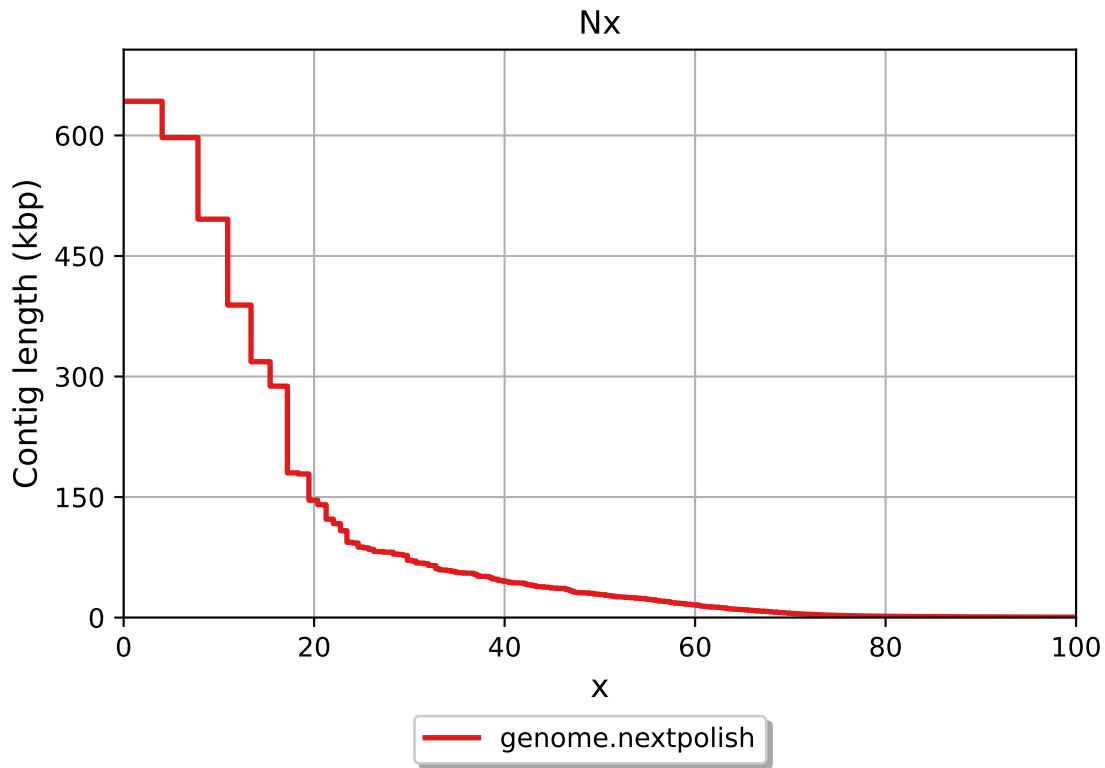


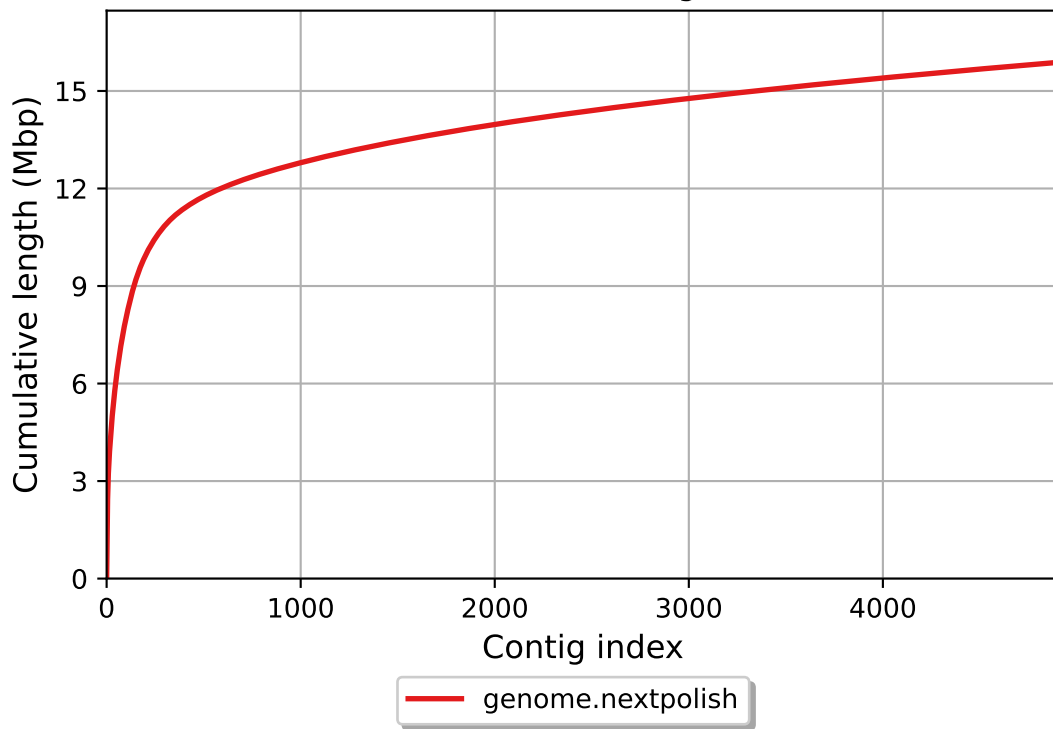
Report

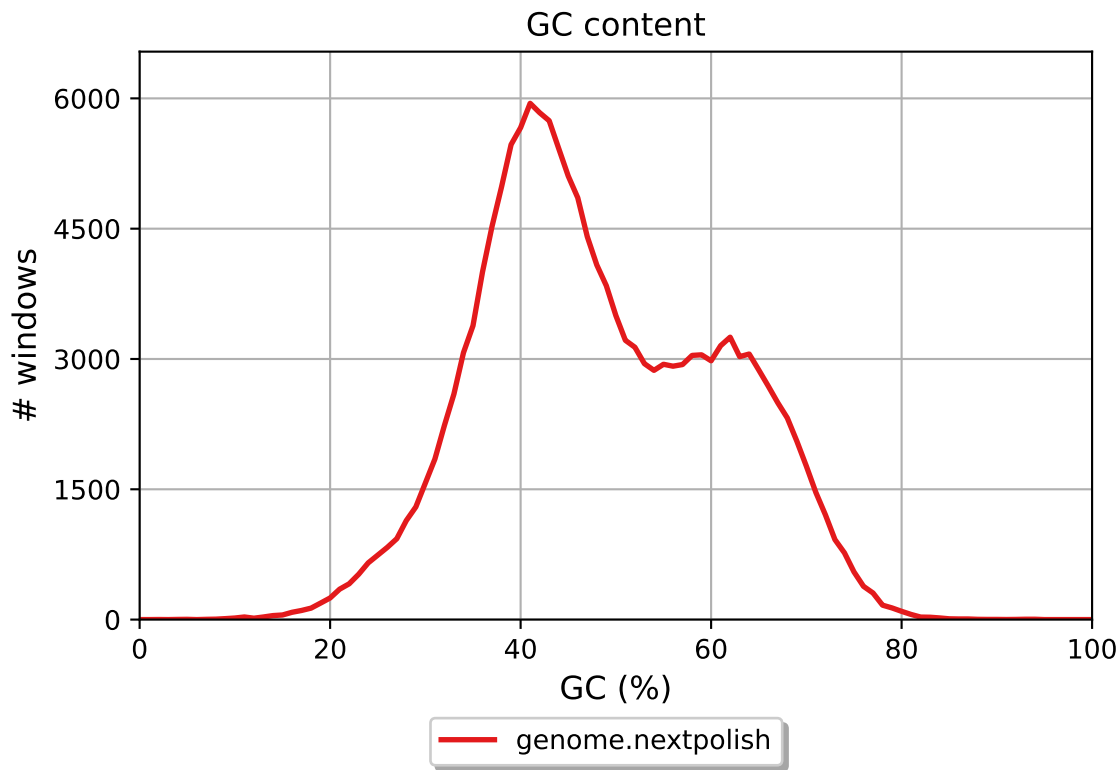
	genome.nextpolish
# contigs (>= 0 bp)	48488
# contigs (>= 1000 bp)	1825
# contigs (>= 5000 bp)	356
# contigs (>= 10000 bp)	232
# contigs (>= 25000 bp)	114
# contigs (>= 50000 bp)	48
Total length (>= 0 bp)	23818298
Total length (>= 1000 bp)	13802049
Total length (>= 5000 bp)	11190335
Total length (>= 10000 bp)	10293239
Total length (>= 25000 bp)	8388442
Total length (>= 50000 bp)	6095765
# contigs	4908
Largest contig	642539
Total length	15882941
GC (%)	48.29
N50	28634
N90	823
auN	112927.2
L50	98
L90	2371
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 6 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).

