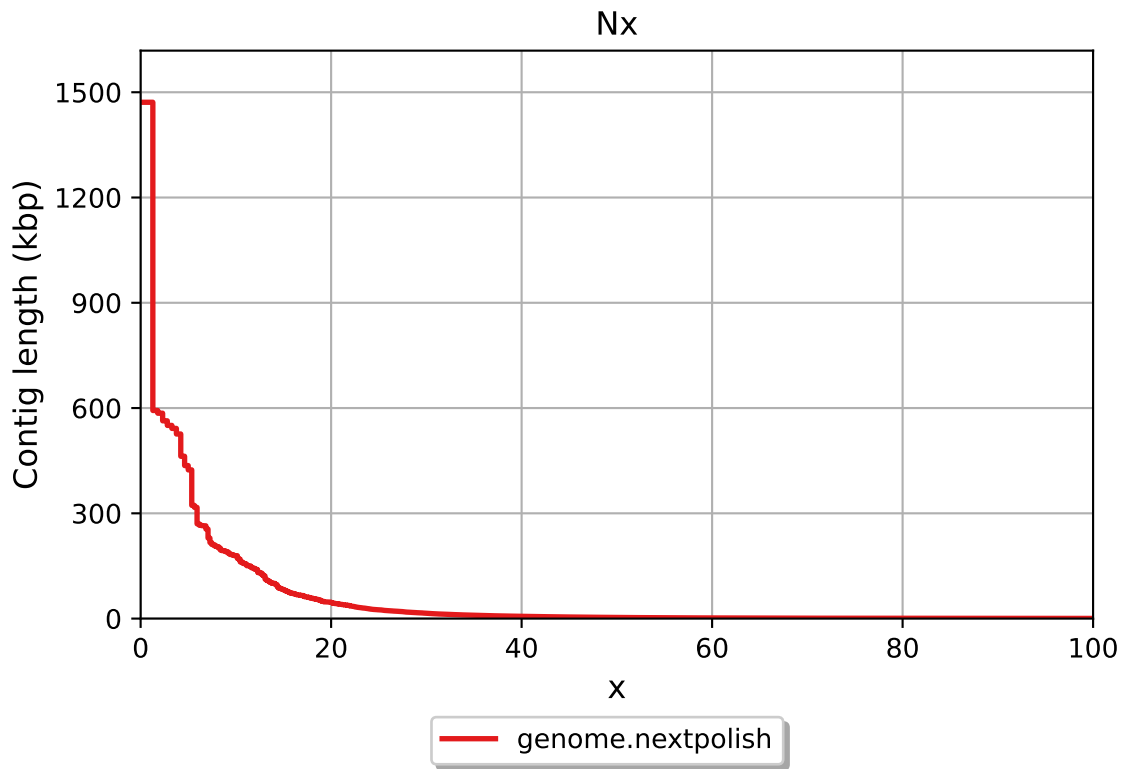
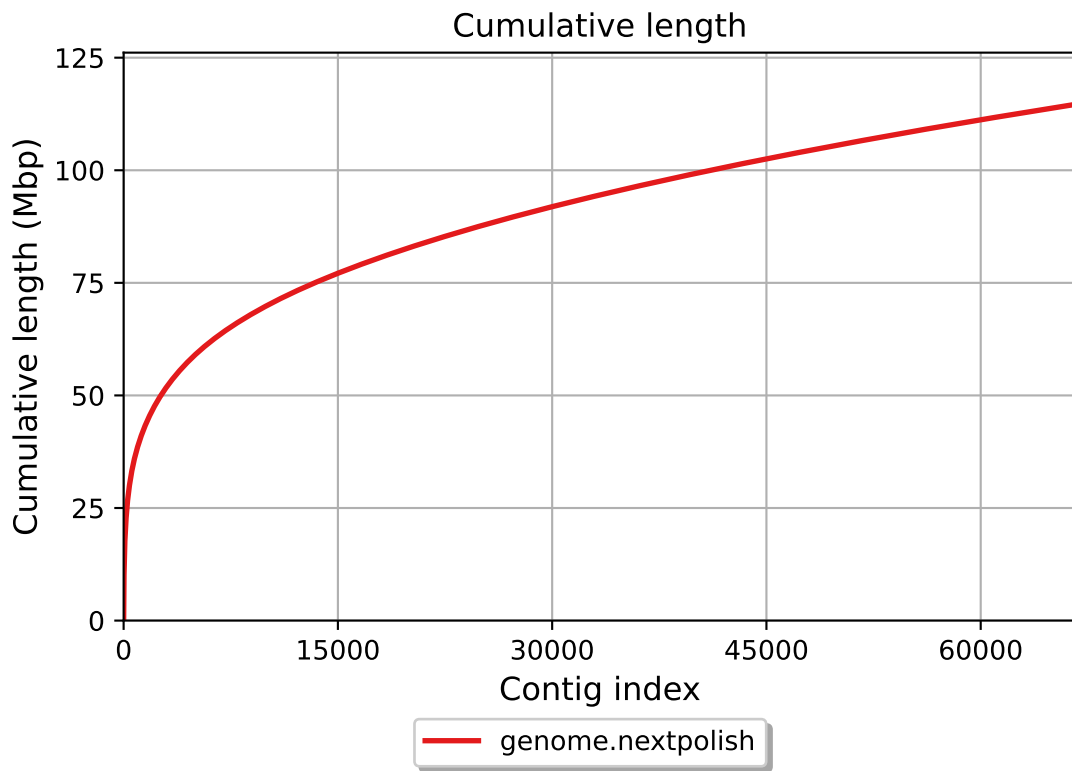


# Report

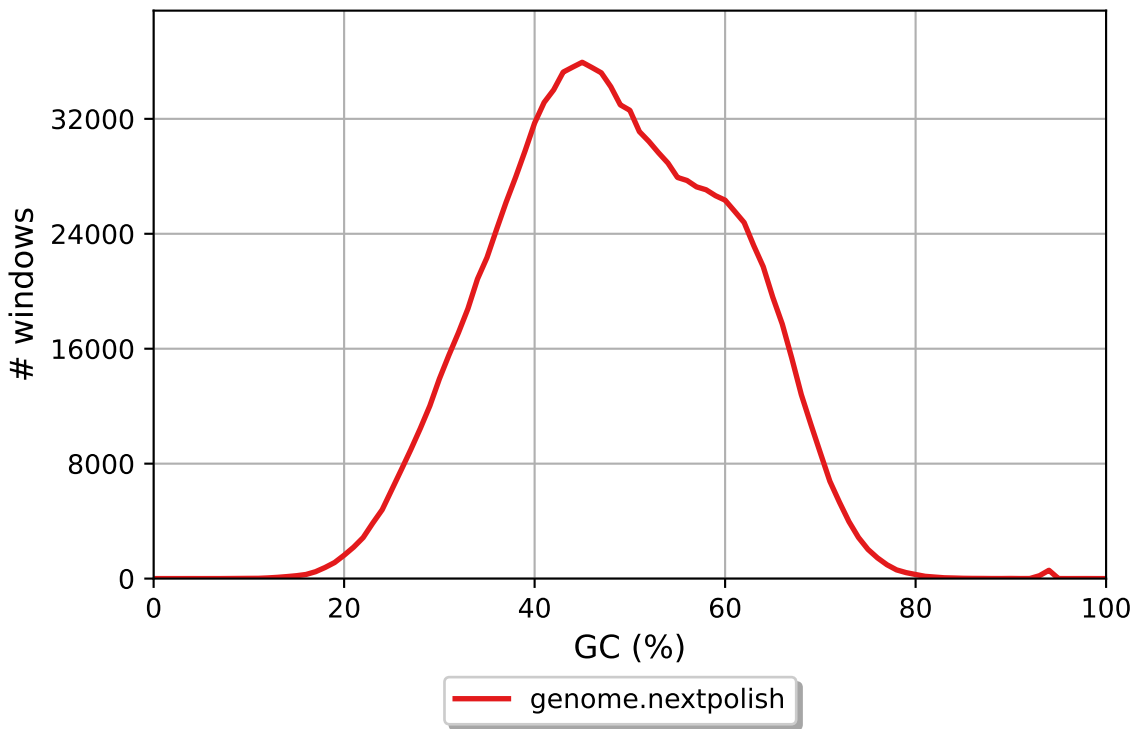
	genome.nextpolish
# contigs (>= 0 bp)	317384
# contigs (>= 1000 bp)	21240
# contigs (>= 5000 bp)	2577
# contigs (>= 10000 bp)	1065
# contigs (>= 25000 bp)	349
# contigs (>= 50000 bp)	153
Total length (>= 0 bp)	187589439
Total length (>= 1000 bp)	84084492
Total length (>= 5000 bp)	49849473
Total length (>= 10000 bp)	39458714
Total length (>= 25000 bp)	28649183
Total length (>= 50000 bp)	21808162
# contigs	66678
Largest contig	1471437
Total length	114653844
GC (%)	48.38
N50	3243
N90	623
auN	65299.1
L50	4455
L90	46044
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	64 + 23 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).

