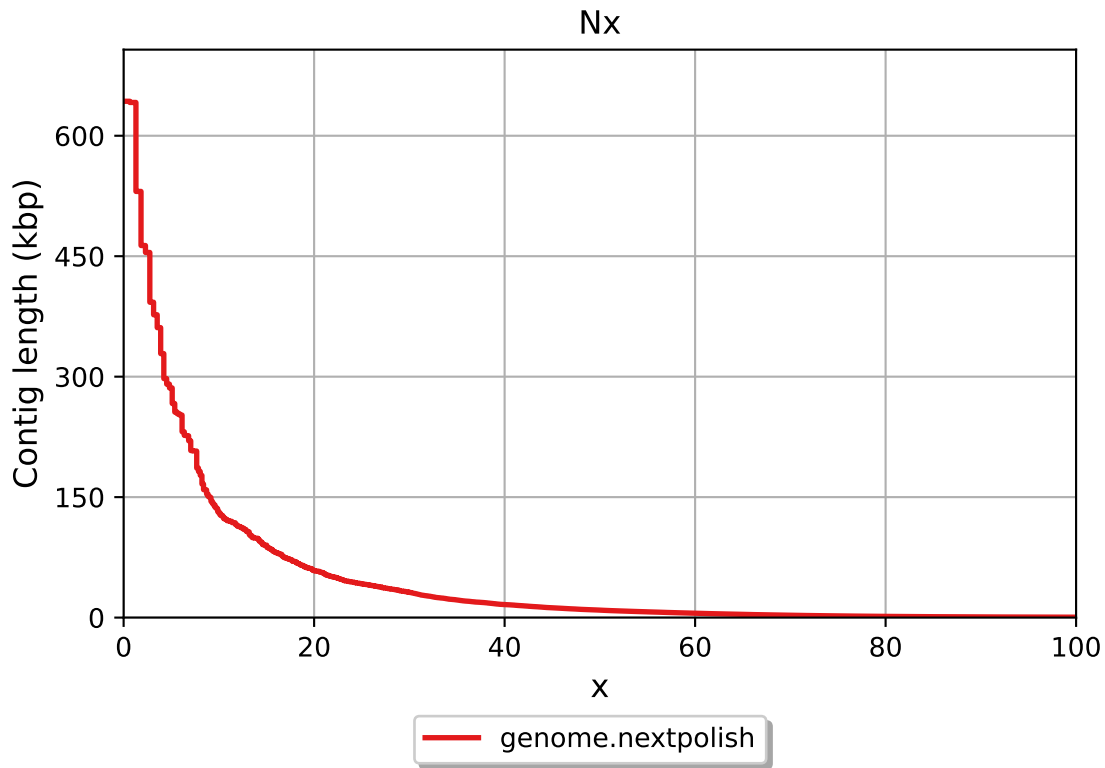
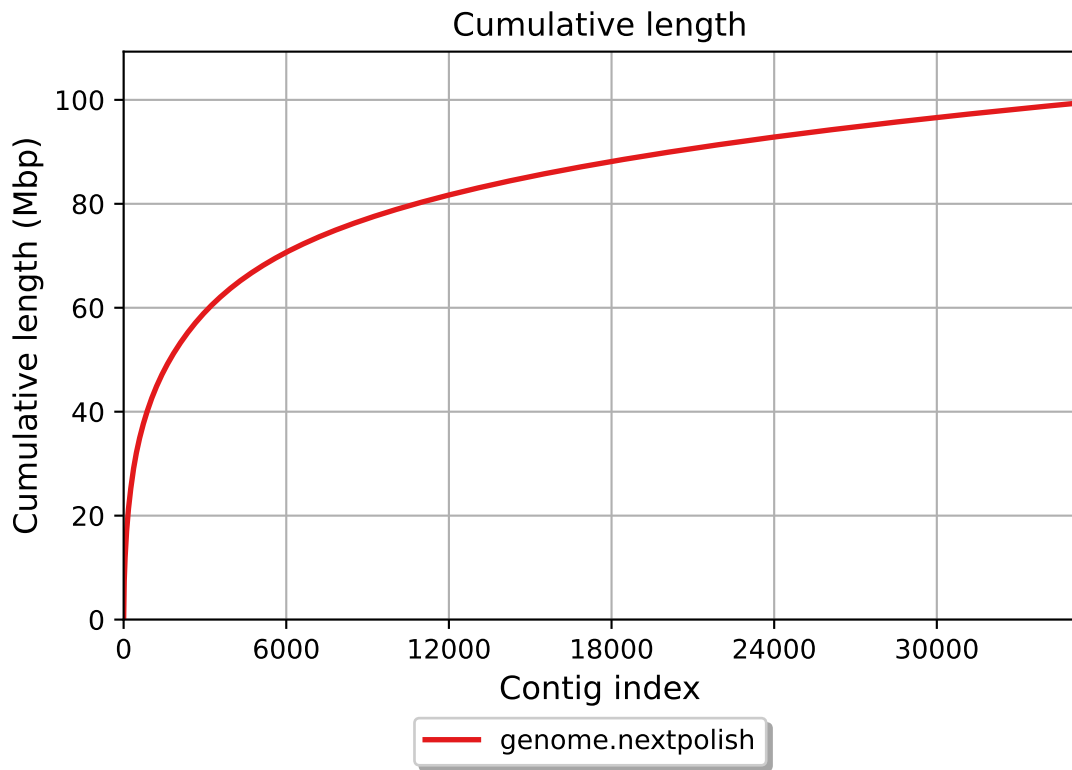


Report

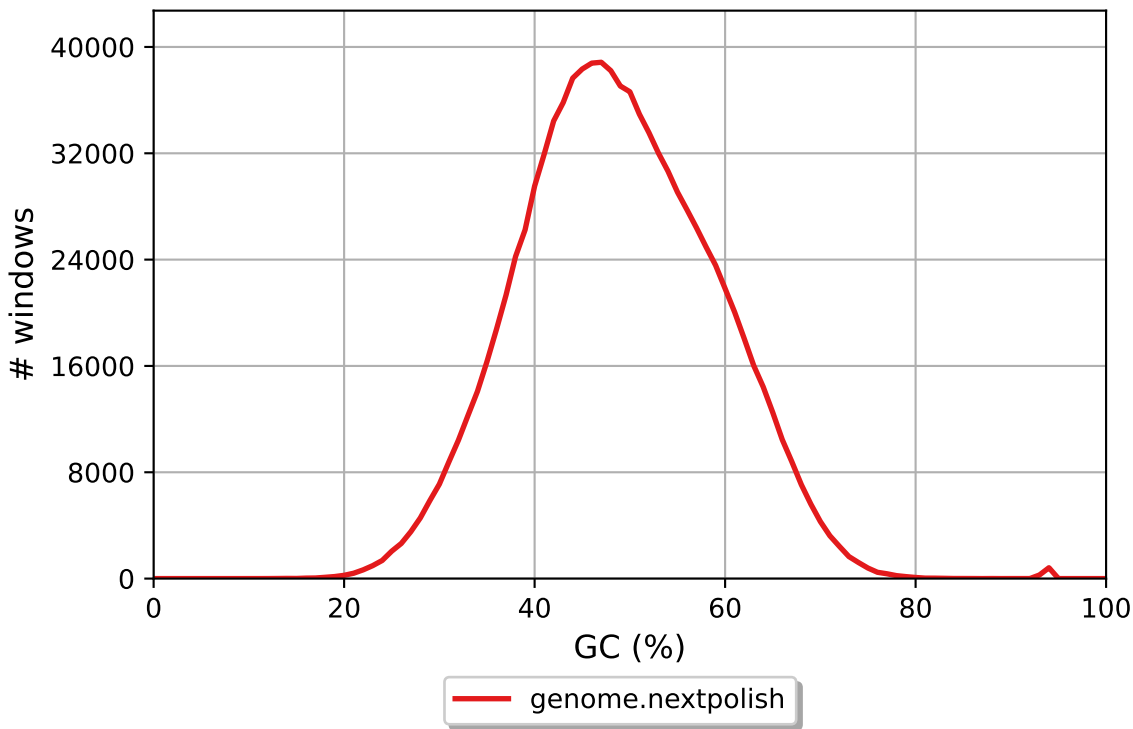
	genome.nextpolish
# contigs (>= 0 bp)	126310
# contigs (>= 1000 bp)	15949
# contigs (>= 5000 bp)	3341
# contigs (>= 10000 bp)	1512
# contigs (>= 25000 bp)	492
# contigs (>= 50000 bp)	194
Total length (>= 0 bp)	125561883
Total length (>= 1000 bp)	86210648
Total length (>= 5000 bp)	60970322
Total length (>= 10000 bp)	48153659
Total length (>= 25000 bp)	32638541
Total length (>= 50000 bp)	22085651
# contigs	35138
Largest contig	642930
Total length	99345239
GC (%)	48.67
N50	9308
N90	835
auN	51433.0
L50	1670
L90	19460
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	37 + 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).

