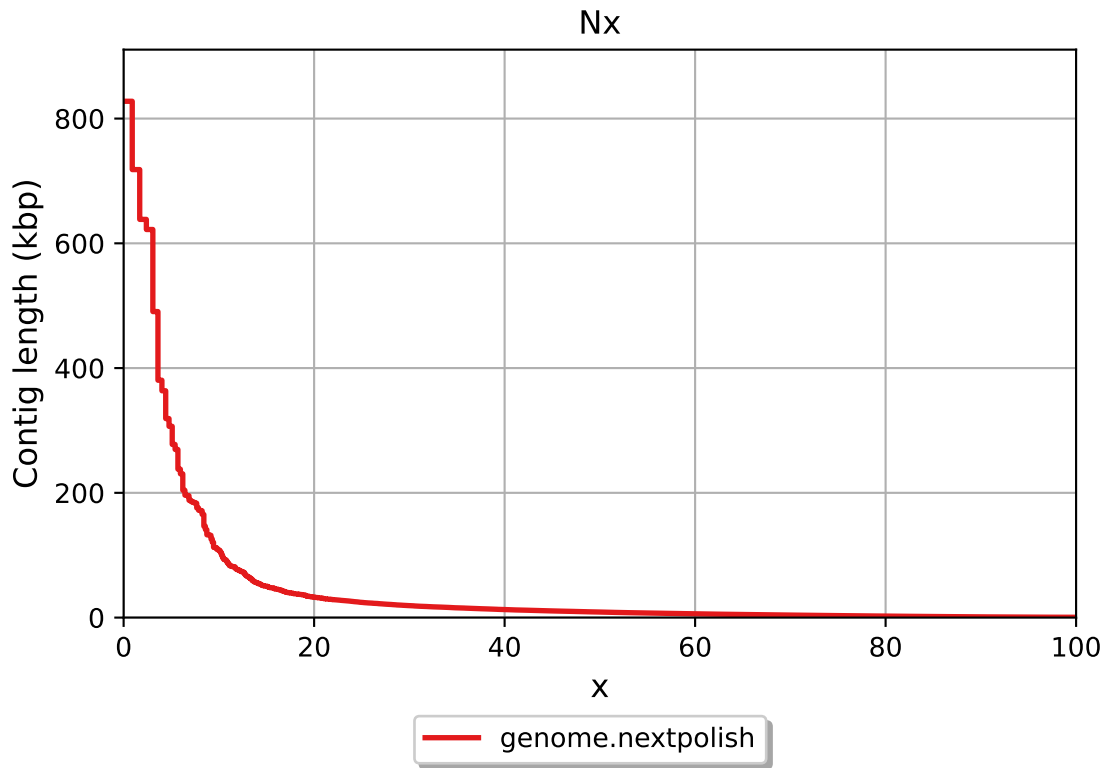
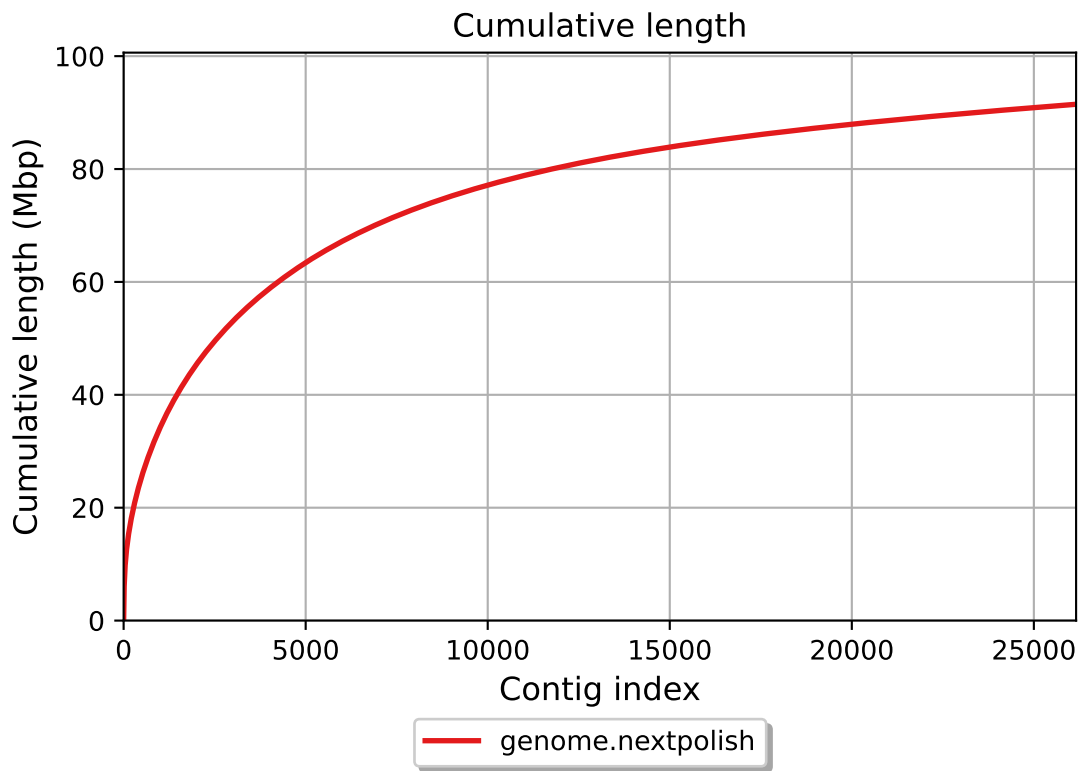


# Report

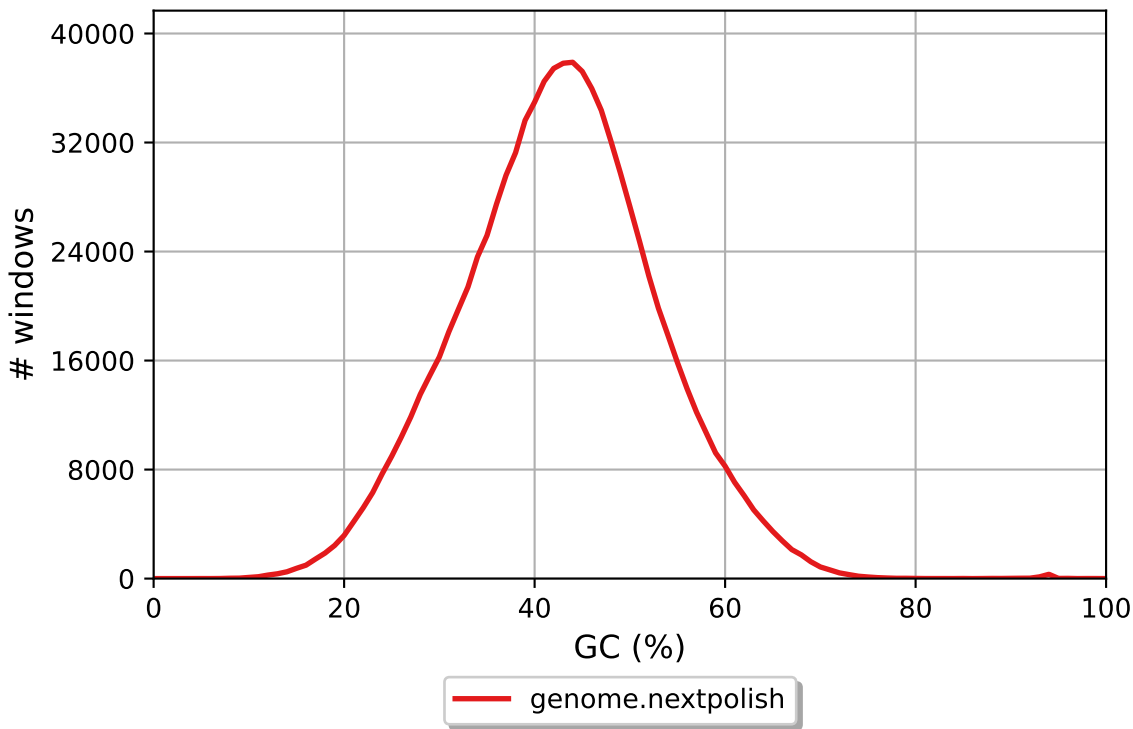
	genome.nextpolish
# contigs (>= 0 bp)	153996
# contigs (>= 1000 bp)	14998
# contigs (>= 5000 bp)	4105
# contigs (>= 10000 bp)	1737
# contigs (>= 25000 bp)	365
# contigs (>= 50000 bp)	103
Total length (>= 0 bp)	121286256
Total length (>= 1000 bp)	83891348
Total length (>= 5000 bp)	59334523
Total length (>= 10000 bp)	42899064
Total length (>= 25000 bp)	22522627
Total length (>= 50000 bp)	13790395
# contigs	26160
Largest contig	827733
Total length	91474401
GC (%)	42.58
N50	8878
N90	1162
auN	51127.6
L50	2039
L90	13547
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	19 + 17 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).

