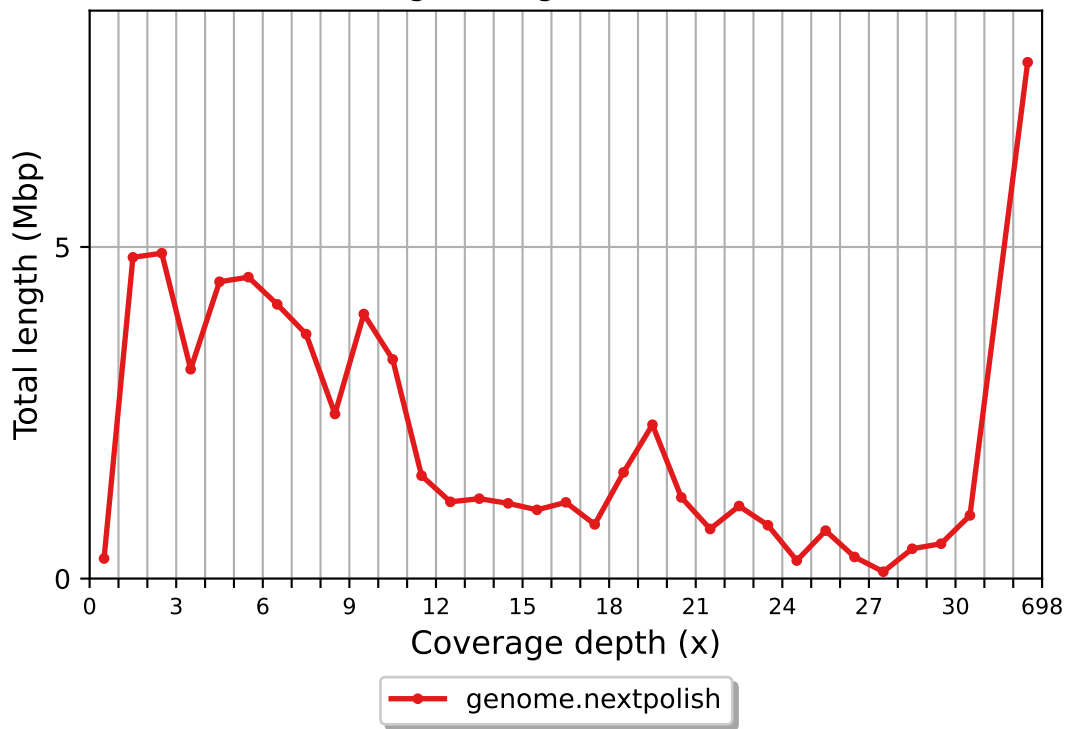
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

