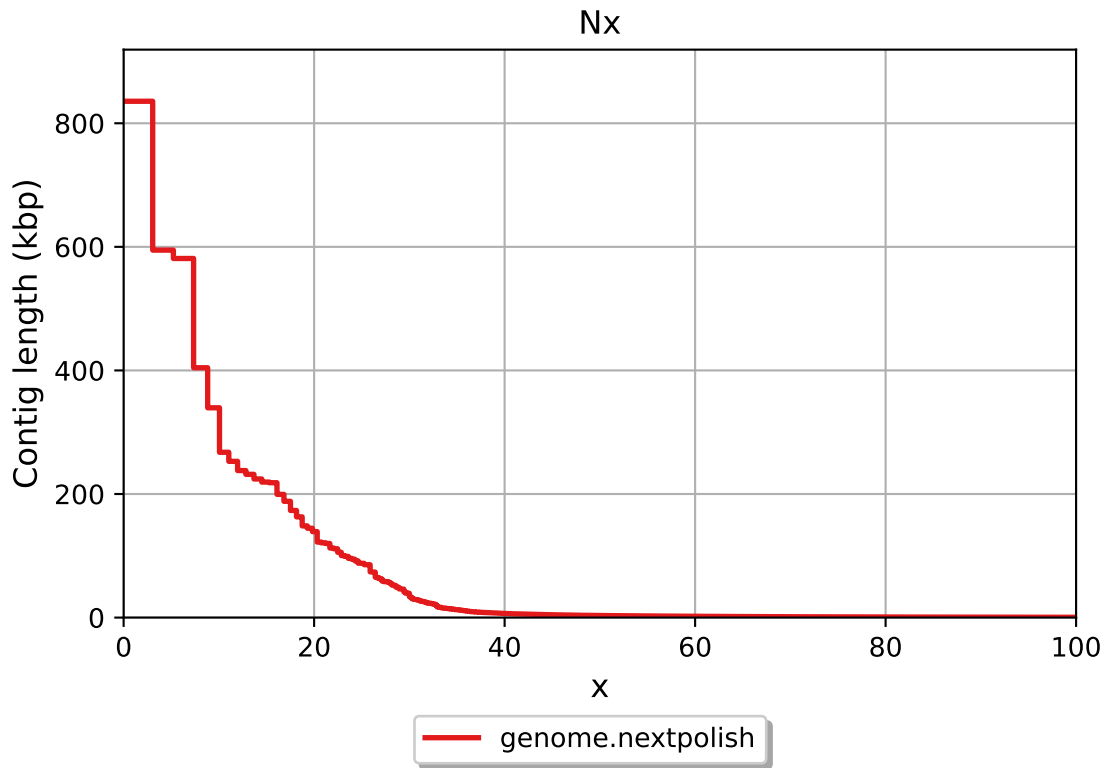


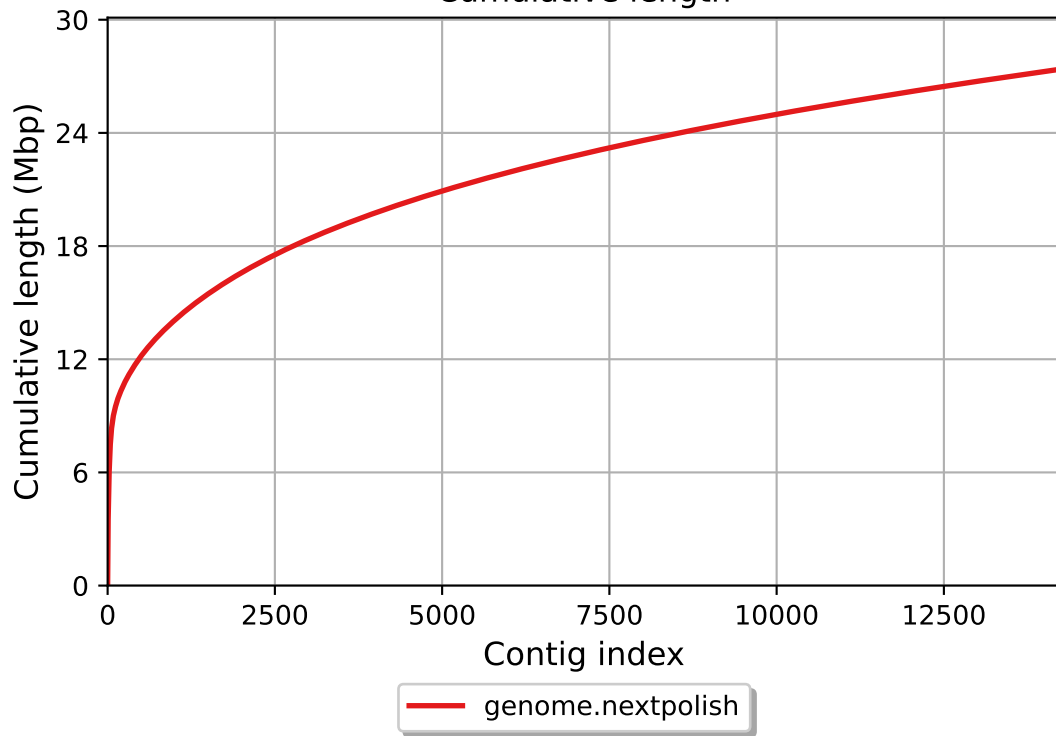
Report

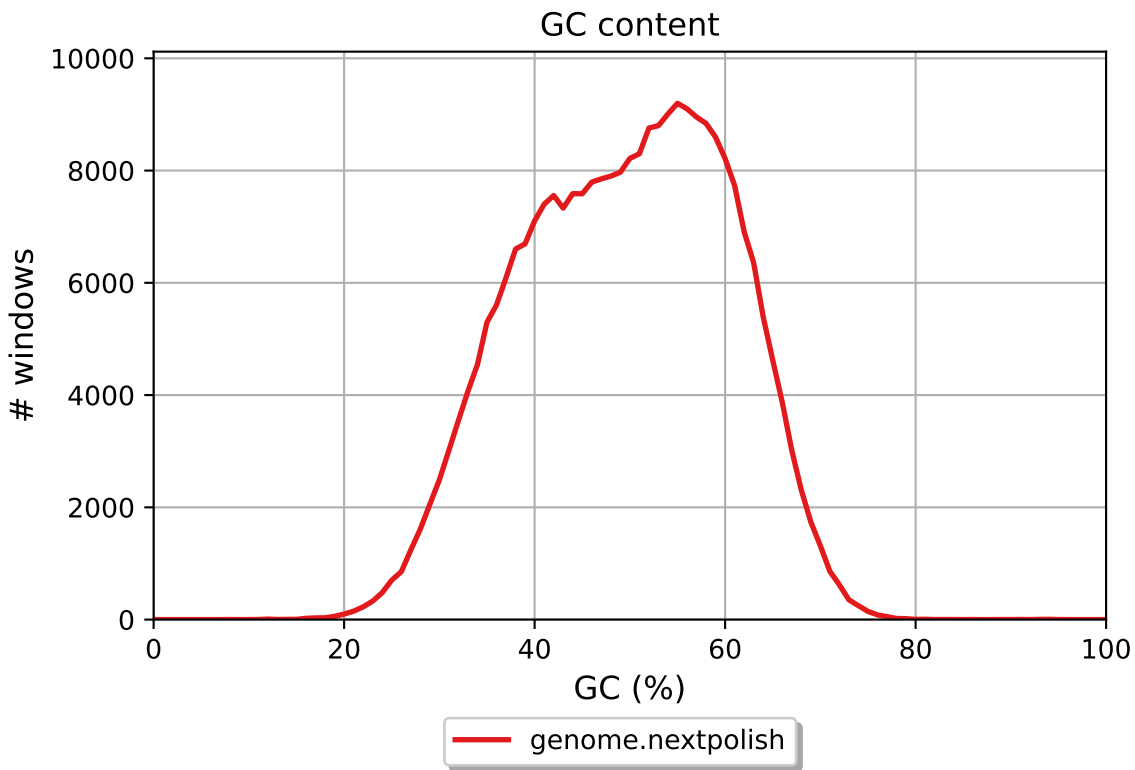
	genome.nextpolish
# contigs (>= 0 bp)	71244
# contigs (>= 1000 bp)	5494
# contigs (>= 5000 bp)	475
# contigs (>= 10000 bp)	156
# contigs (>= 25000 bp)	72
# contigs (>= 50000 bp)	46
Total length (>= 0 bp)	43120481
Total length (>= 1000 bp)	21431087
Total length (>= 5000 bp)	12046865
Total length (>= 10000 bp)	9937529
Total length (>= 25000 bp)	8696051
Total length (>= 50000 bp)	7816665
# contigs	14237
Largest contig	835604
Total length	27373704
GC (%)	49.44
N50	3364
N90	667
auN	93067.9
L50	884
L90	9465
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	21 + 14 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).

