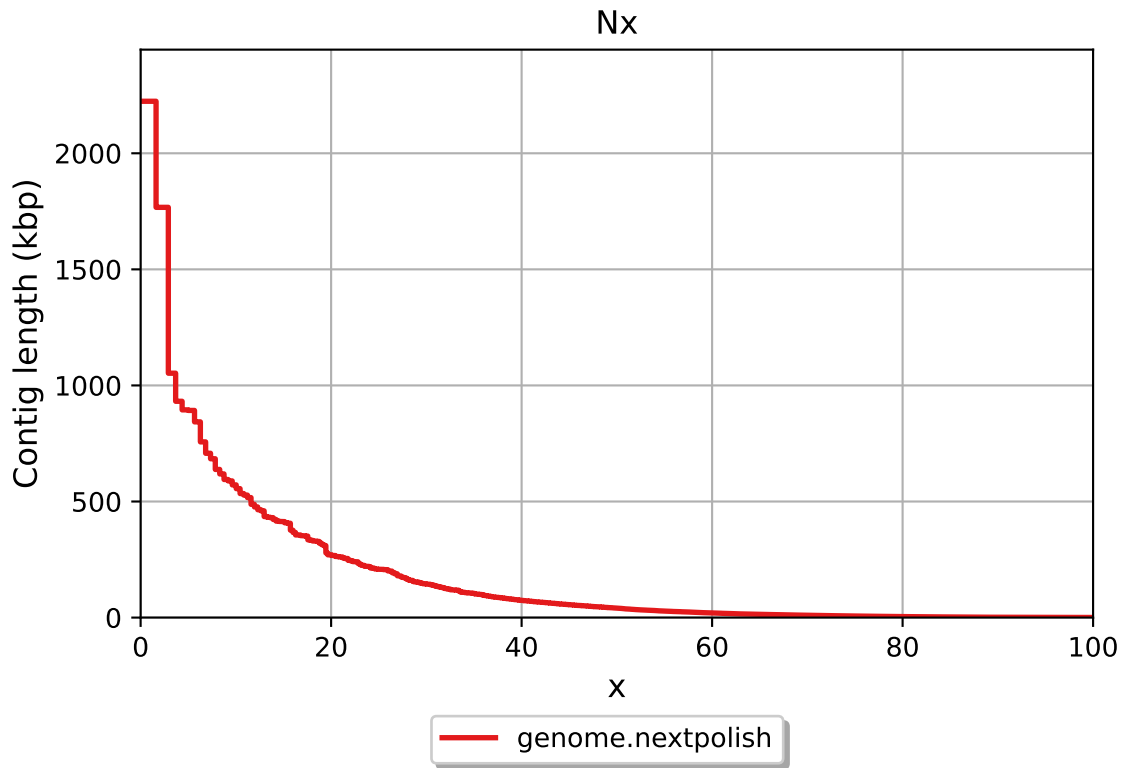
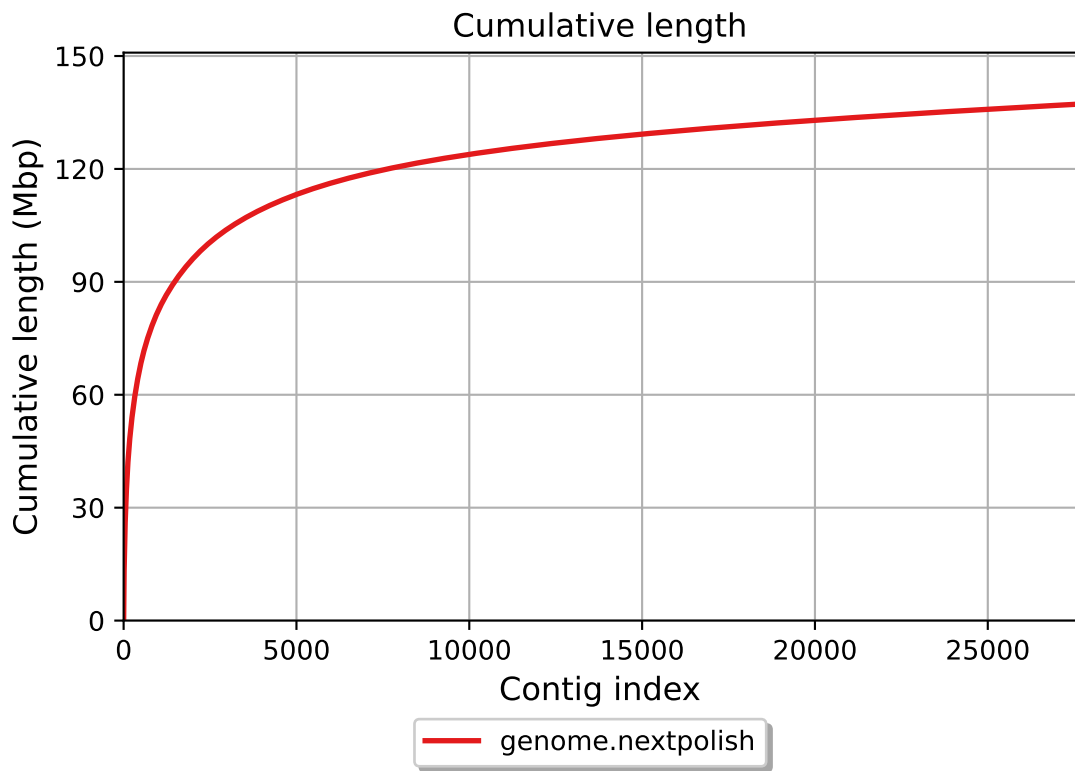


Report

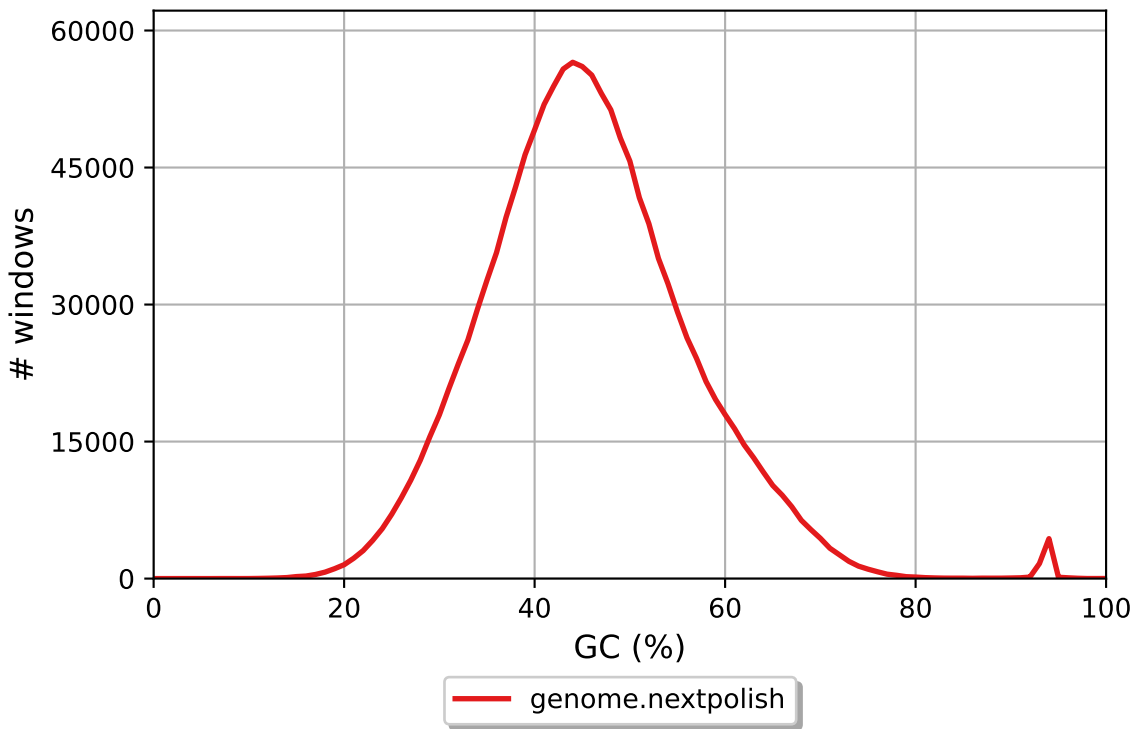
	genome.nextpolish
# contigs (>= 0 bp)	151827
# contigs (>= 1000 bp)	13071
# contigs (>= 5000 bp)	3638
# contigs (>= 10000 bp)	2011
# contigs (>= 25000 bp)	804
# contigs (>= 50000 bp)	409
Total length (>= 0 bp)	171665782
Total length (>= 1000 bp)	127466016
Total length (>= 5000 bp)	107598573
Total length (>= 10000 bp)	96216921
Total length (>= 25000 bp)	77945422
Total length (>= 50000 bp)	64286408
# contigs	27555
Largest contig	2224038
Total length	137179232
GC (%)	45.65
N50	40764
N90	1430
auN	197467.7
L50	505
L90	9711
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	119 + 27 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).

