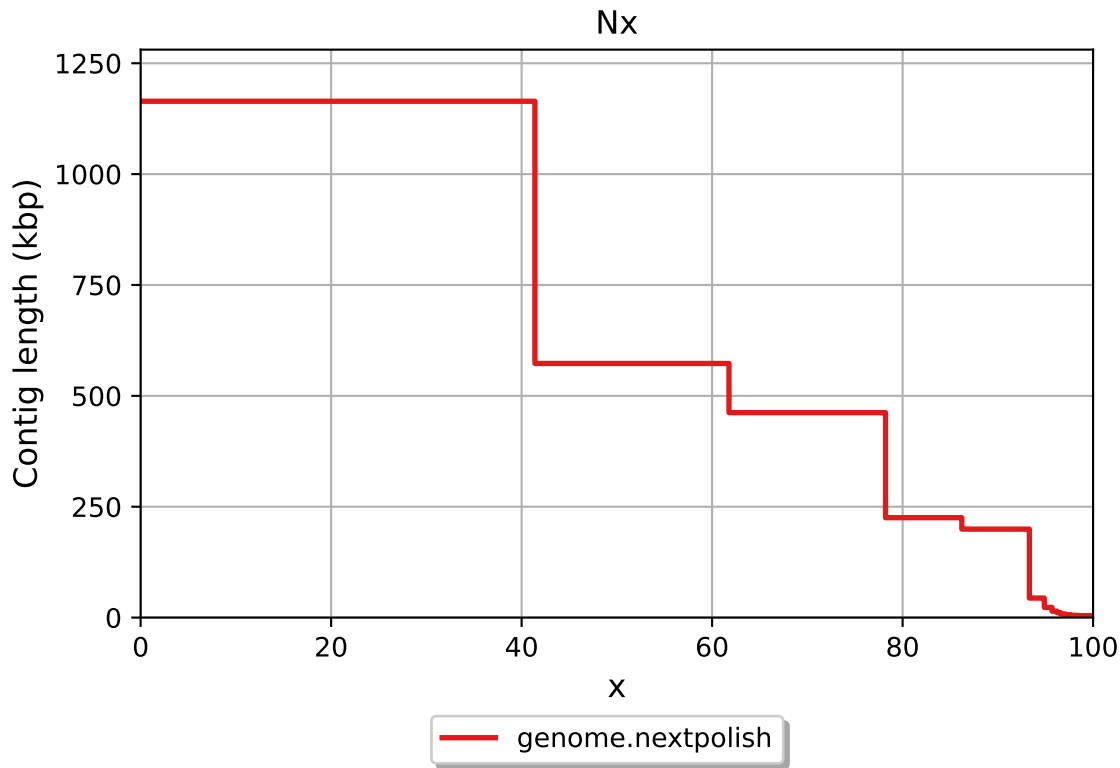
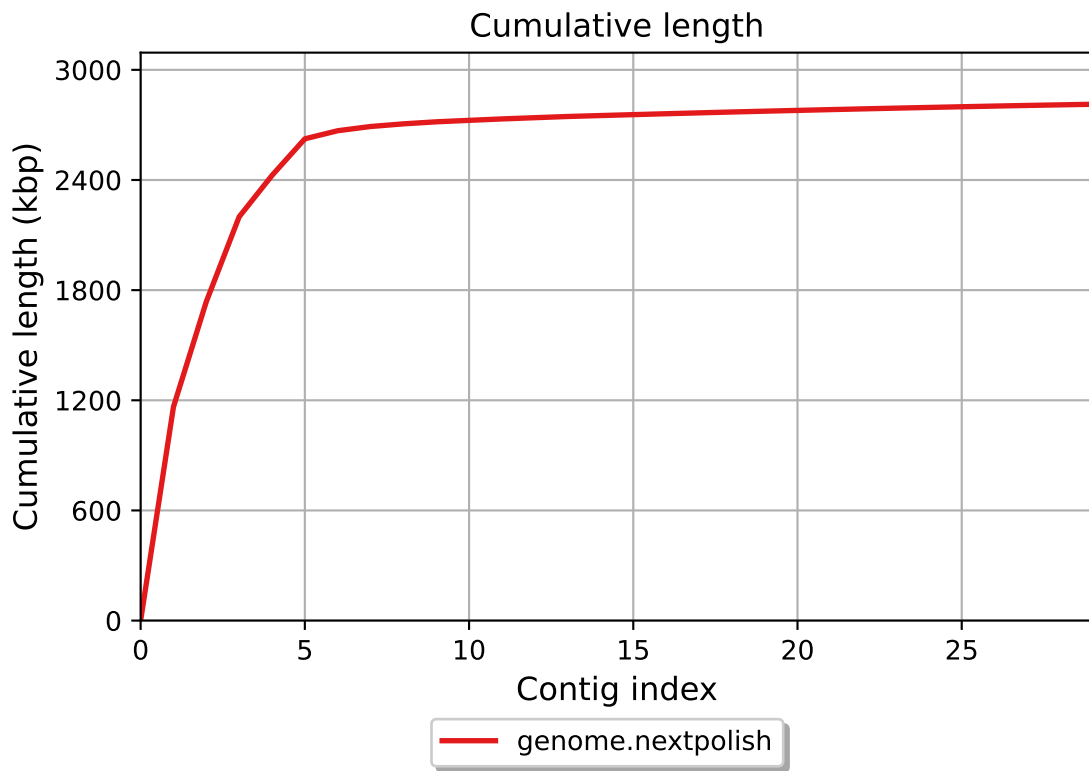


