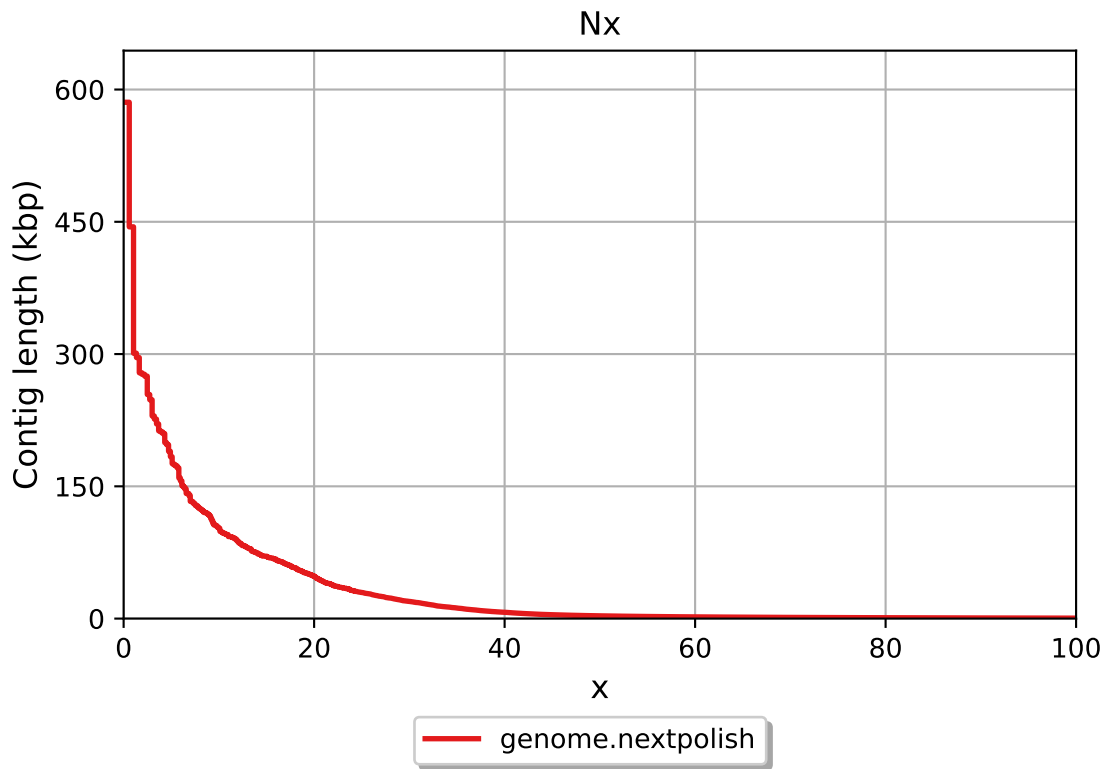
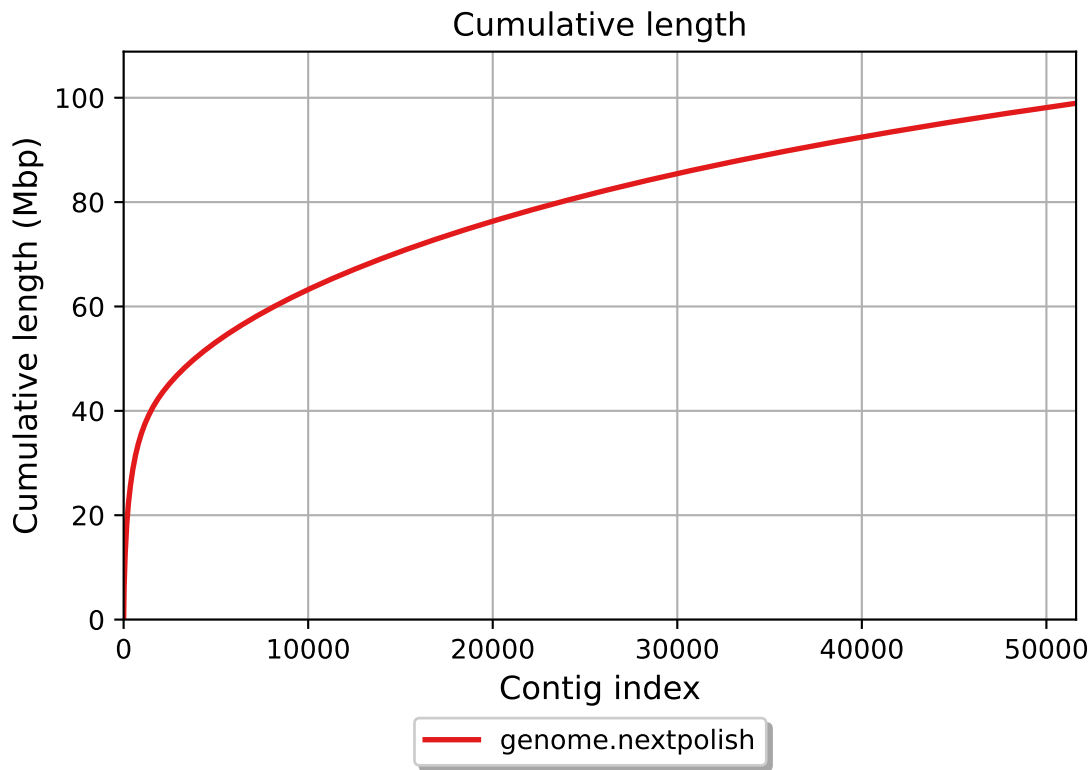


Report

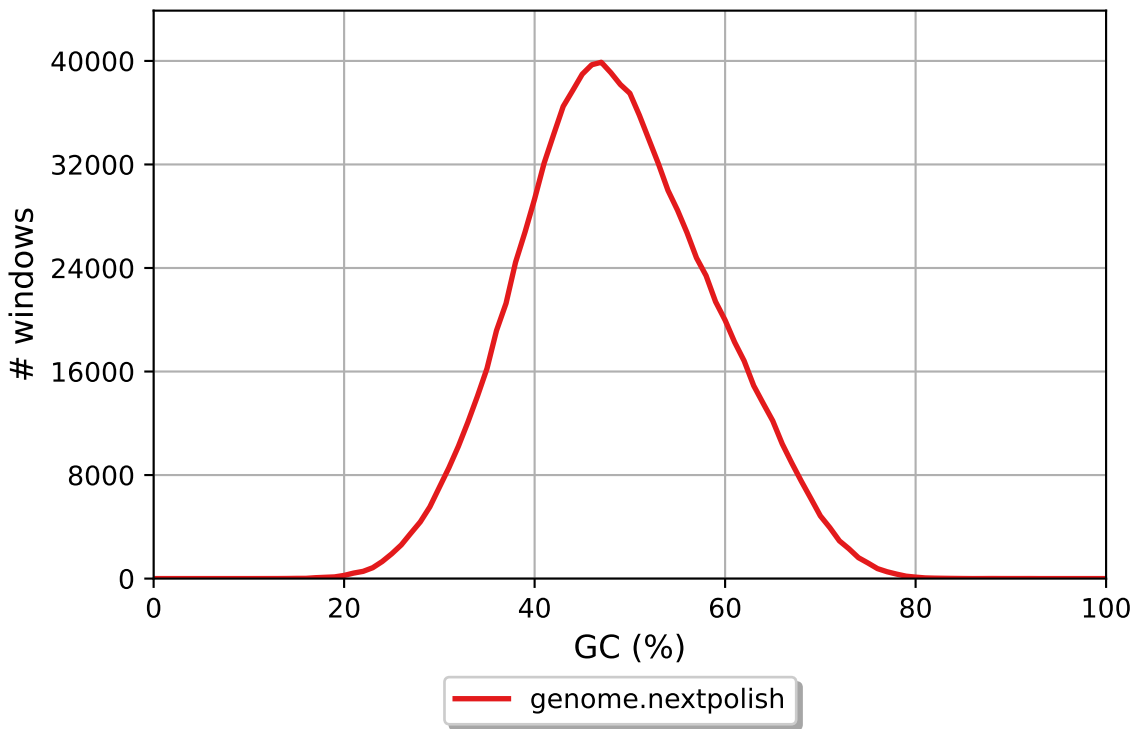
	genome.nextpolish
# contigs (>= 0 bp)	139352
# contigs (>= 1000 bp)	21854
# contigs (>= 5000 bp)	1991
# contigs (>= 10000 bp)	1021
# contigs (>= 25000 bp)	417
# contigs (>= 50000 bp)	191
Total length (>= 0 bp)	125384662
Total length (>= 1000 bp)	78254575
Total length (>= 5000 bp)	42972860
Total length (>= 10000 bp)	36274078
Total length (>= 25000 bp)	26937584
Total length (>= 50000 bp)	19370355
# contigs	51609
Largest contig	585551
Total length	98941228
GC (%)	48.60
N50	3117
N90	700
auN	34470.3
L50	3680
L90	34825
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	24 + 18 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).

