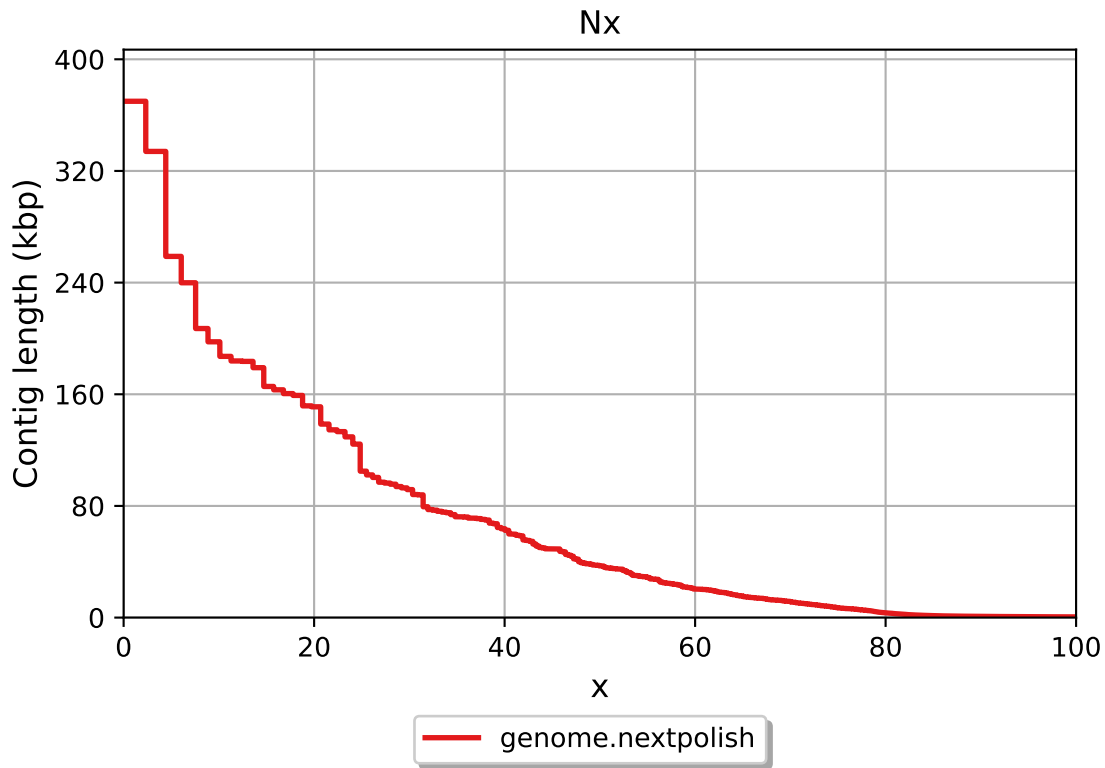


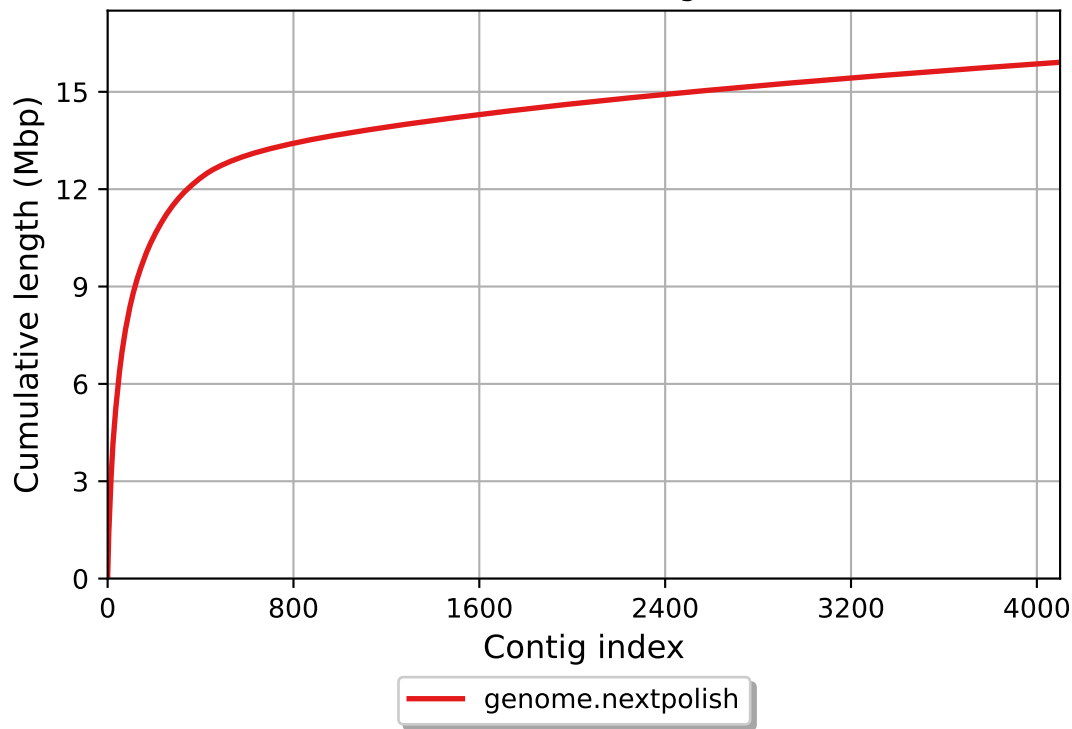
Report

	genome.nextpolish
# contigs (>= 0 bp)	16173
# contigs (>= 1000 bp)	1313
# contigs (>= 5000 bp)	419
# contigs (>= 10000 bp)	269
# contigs (>= 25000 bp)	118
# contigs (>= 50000 bp)	62
Total length (>= 0 bp)	19510555
Total length (>= 1000 bp)	14029543
Total length (>= 5000 bp)	12447777