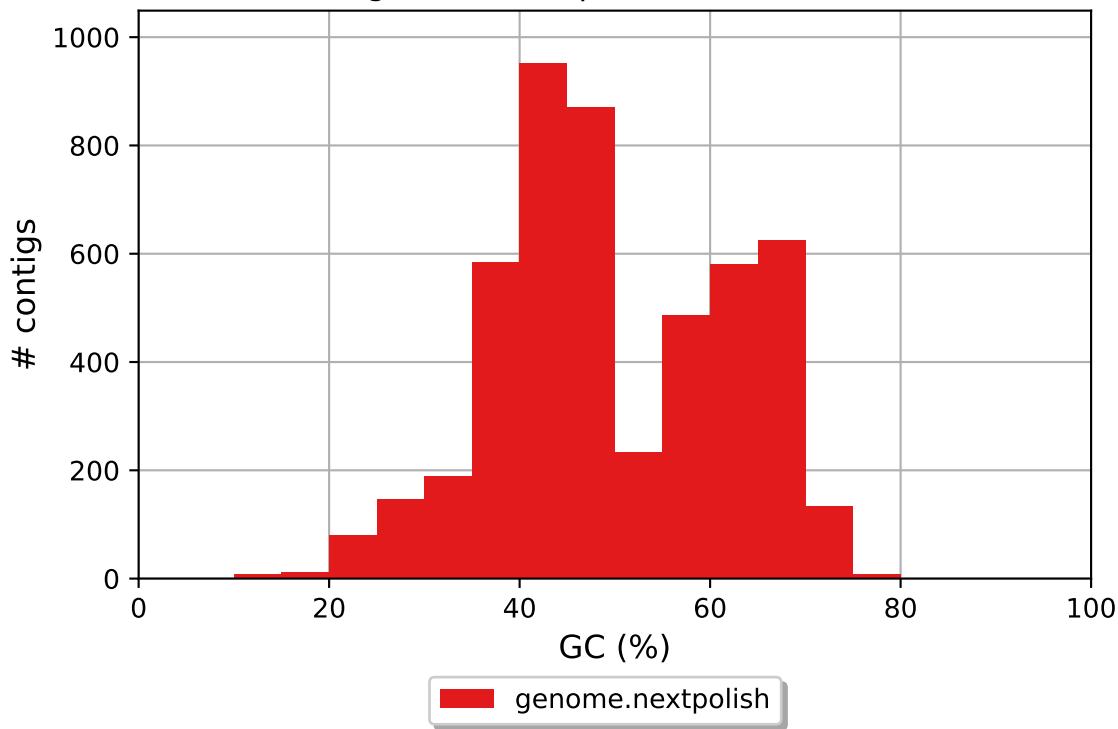


Cumulative length

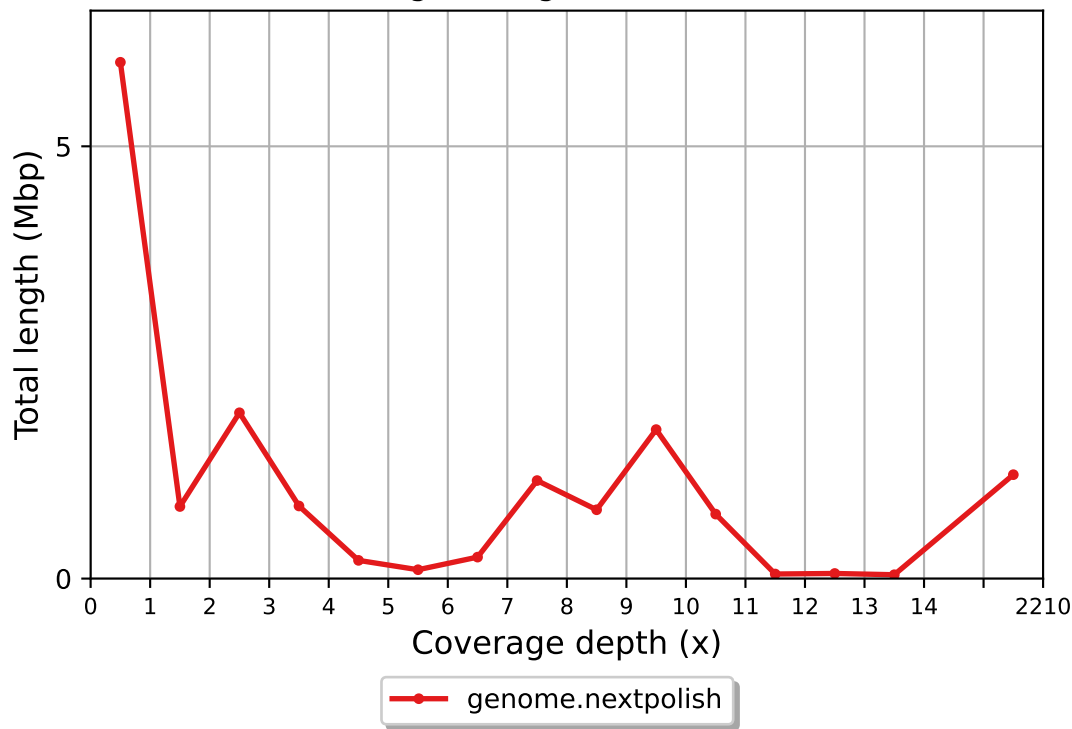




genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