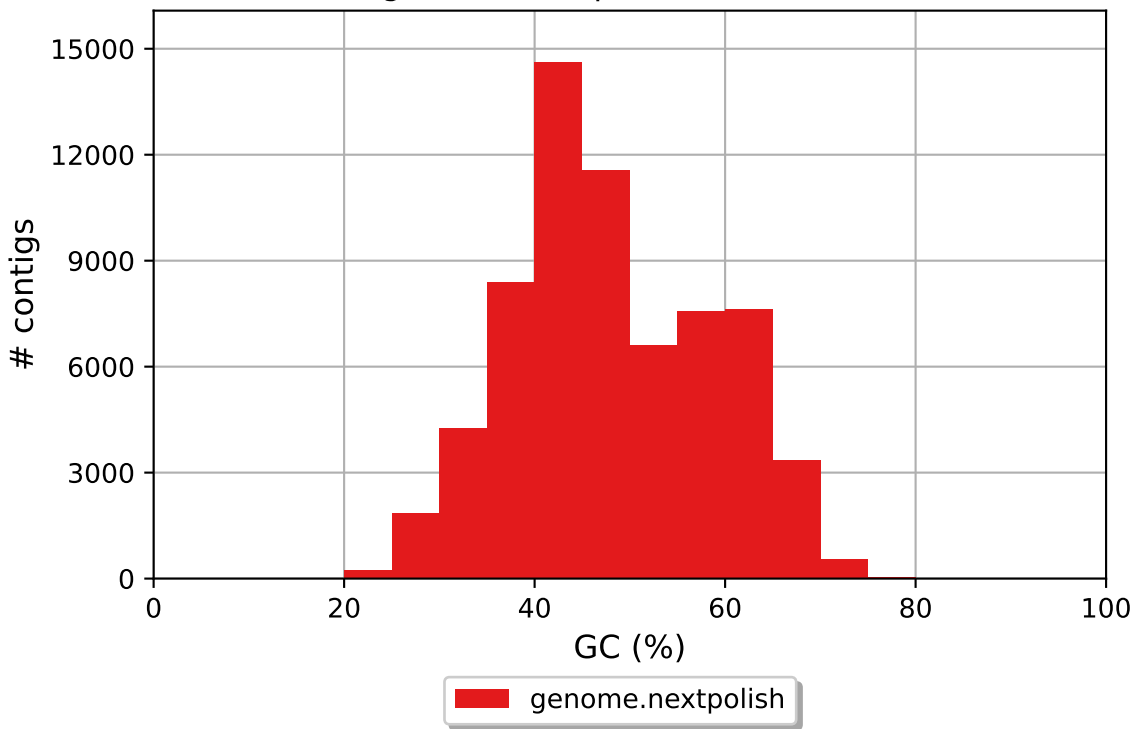




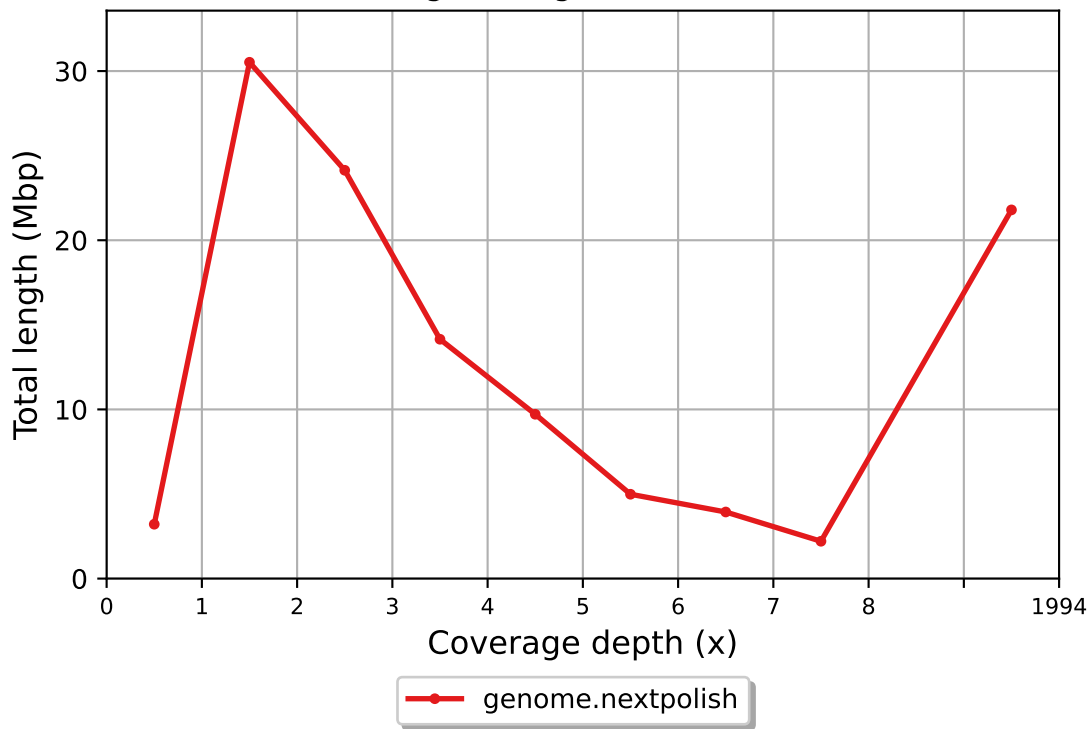
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