Total length (>= 10000 bp)	11386791
Total length (>= 25000 bp)	9009700
Total length (>= 50000 bp)	6990148
# contigs	4100
Largest contig	369920
Total length	15910652
GC (%)	48.56
N50	37076
N90	881
auN	74831.0
L50	85
L90	1624
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 1 part

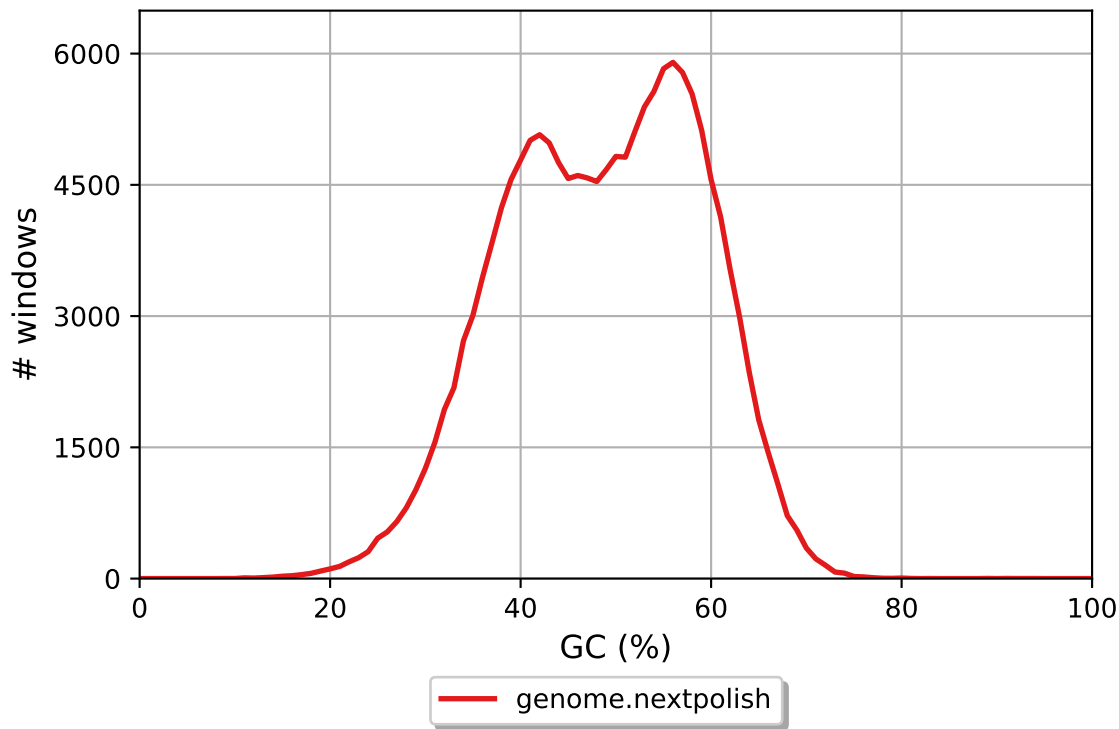
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