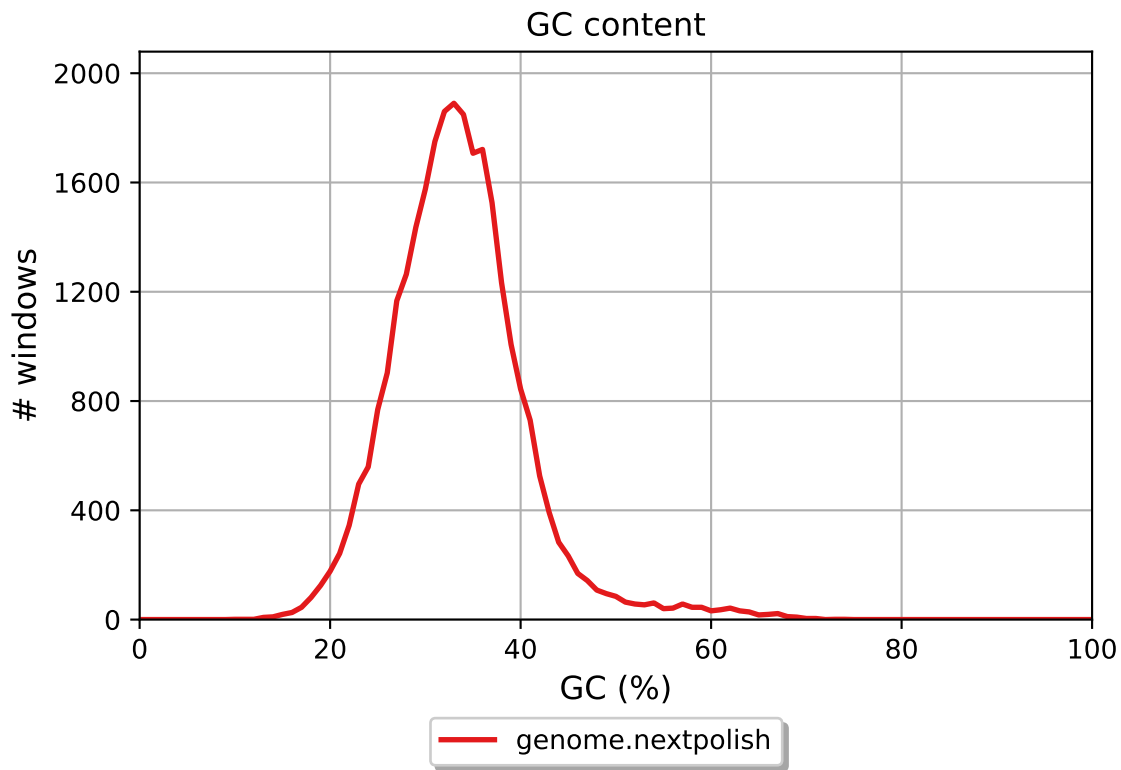
Report

	genome.nextpolish
# contigs (>= 0 bp)	29
# contigs (>= 1000 bp)	29
# contigs (>= 5000 bp)	15
# contigs (>= 10000 bp)	9
# contigs (>= 25000 bp)	6
# contigs (>= 50000 bp)	5
Total length (>= 0 bp)	2812436
Total length (>= 1000 bp)	2812436
Total length (>= 5000 bp)	2755966
Total length (>= 10000 bp)	2716856
Total length (>= 25000 bp)	2668304
Total length (>= 50000 bp)	2624247
# contigs	29
Largest contig	1164283
Total length	2812436
GC (%)	33.58
N50	573044
N90	199487
auN	708051.4
L50	2
L90	5
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 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).







genome.nextpolish GC content