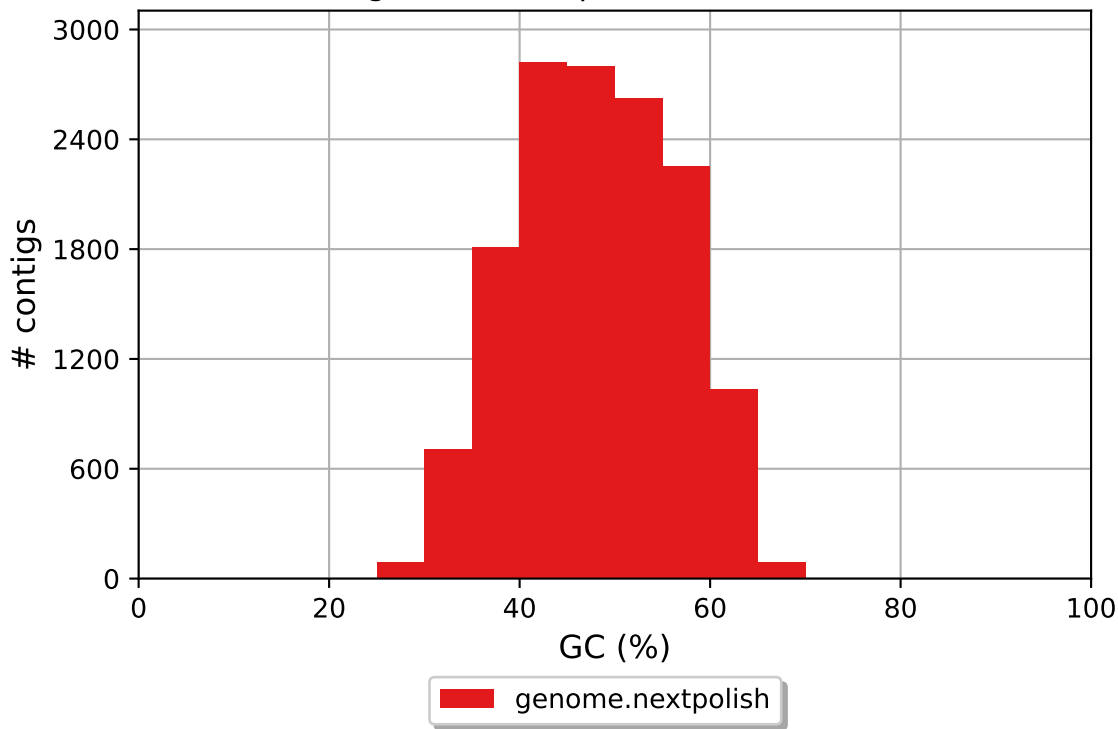


Cumulative length

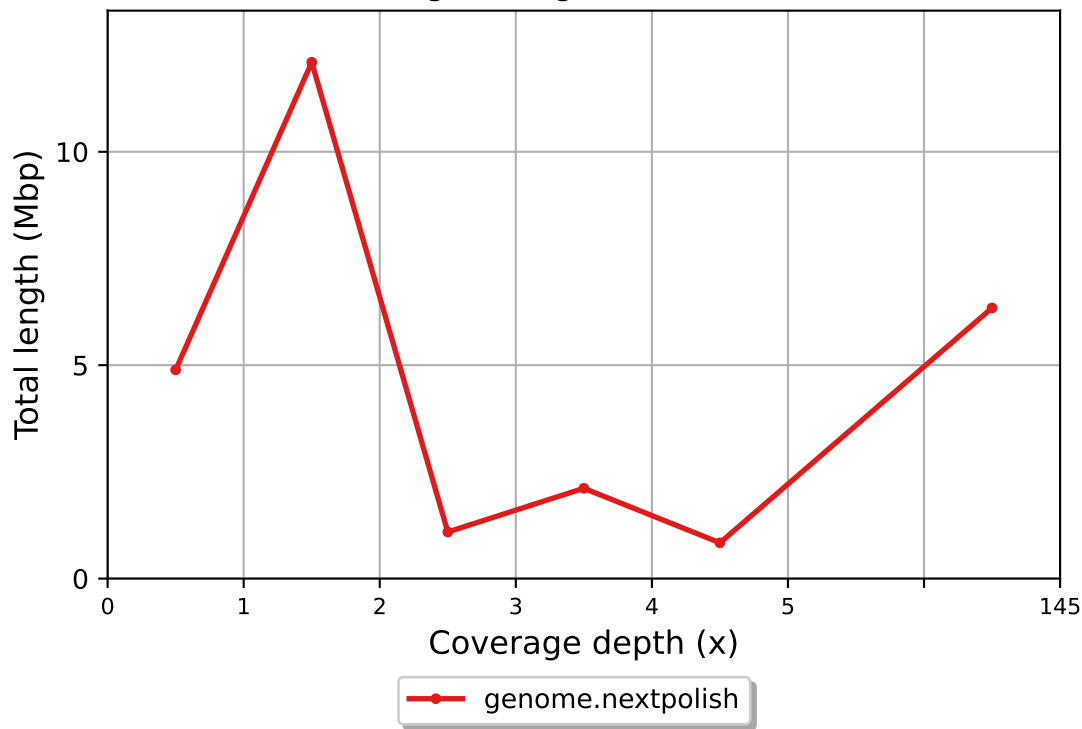




genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