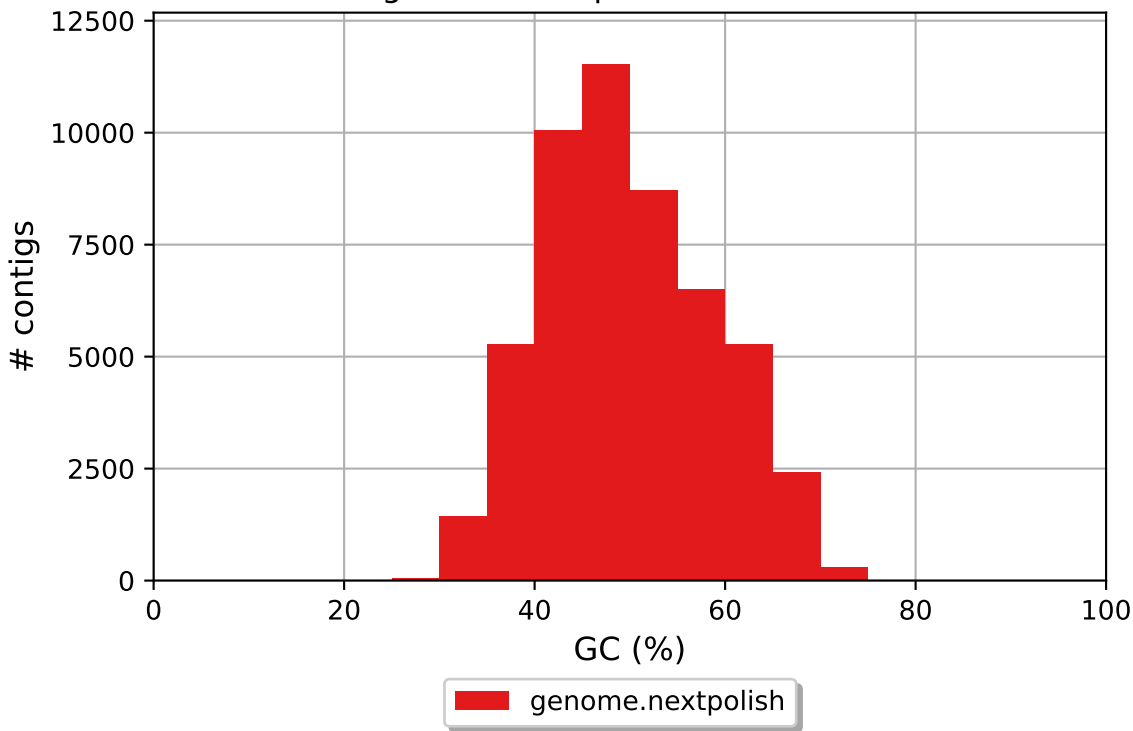




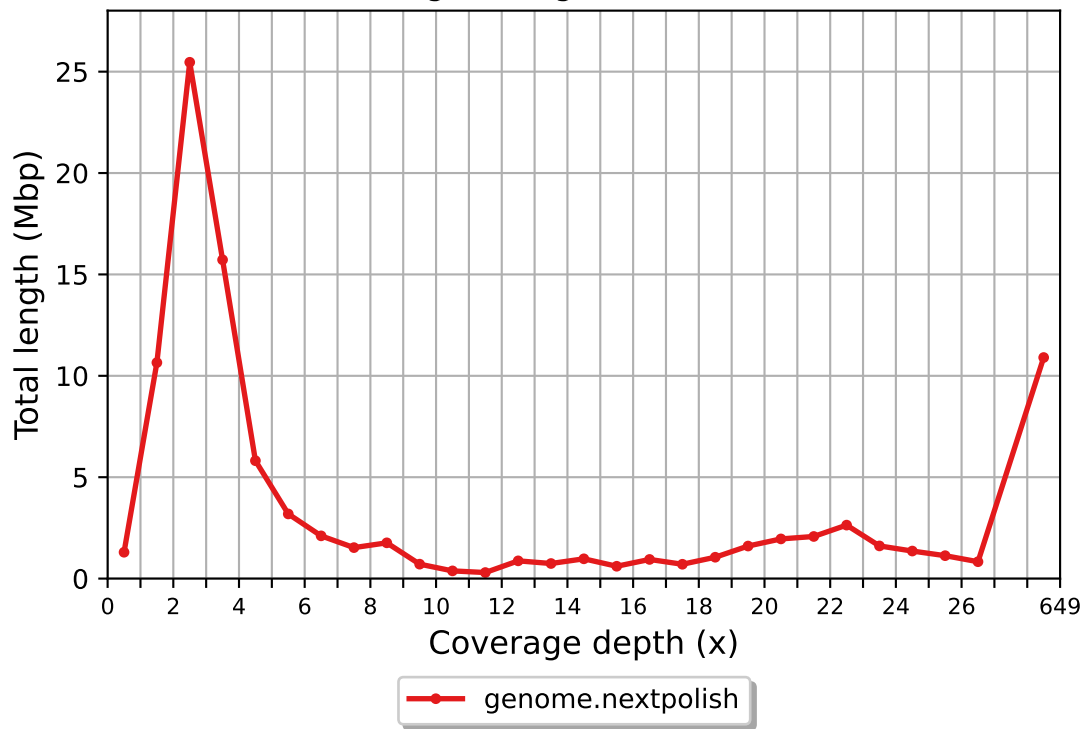
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

