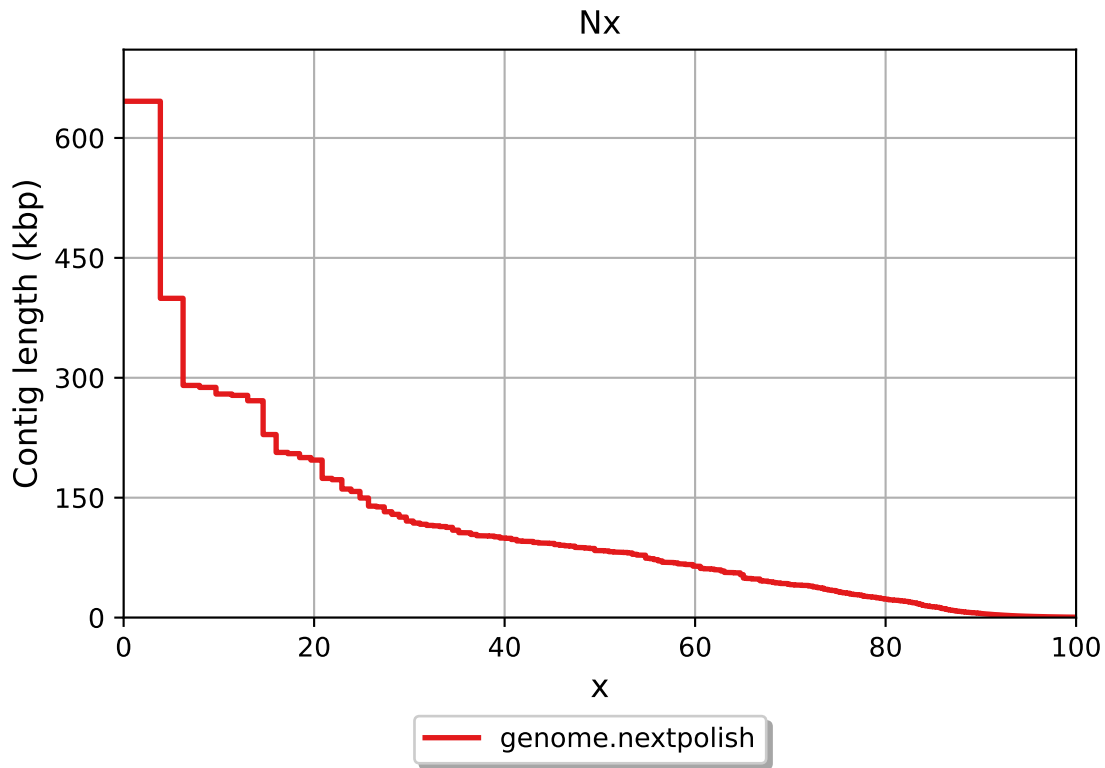


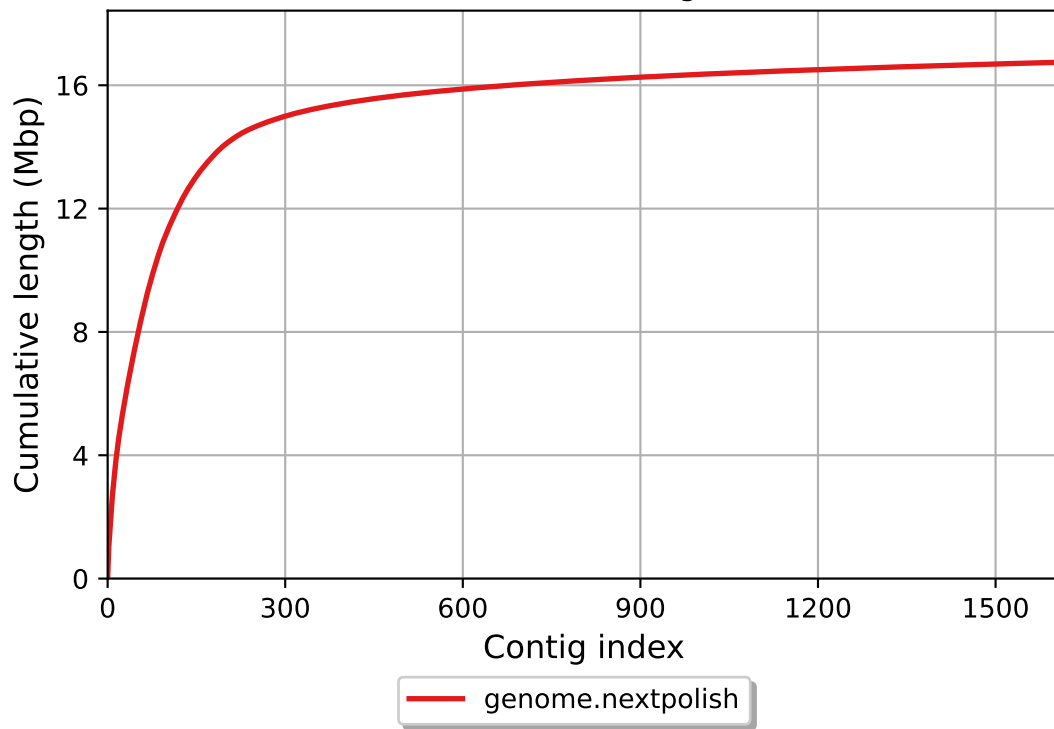
Report

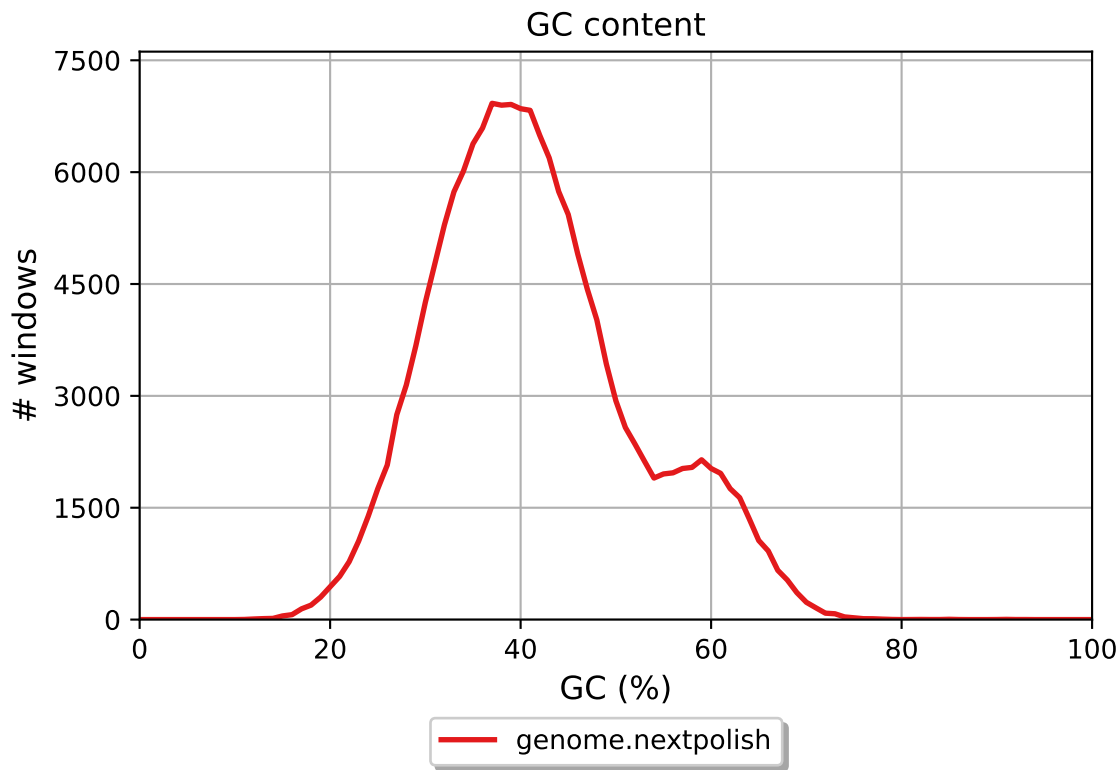
	genome.nextpolish
# contigs (>= 0 bp)	5443
# contigs (>= 1000 bp)	892
# contigs (>= 5000 bp)	317
# contigs (>= 10000 bp)	230
# contigs (>= 25000 bp)	157
# contigs (>= 50000 bp)	93
Total length (>= 0 bp)	17760165
Total length (>= 1000 bp)	16255046
Total length (>= 5000 bp)	15089850
Total length (>= 10000 bp)	14482707
Total length (>= 25000 bp)	13231224
Total length (>= 50000 bp)	10895824
# contigs	1609
Largest contig	645898
Total length	16746011
GC (%)	41.59
N50	83732
N90	5133
auN	122356.8
L50	57
L90	314
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	11 + 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).