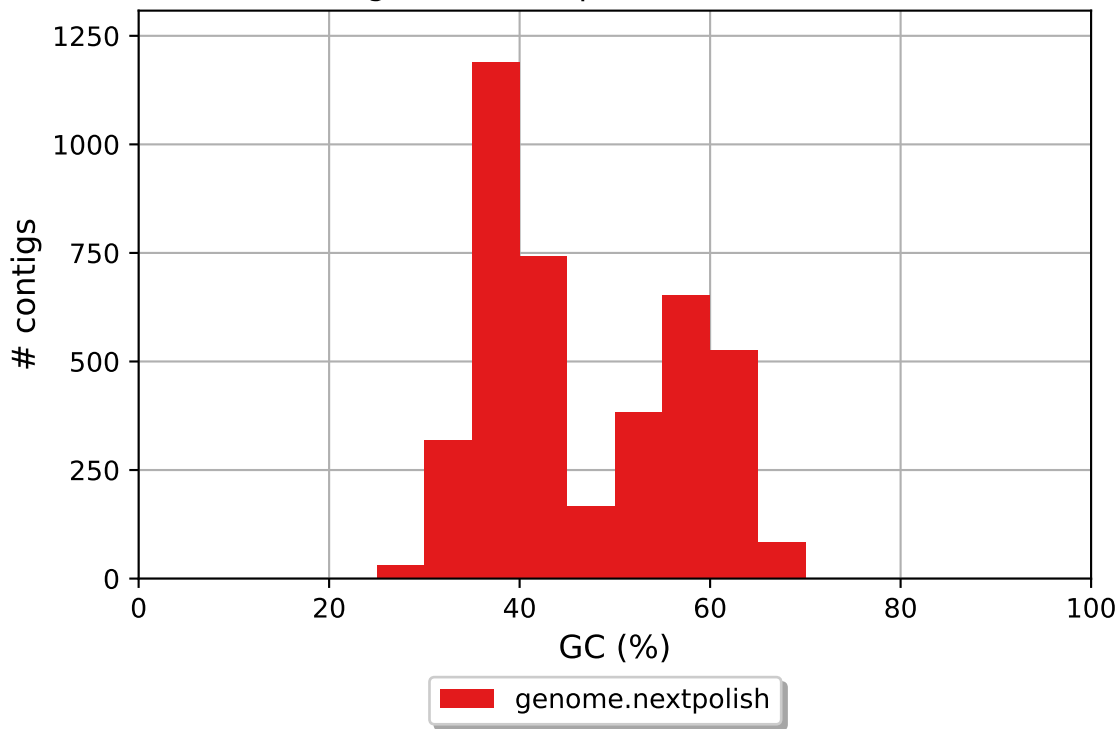
Cumulative length



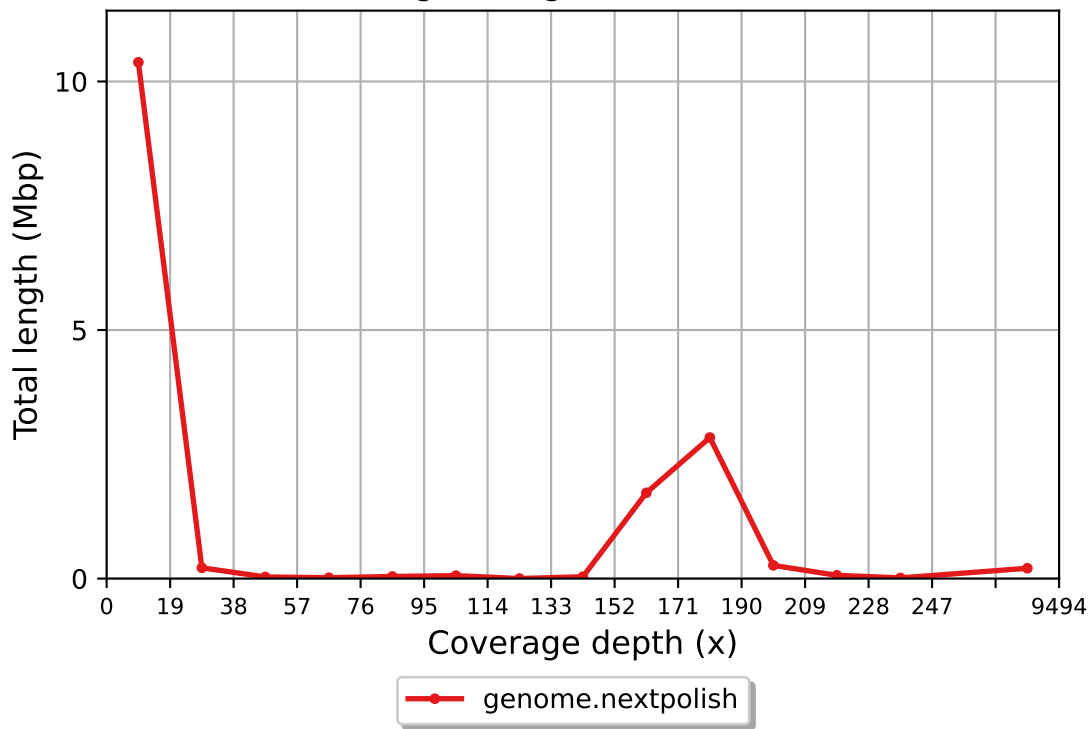
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 19x)



genome.nextpolish coverage histogram (bin size: 19x)

