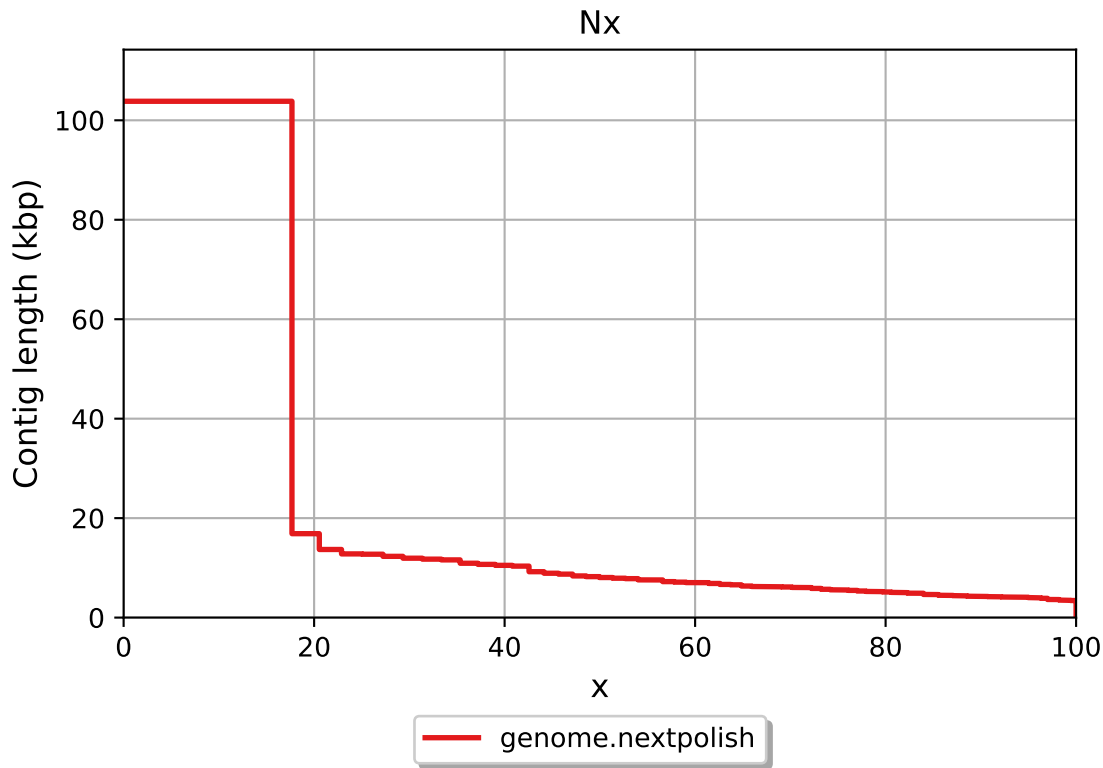
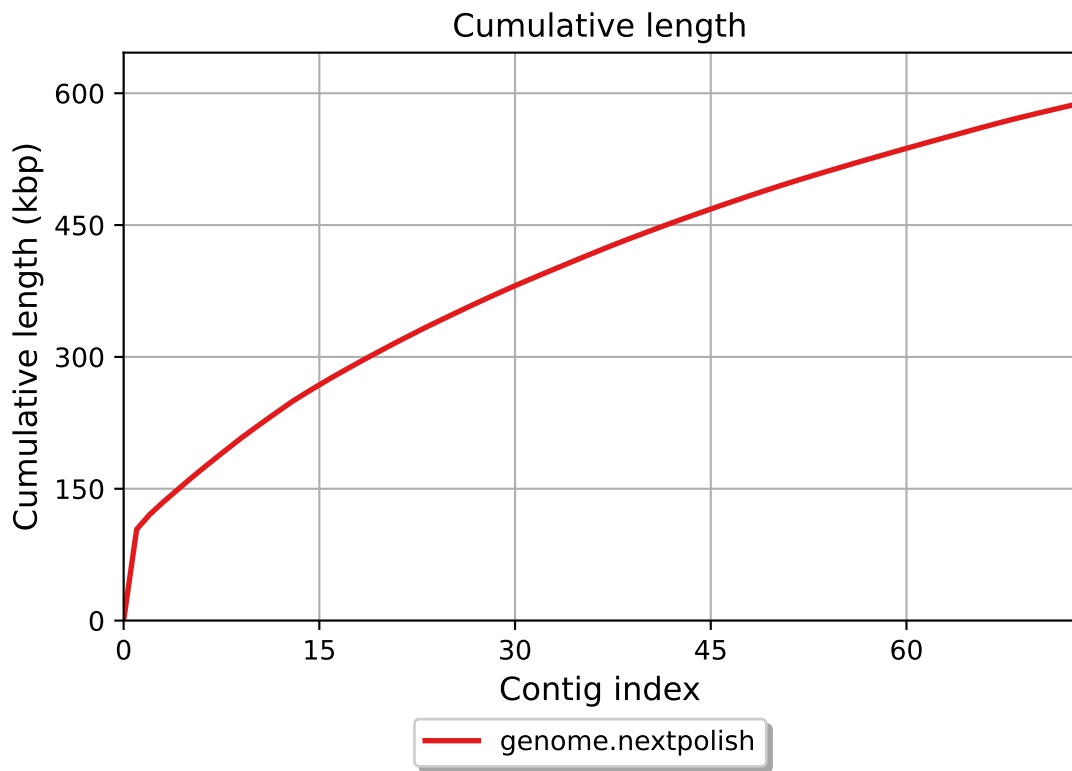


