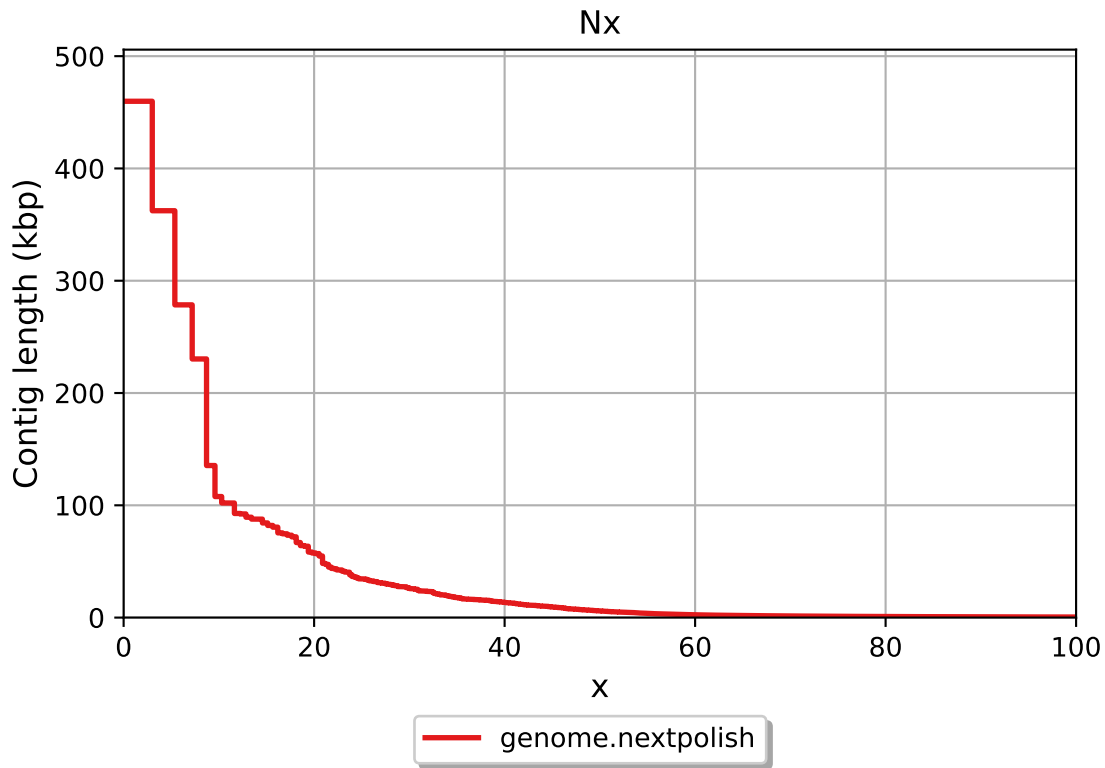


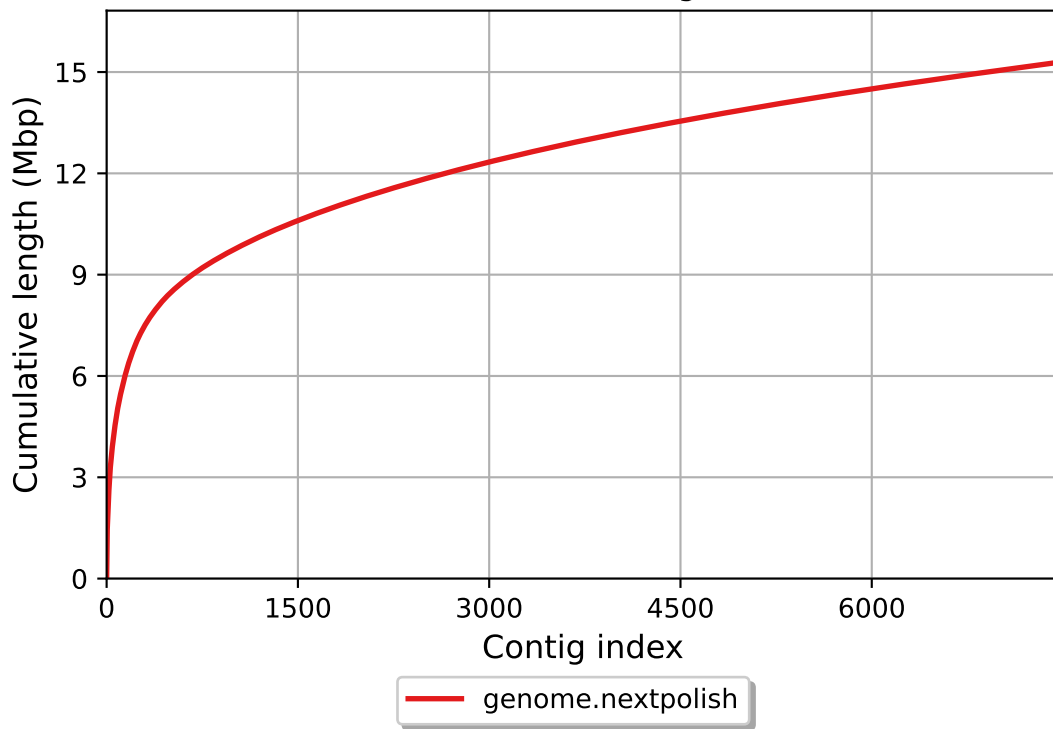
# Report

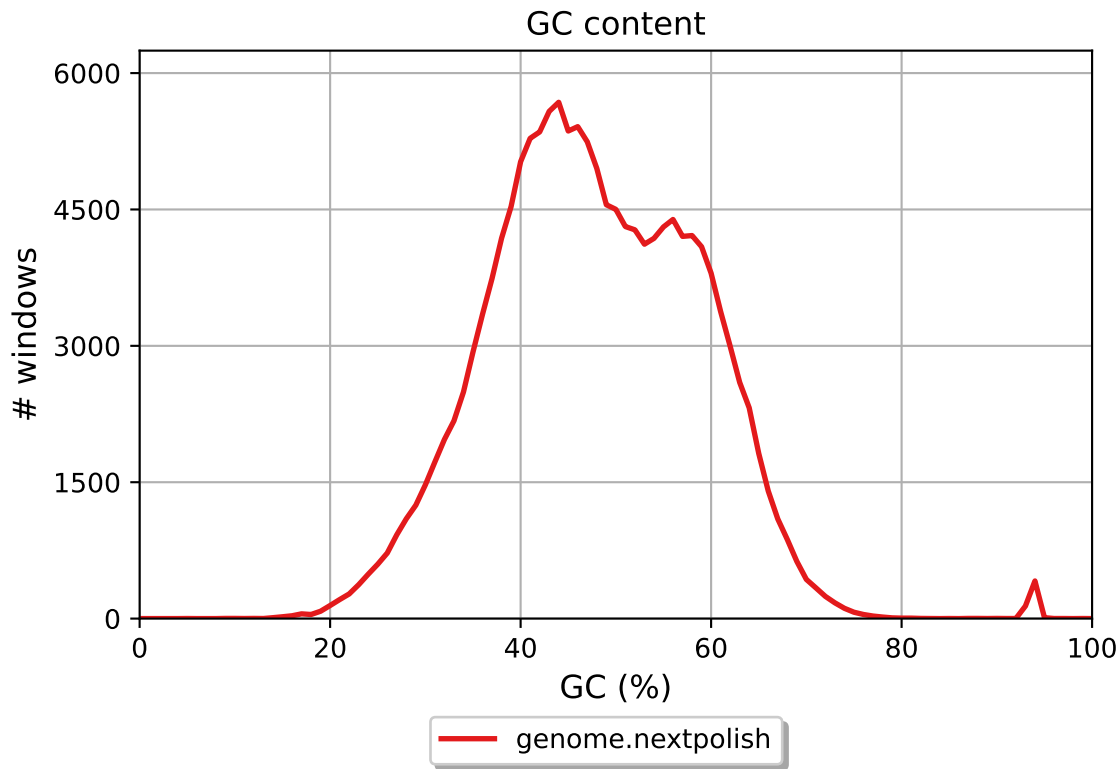
	genome.nextpolish
# contigs (>= 0 bp)	55912
# contigs (>= 1000 bp)	2707
# contigs (>= 5000 bp)	376
# contigs (>= 10000 bp)	212
# contigs (>= 25000 bp)	73
# contigs (>= 50000 bp)	27
Total length (>= 0 bp)	28356410
Total length (>= 1000 bp)	12053954
Total length (>= 5000 bp)	7929634
Total length (>= 10000 bp)	6801408
Total length (>= 25000 bp)	4720368
Total length (>= 50000 bp)	3192578
# contigs	7469
Largest contig	459827
Total length	15289434
GC (%)	47.79
N50	5939
N90	674
auN	48239.9
L50	324
L90	4810
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	25 + 4 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).