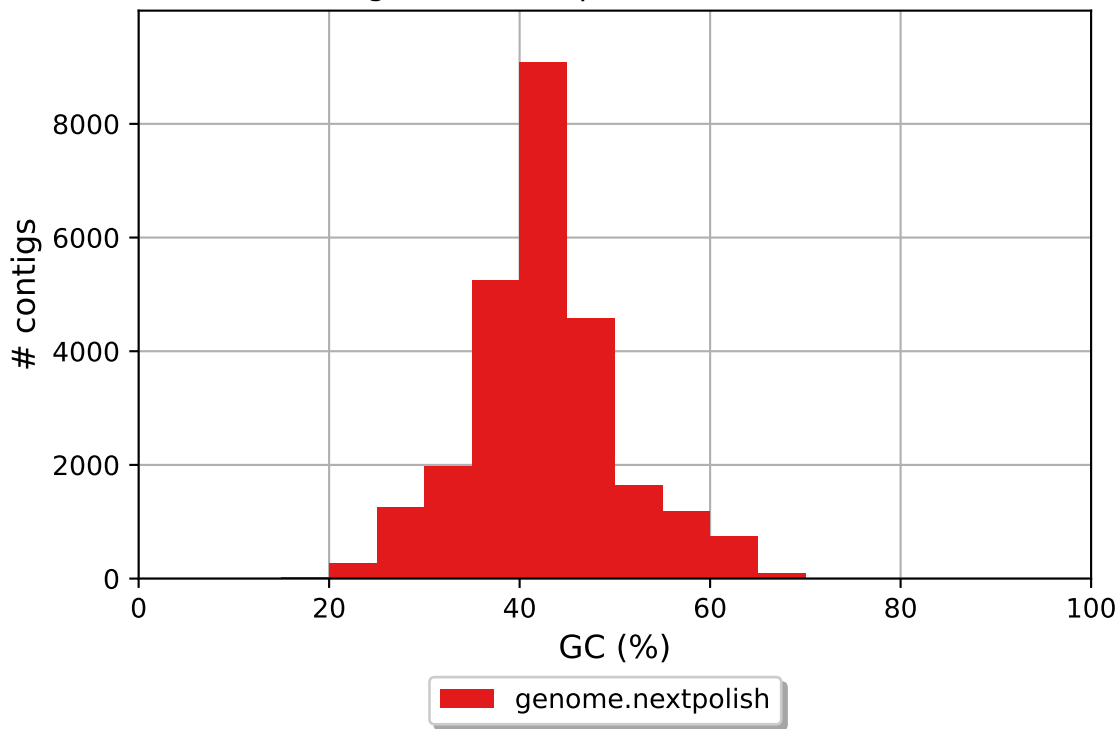




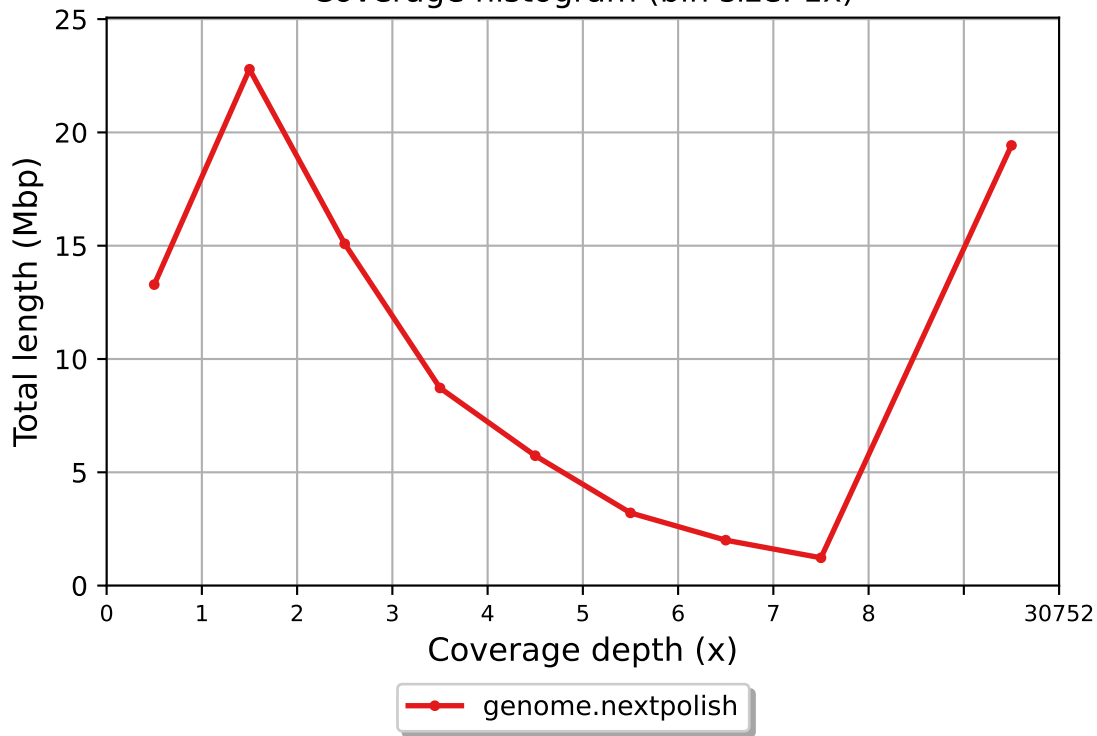
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

