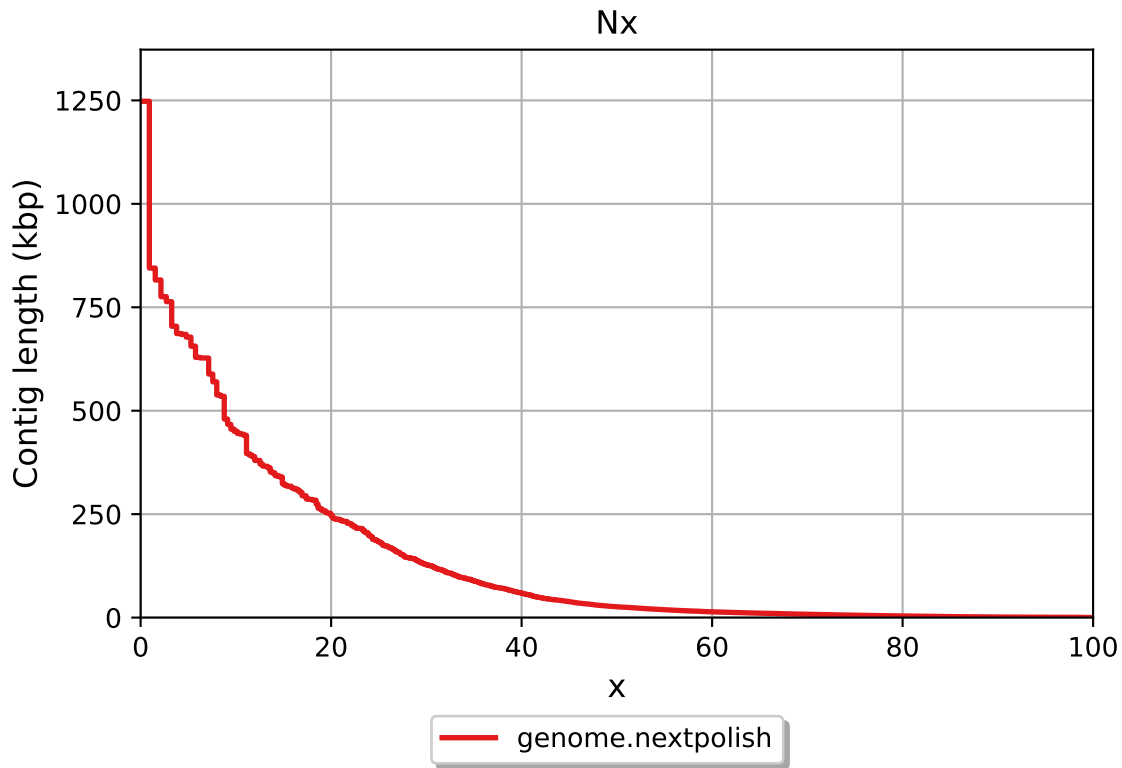


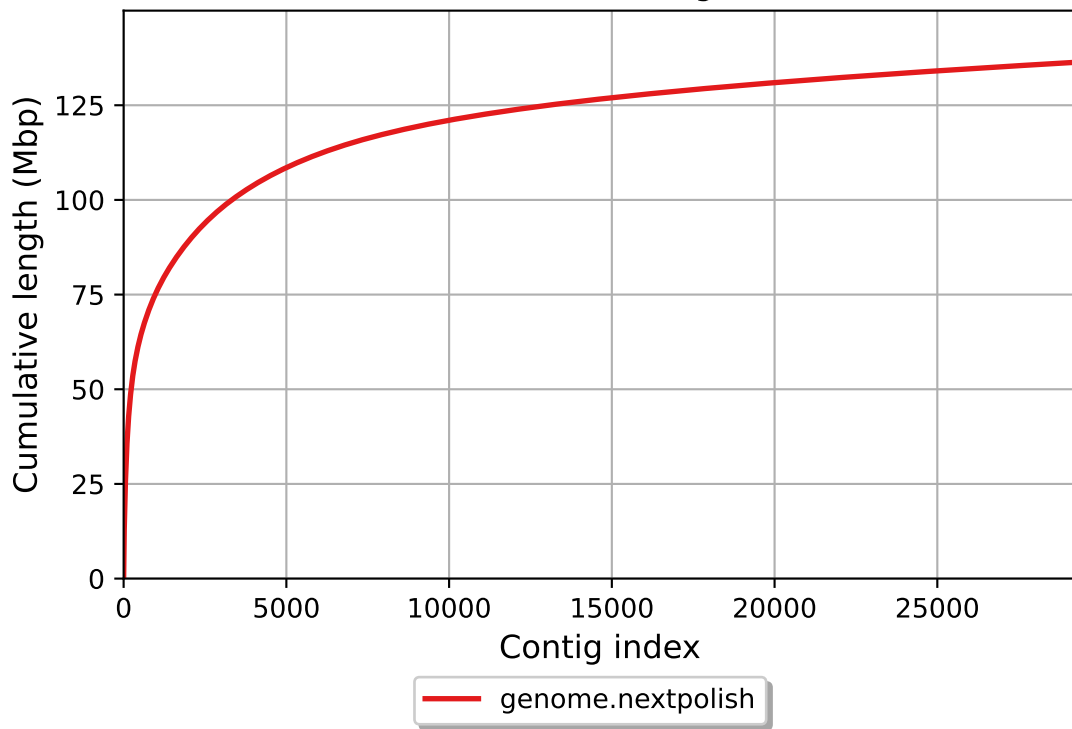
Report

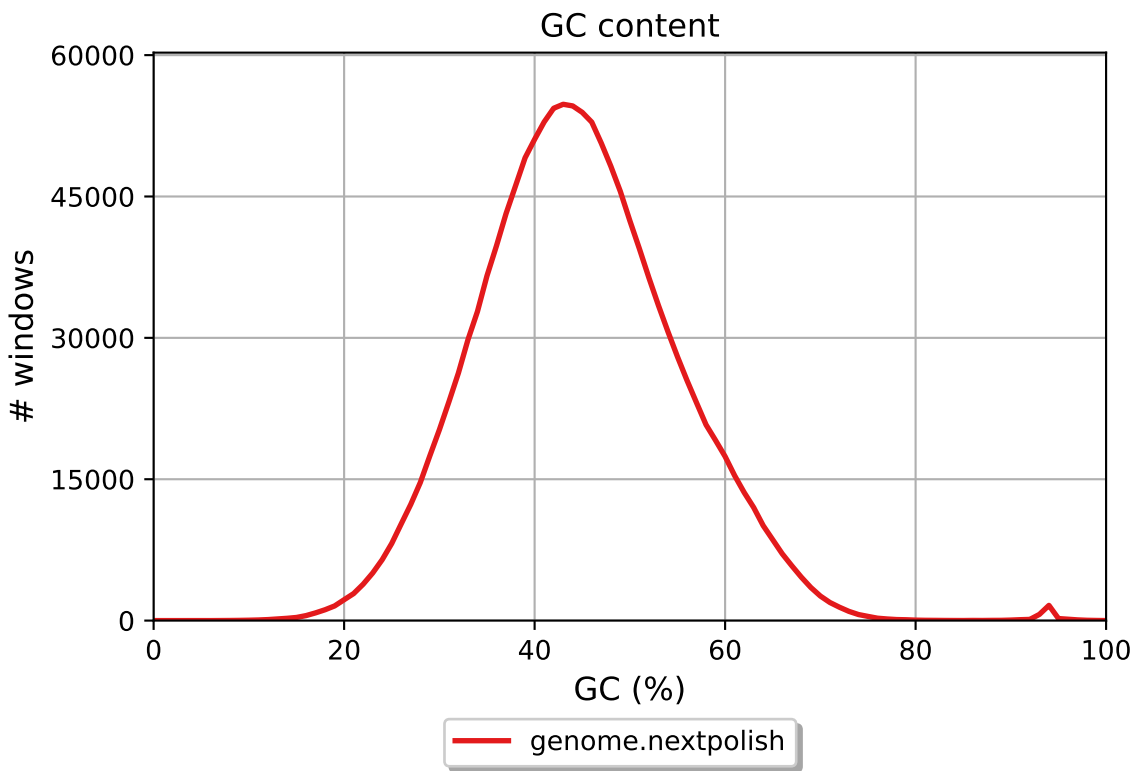
	genome.nextpolish
# contigs (>= 0 bp)	161716
# contigs (>= 1000 bp)	14098
# contigs (>= 5000 bp)	4176
# contigs (>= 10000 bp)	2124
# contigs (>= 25000 bp)	706
# contigs (>= 50000 bp)	337
Total length (>= 0 bp)	172685046
Total length (>= 1000 bp)	126116048
Total length (>= 5000 bp)	104826200
Total length (>= 10000 bp)	90433134
Total length (>= 25000 bp)	69284297
Total length (>= 50000 bp)	56708582
# contigs	29264
Largest contig	1247910
Total length	136349461
GC (%)	44.56
N50	26091
N90	1351
auN	139870.1
L50	663
L90	11157
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	125 + 22 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).

