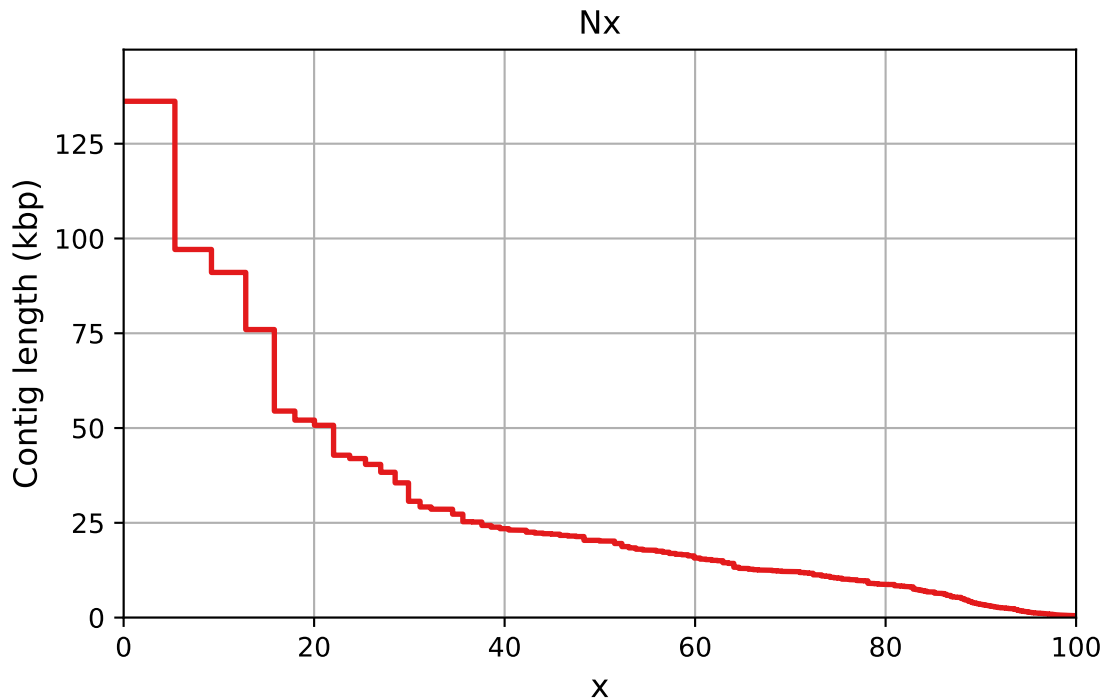


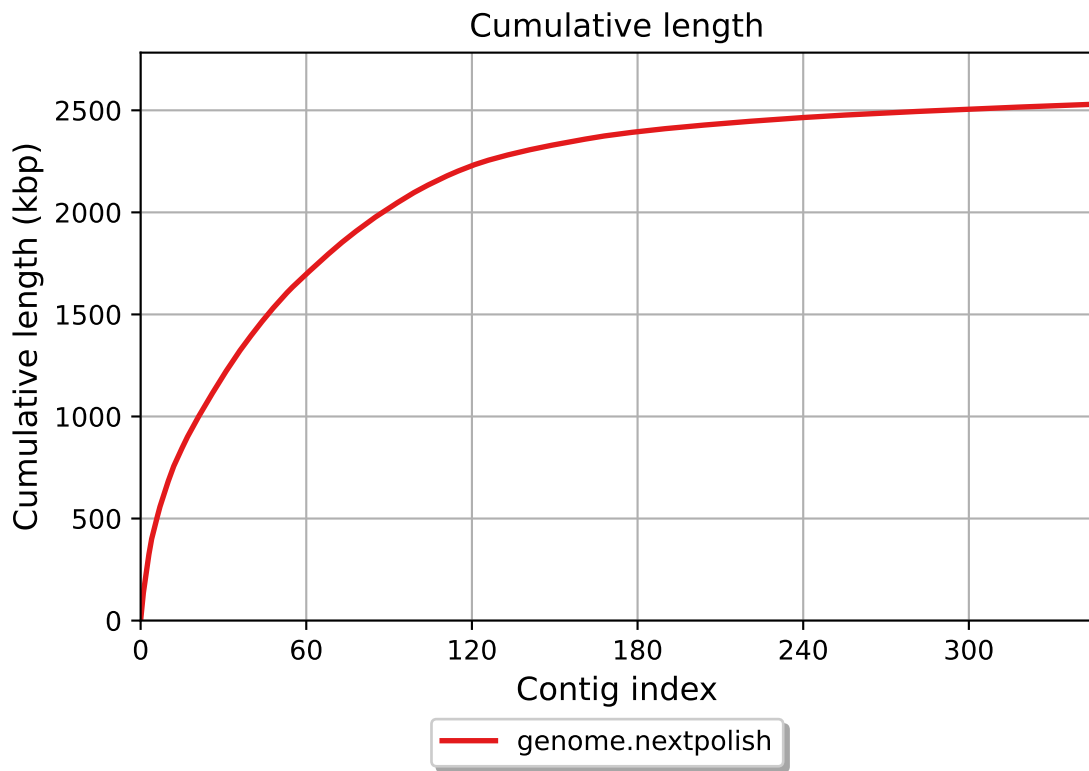
Report

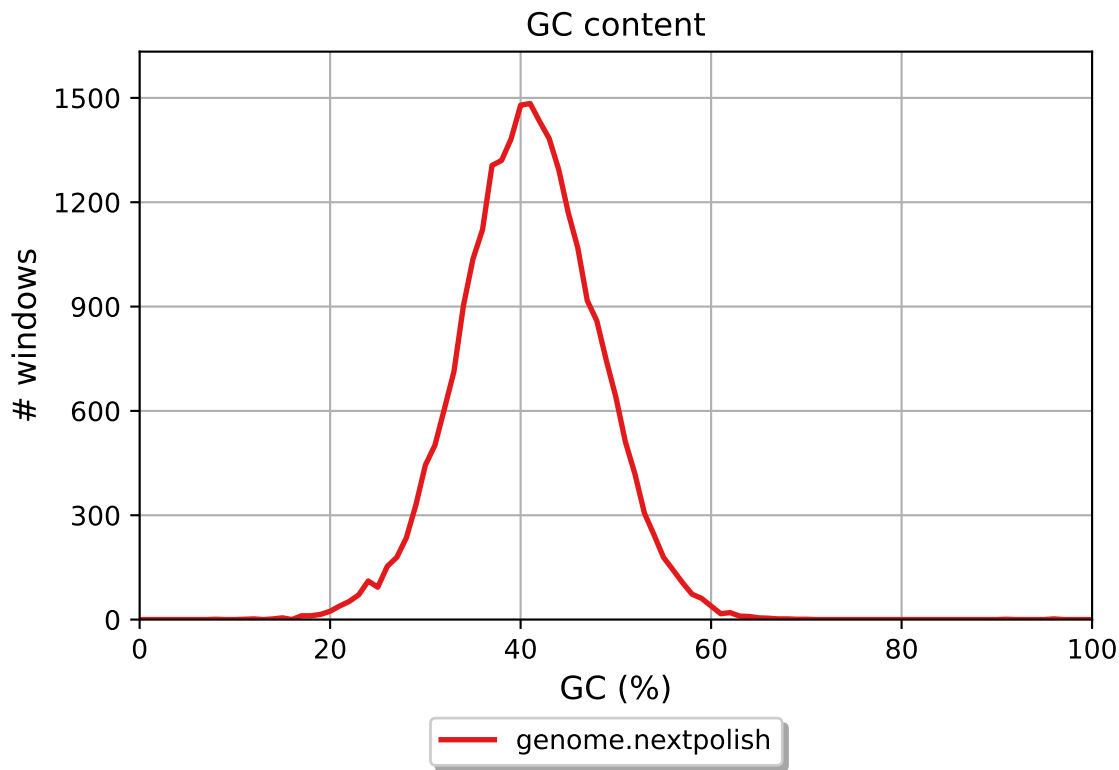
	genome.nextpolish
# contigs (>= 0 bp)	811
# contigs (>= 1000 bp)	223
# contigs (>= 5000 bp)	120
# contigs (>= 10000 bp)	81
# contigs (>= 25000 bp)	19
# contigs (>= 50000 bp)	7
Total length (>= 0 bp)	2667530
Total length (>= 1000 bp)	2448496
Total length (>= 5000 bp)	2228937
Total length (>= 10000 bp)	1937485
Total length (>= 25000 bp)	951456
Total length (>= 50000 bp)	557600
# contigs	345
Largest contig	136195
Total length	2529631
GC (%)	40.87
N50	20199
N90	3497
auN	32983.1
L50	34
L90	132
# N's per 100 kbp	1.54
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 2 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).

