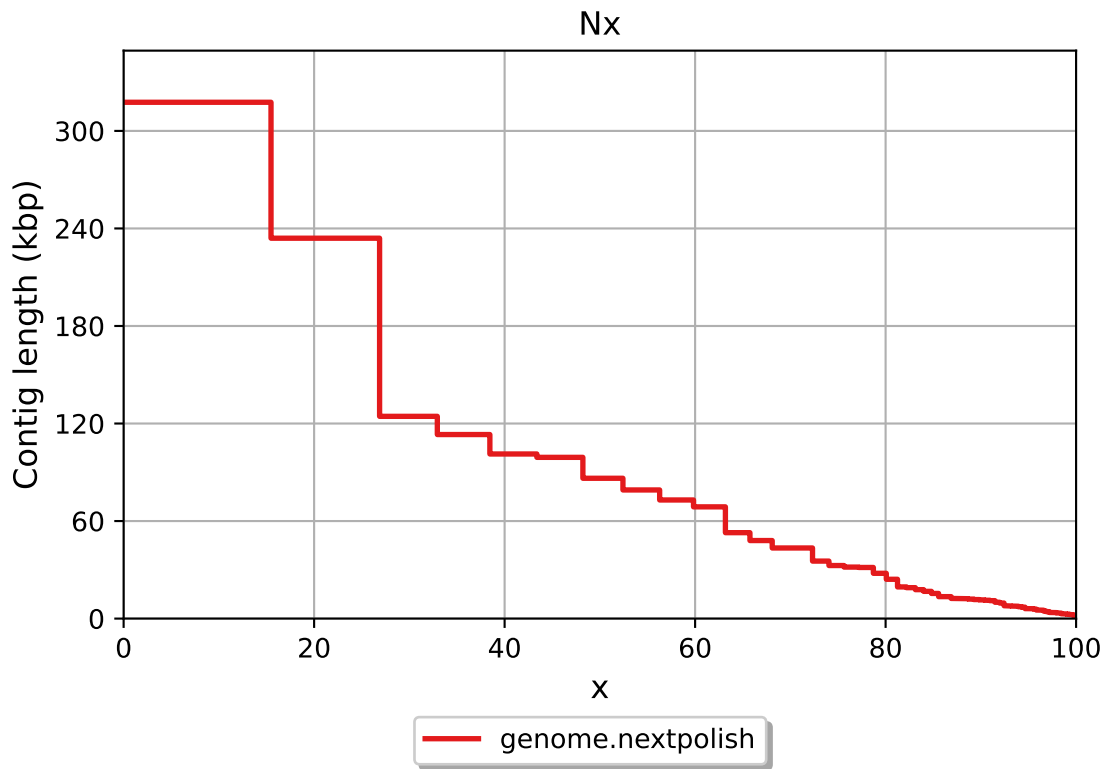
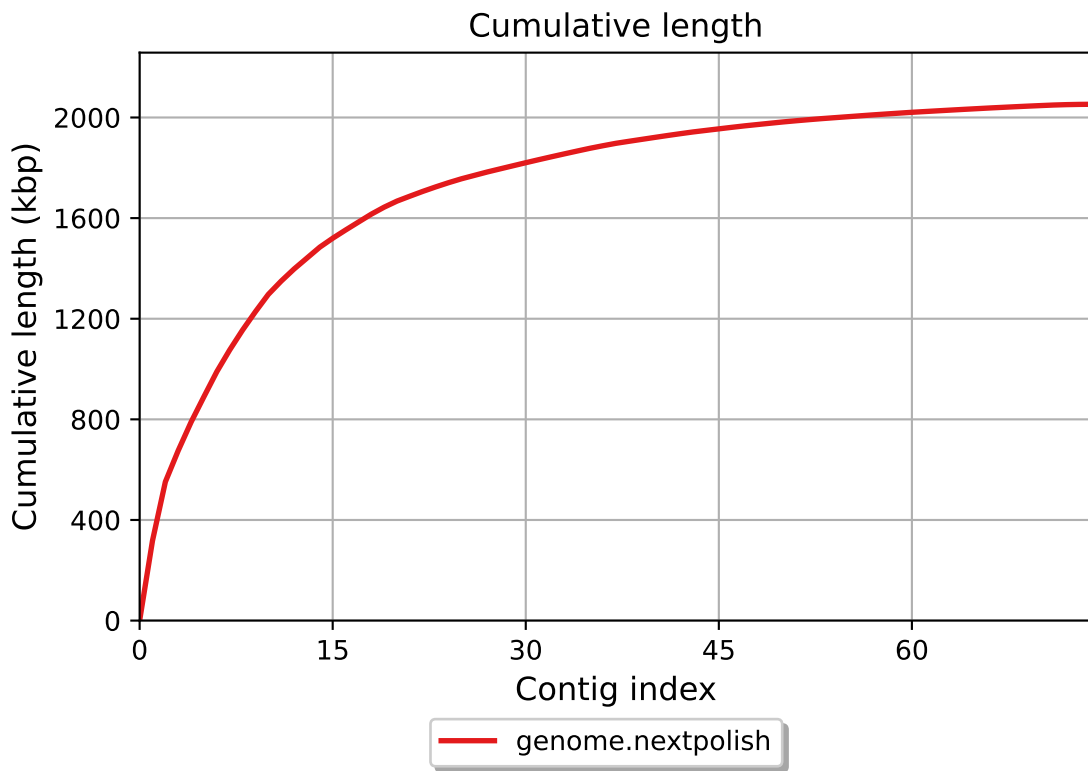


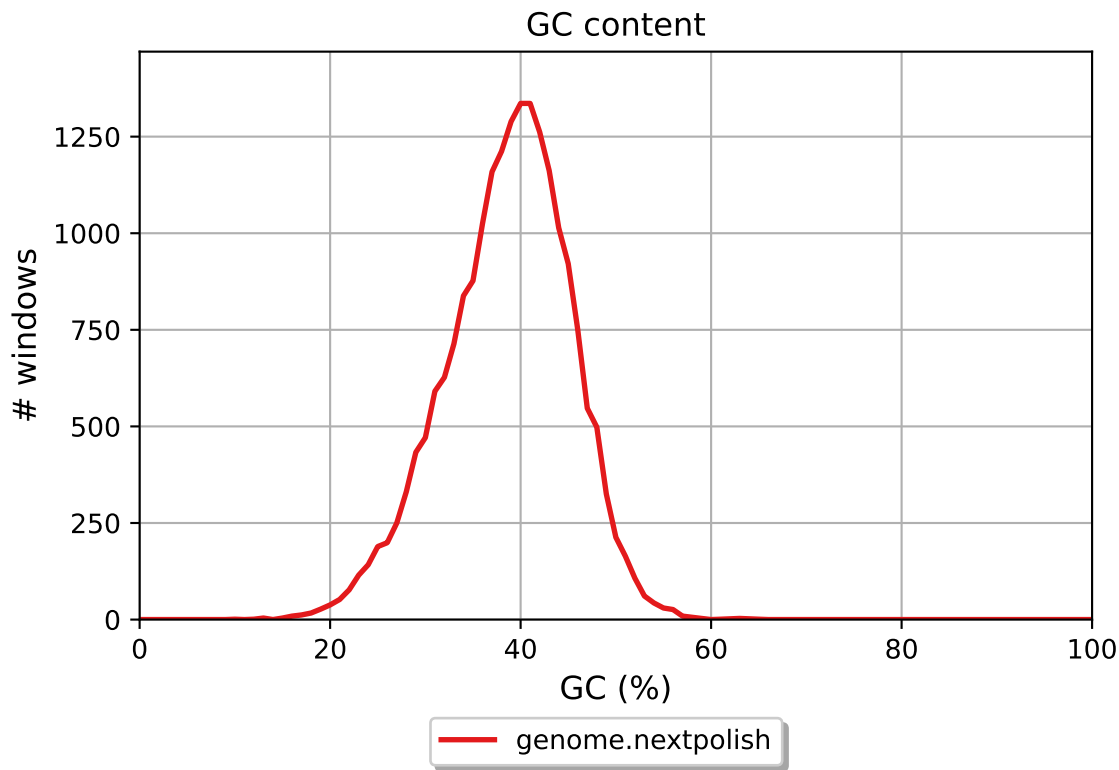
Report

	genome.nextpolish
# contigs (>= 0 bp)	74
# contigs (>= 1000 bp)	72
# contigs (>= 5000 bp)	50
# contigs (>= 10000 bp)	36
# contigs (>= 25000 bp)	19
# contigs (>= 50000 bp)	11
Total length (>= 0 bp)	2052579
Total length (>= 1000 bp)	2051103
Total length (>= 5000 bp)	1982385
Total length (>= 10000 bp)	1887767
Total length (>= 25000 bp)	1643566
Total length (>= 50000 bp)	1349704
# contigs	74
Largest contig	317621
Total length	2052579
GC (%)	38.63
N50	86318
N90	11493
auN	120033.4
L50	7
L90	33
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 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