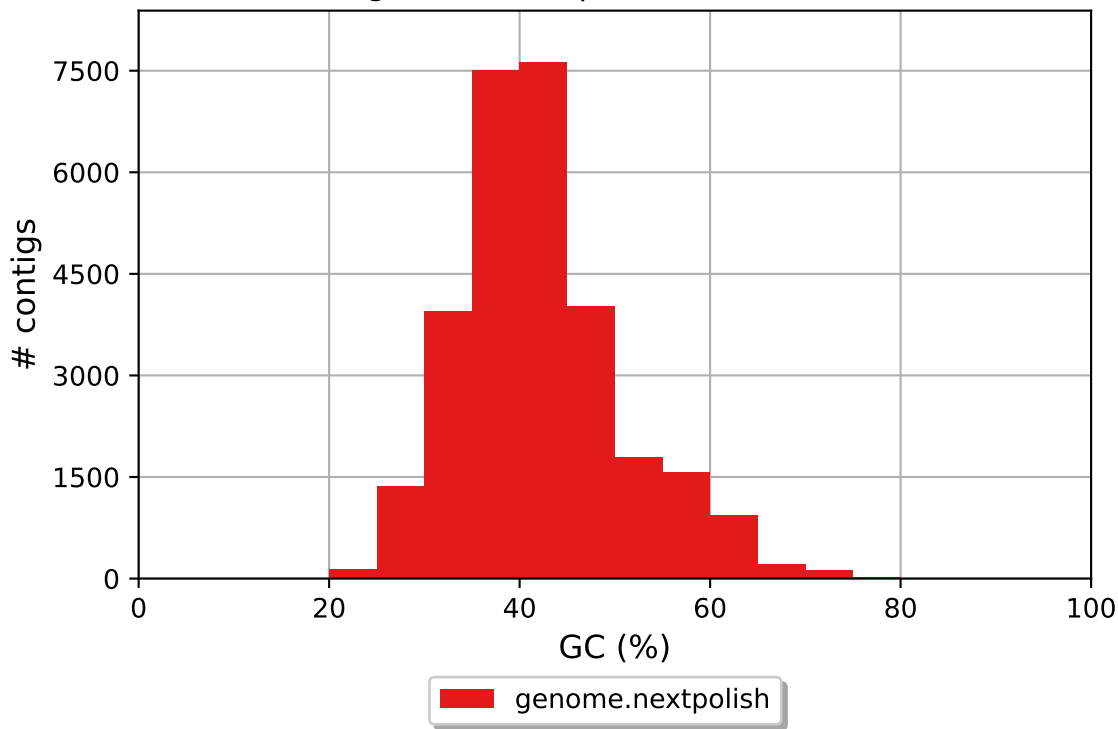


Cumulative length

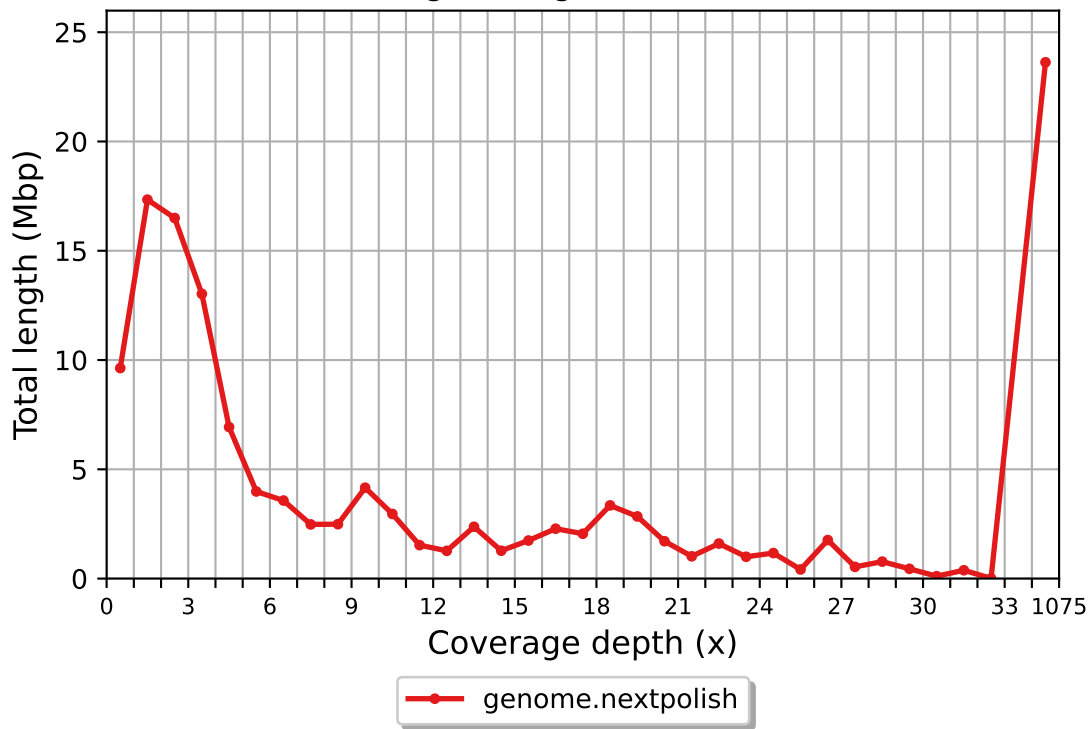




genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