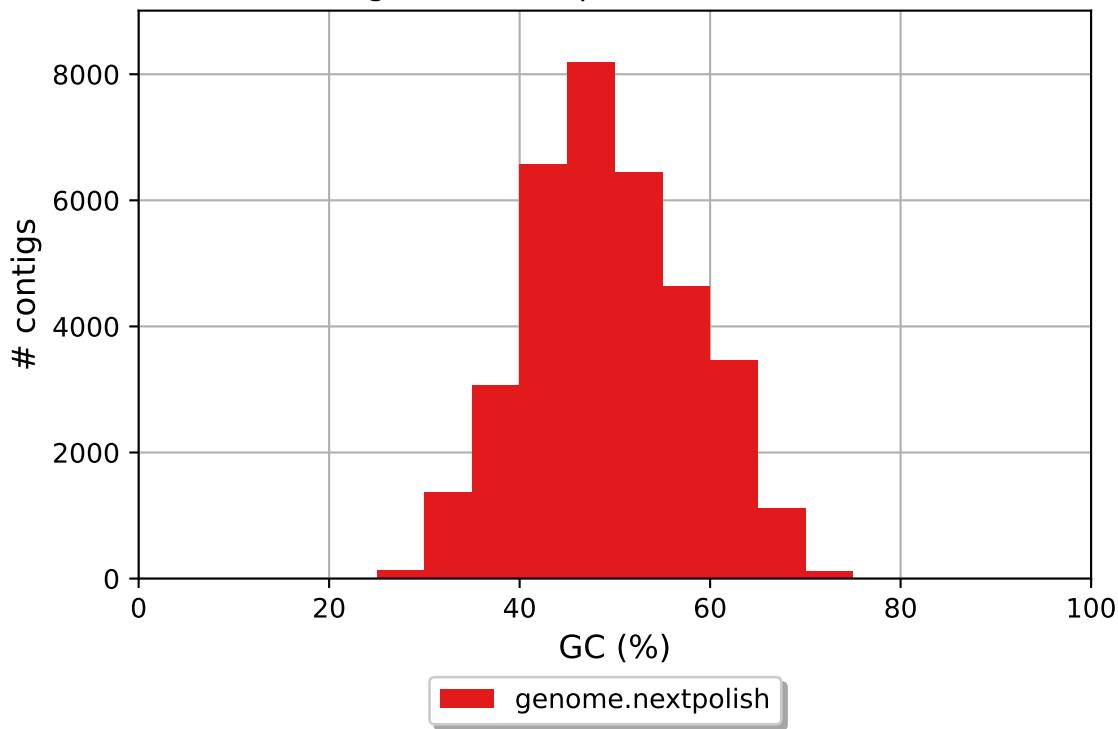




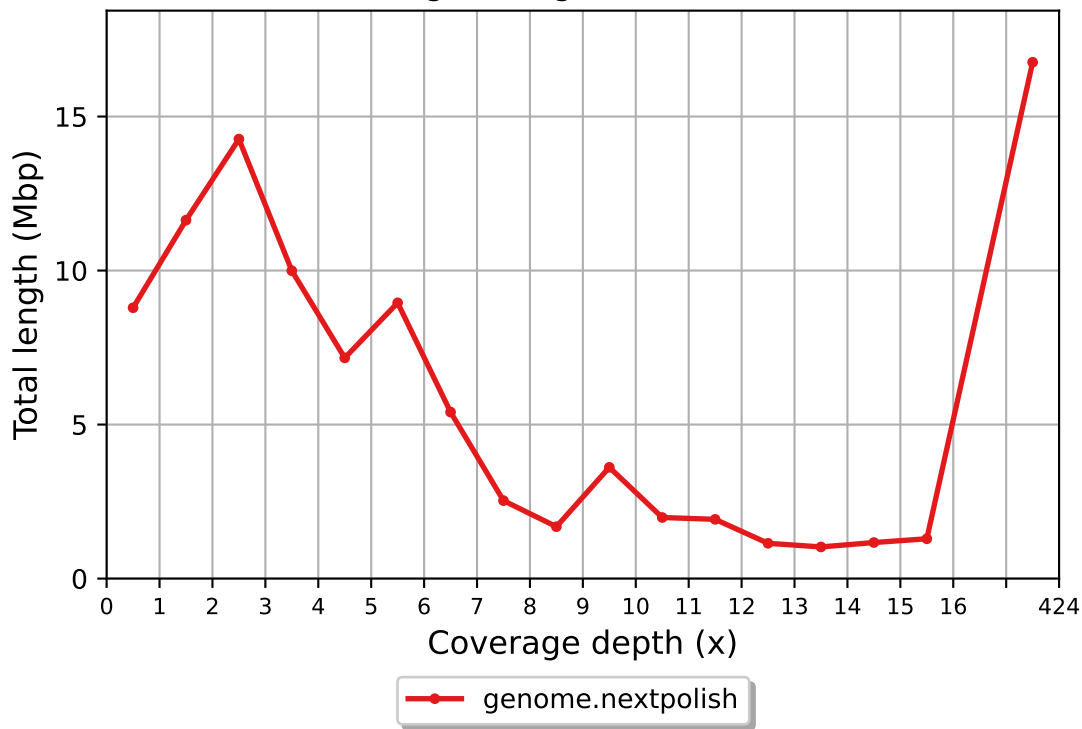
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