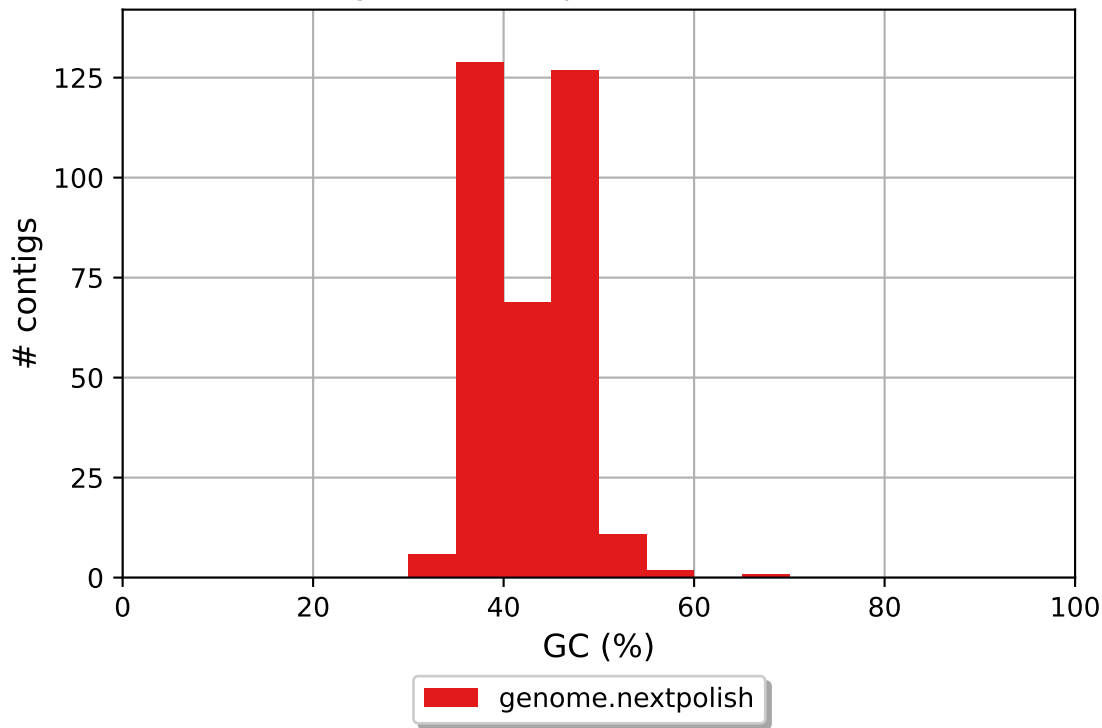


— genome.nextpolish

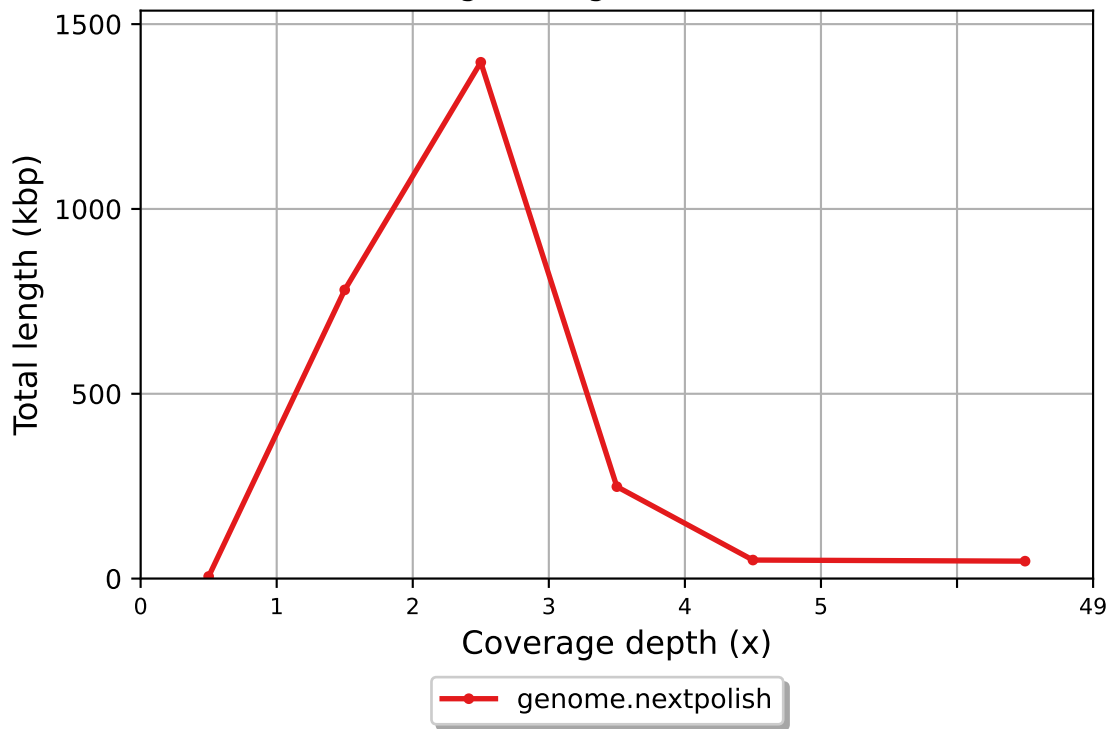




genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