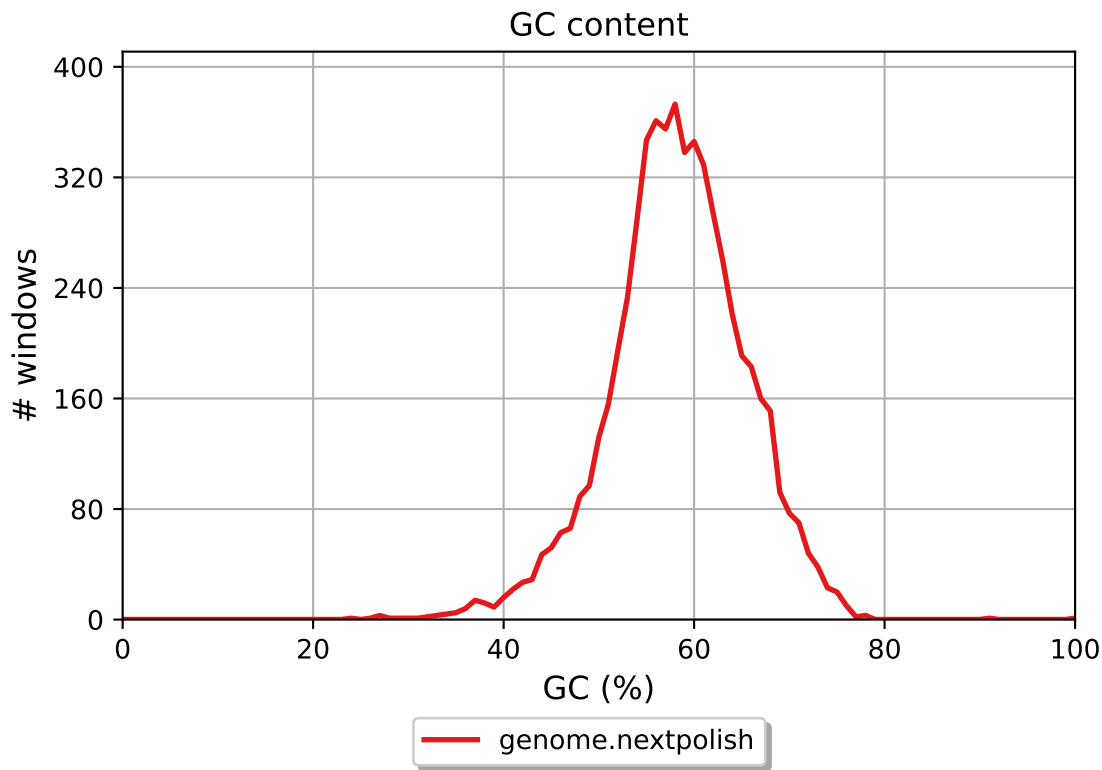
Report

	genome.nextpolish
# contigs (>= 0 bp)	73
# contigs (>= 1000 bp)	73
# contigs (>= 5000 bp)	48
# contigs (>= 10000 bp)	13
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	587458
Total length (>= 1000 bp)	587458
Total length (>= 5000 bp)	483481
Total length (>= 10000 bp)	250035
Total length (>= 25000 bp)	103825
Total length (>= 50000 bp)	103825
# contigs	73
Largest contig	103825
Total length	587458
GC (%)	58.23
N50	8057
N90	4251
auN	24919.9
L50	19
L90	58
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 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).







genome.nextpolish GC content

