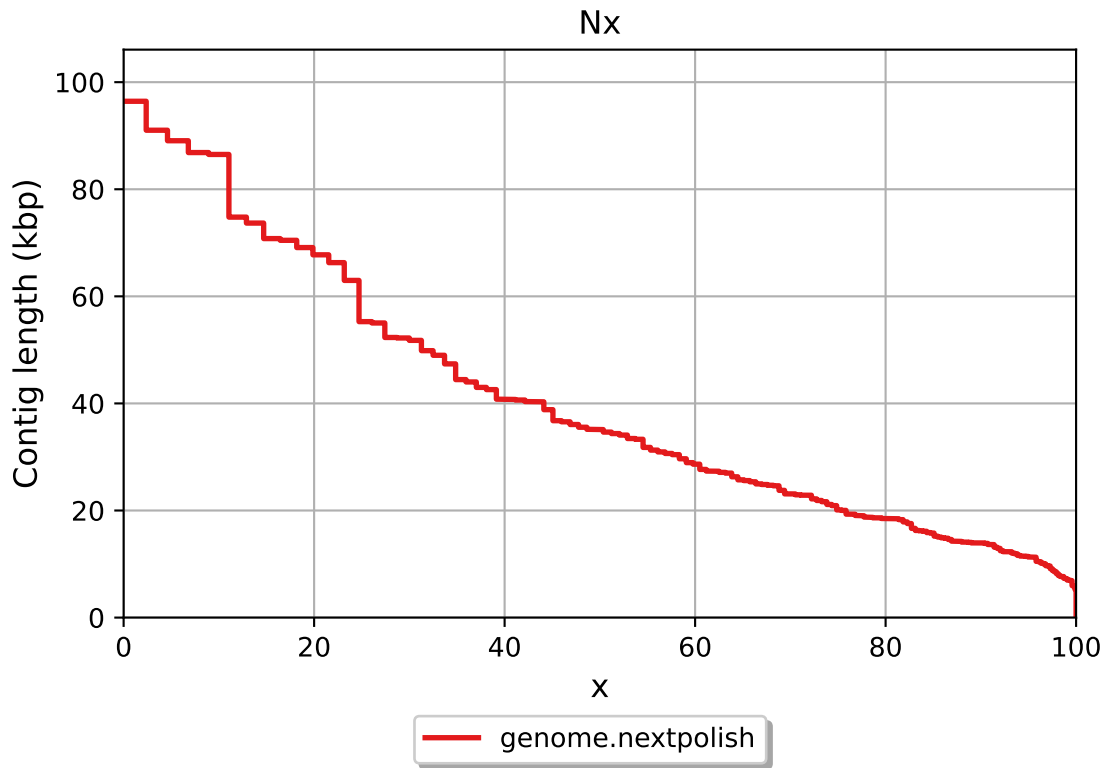
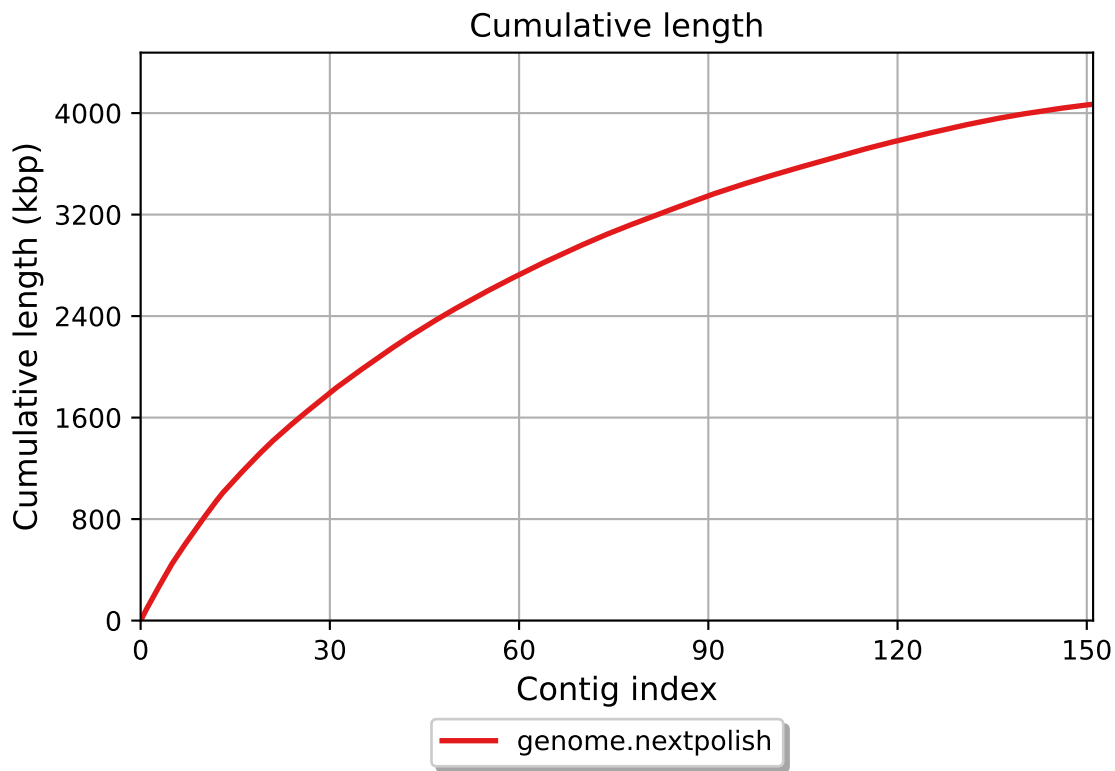


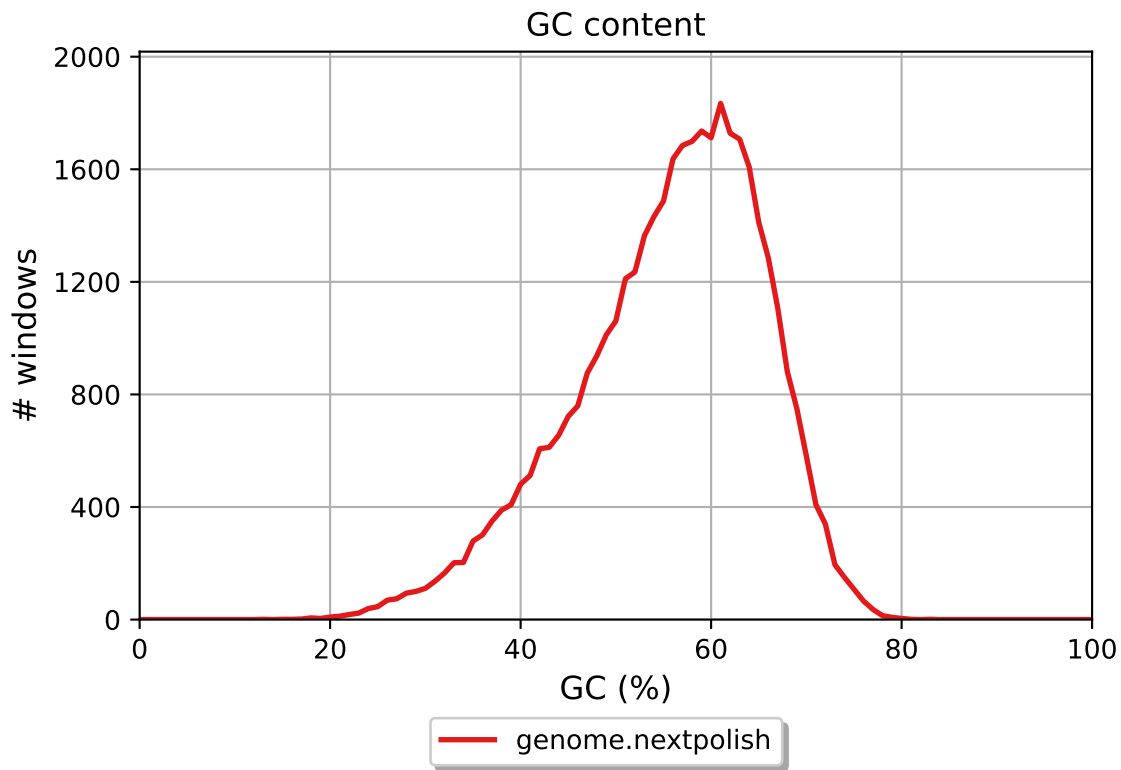
Report

	genome.nextpolish
# contigs (>= 0 bp)	151
# contigs (>= 1000 bp)	151
# contigs (>= 5000 bp)	151
# contigs (>= 10000 bp)	134
# contigs (>= 25000 bp)	59
# contigs (>= 50000 bp)	18
Total length (>= 0 bp)	4069592
Total length (>= 1000 bp)	4069592
Total length (>= 5000 bp)	4069592
Total length (>= 10000 bp)	3939728
Total length (>= 25000 bp)	2701107
Total length (>= 50000 bp)	1272340
# contigs	151
Largest contig	96432
Total length	4069592
GC (%)	55.50
N50	35114
N90	13944
auN	41271.1
L50	37
L90	111
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	13 + 0 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 GC content