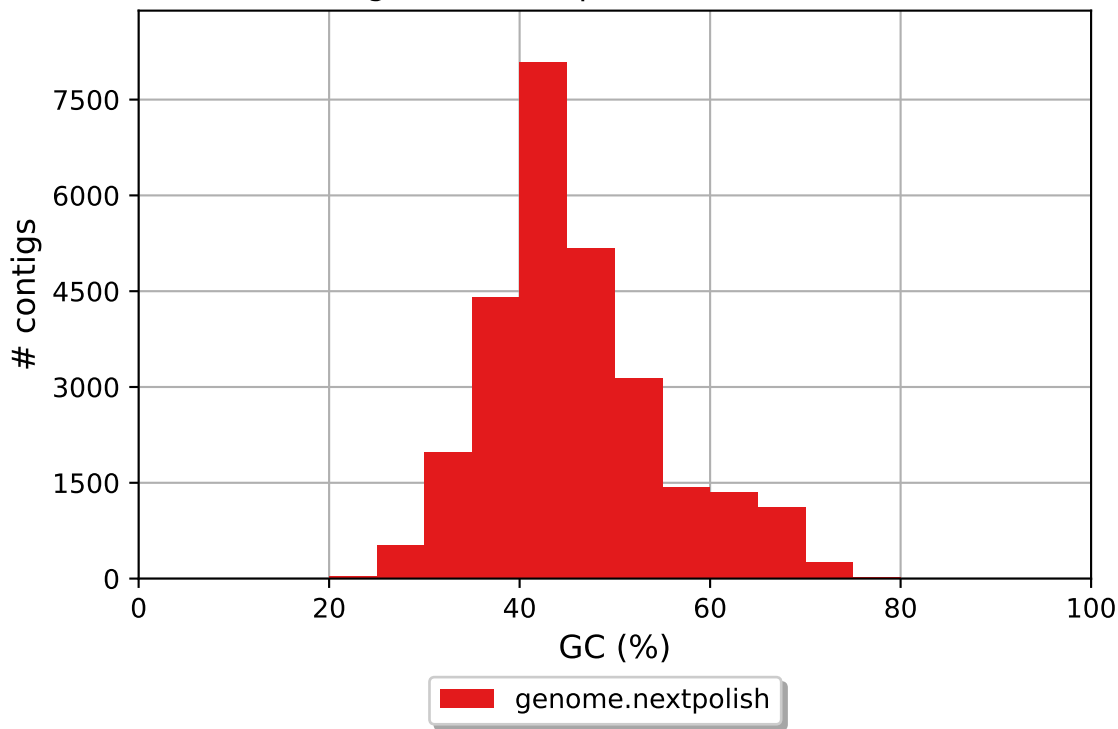




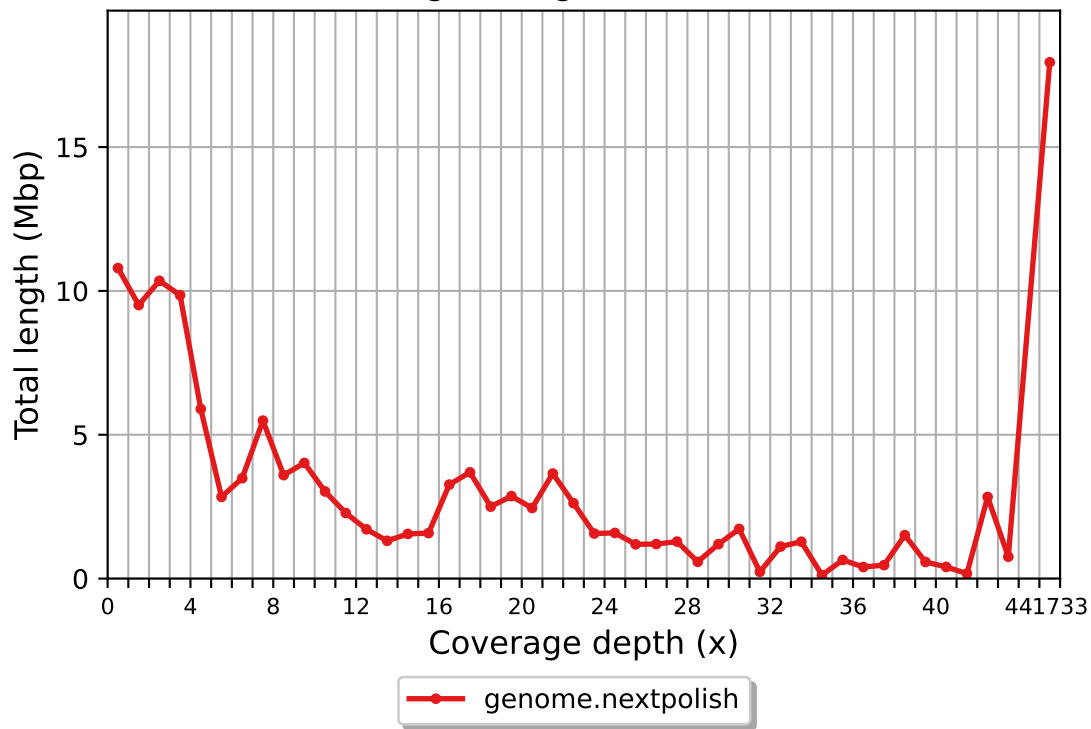
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