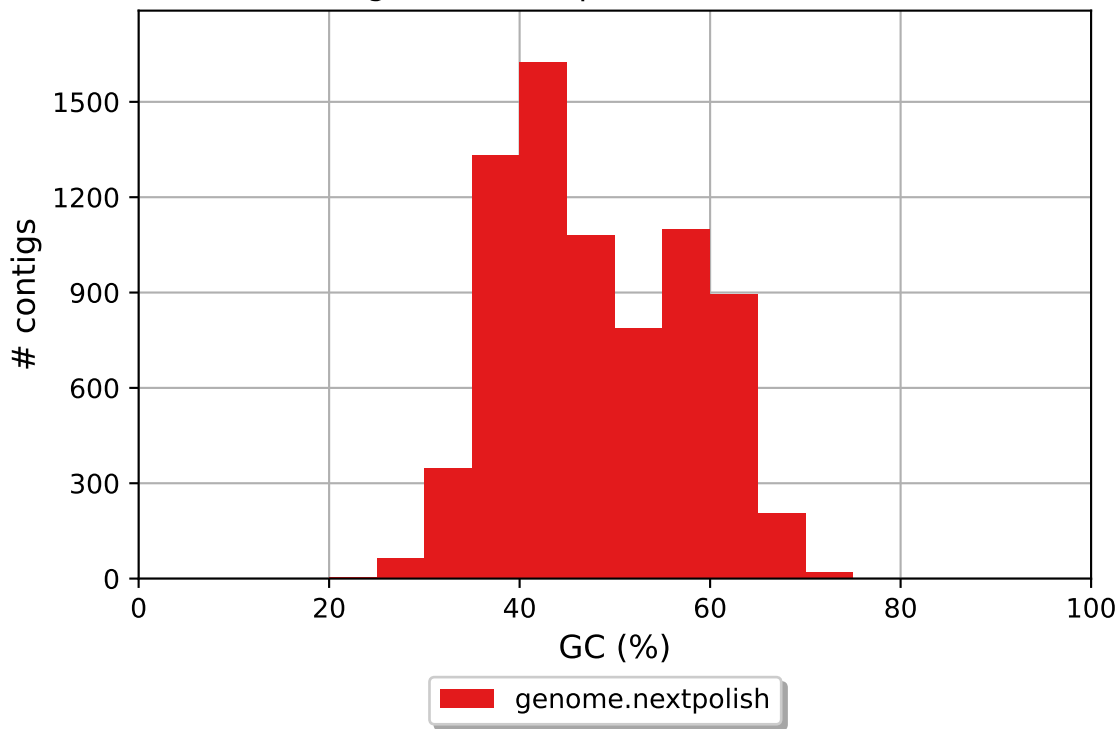


Cumulative length

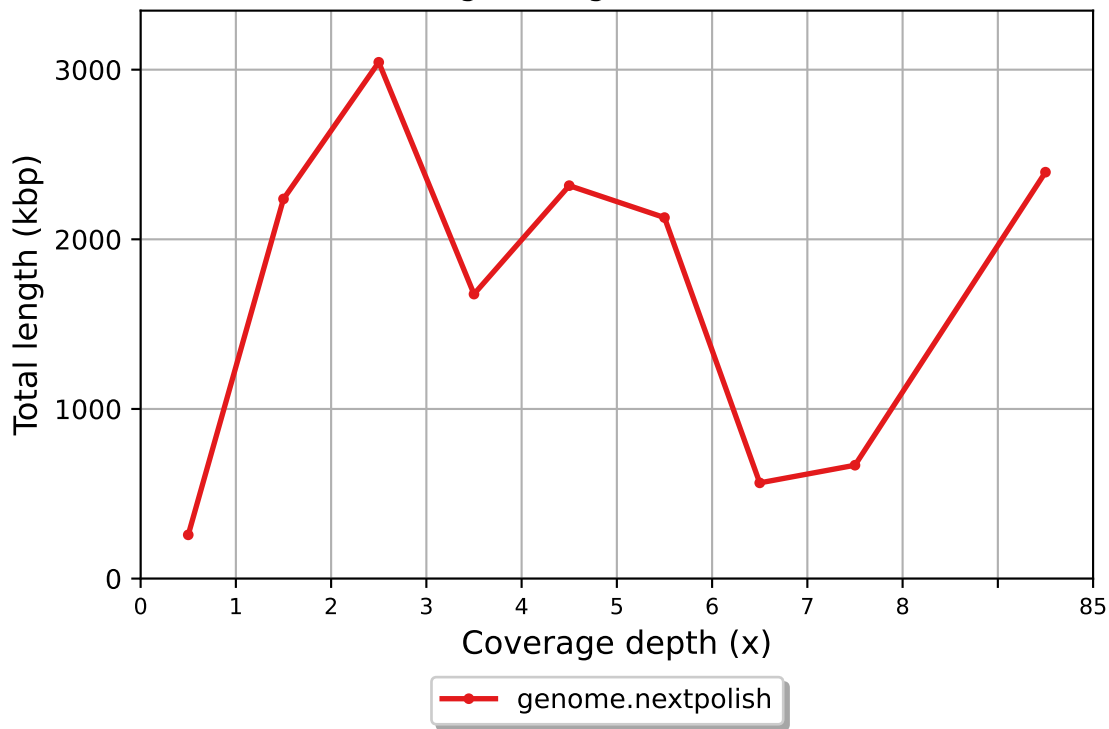




genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