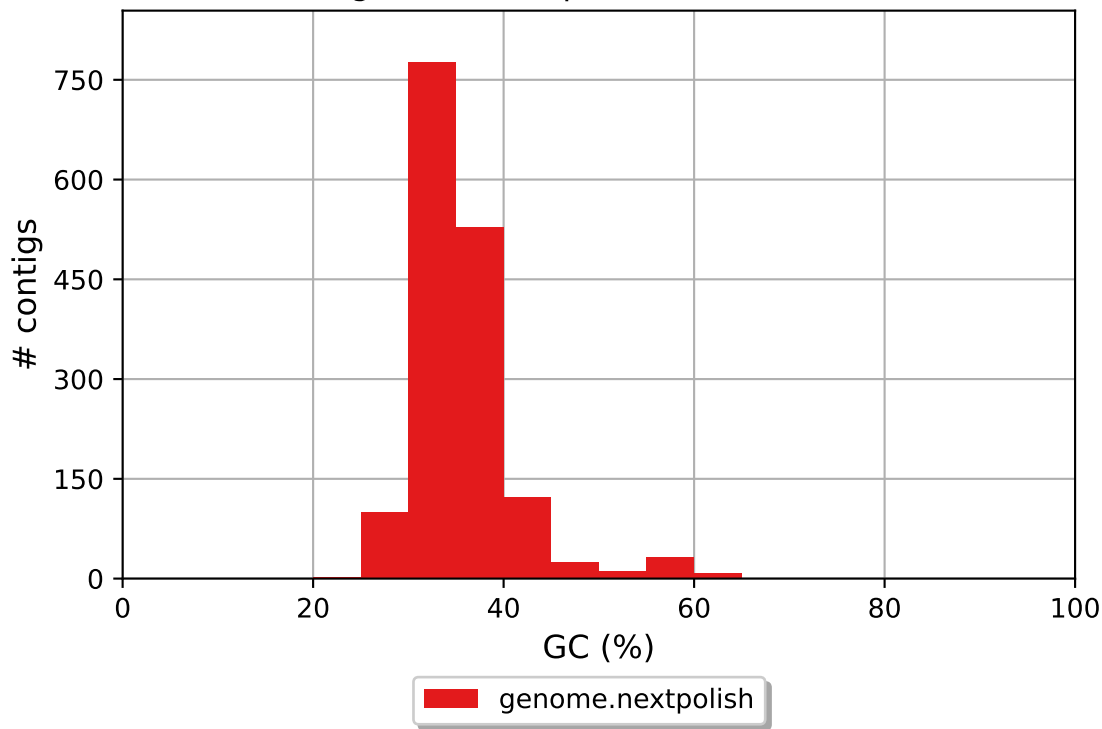


Cumulative length

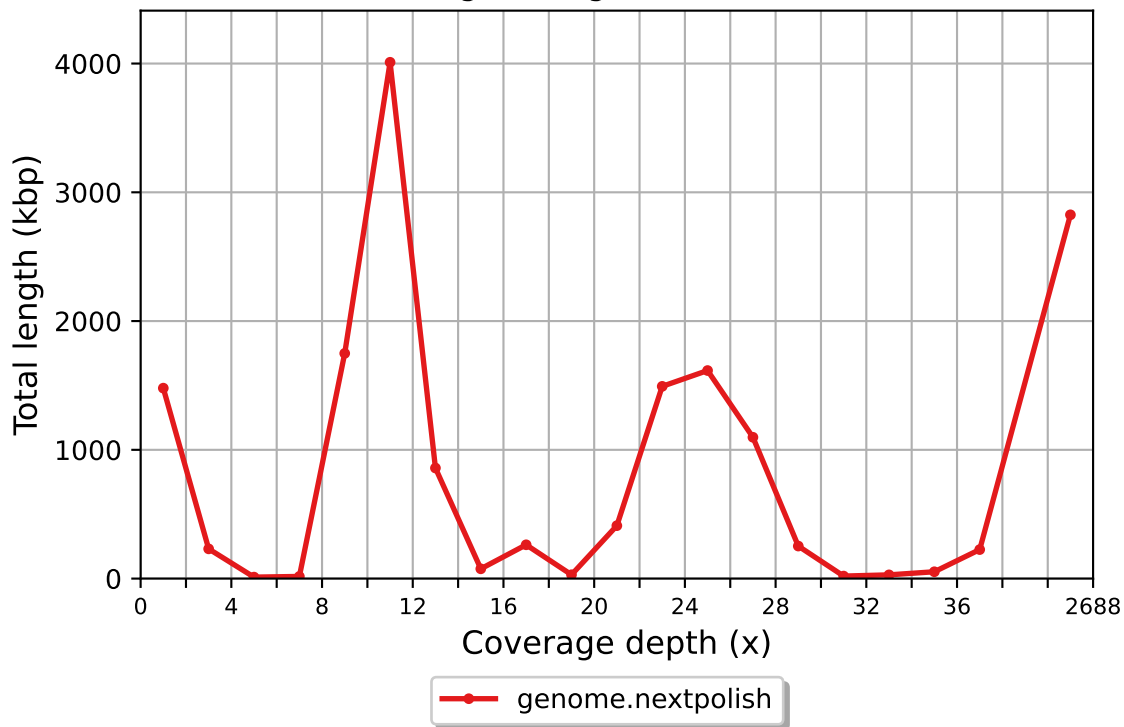




genome.nextpolish GC content



Coverage histogram (bin size: 2x)



genome.nextpolish coverage histogram (bin size: 2x)

