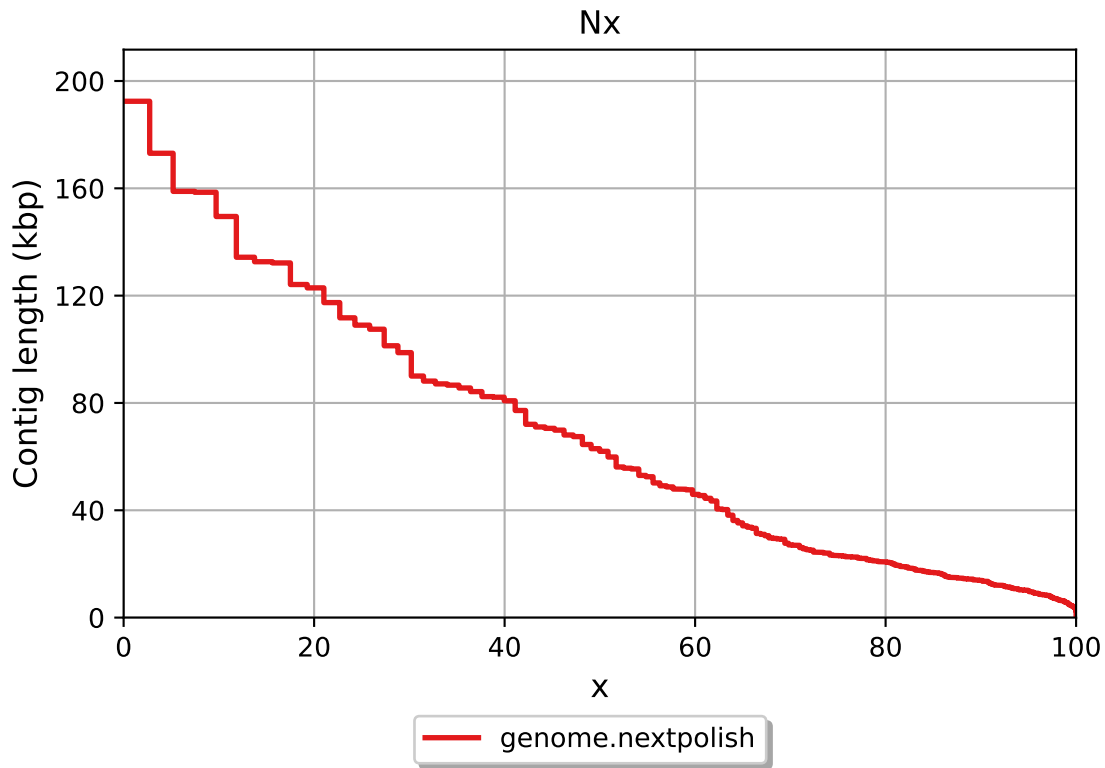
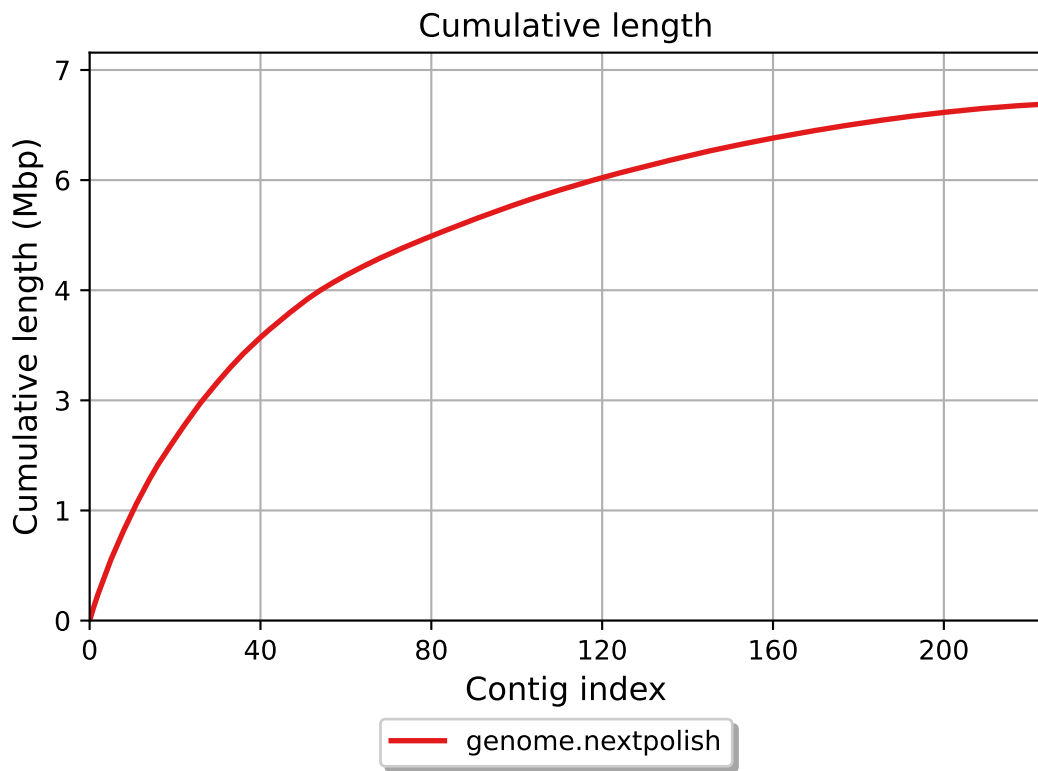


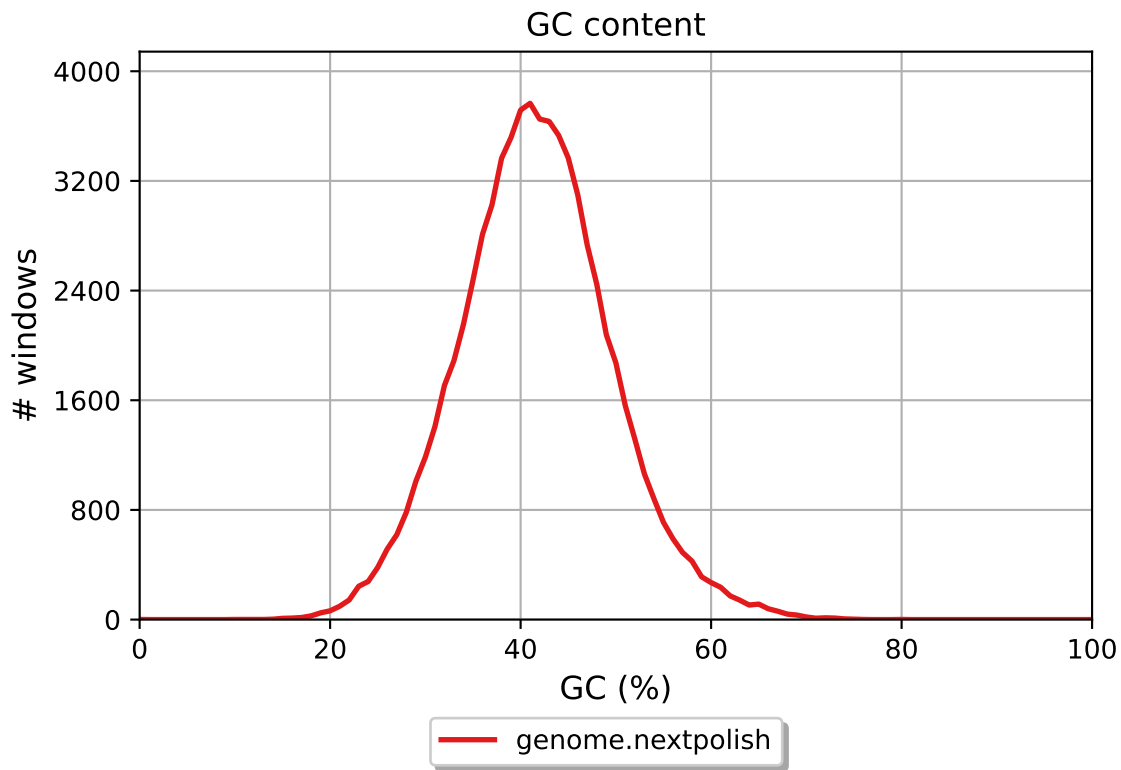
Report

	genome.nextpolish
# contigs (>= 0 bp)	223
# contigs (>= 1000 bp)	223
# contigs (>= 5000 bp)	209
# contigs (>= 10000 bp)	171
# contigs (>= 25000 bp)	74
# contigs (>= 50000 bp)	42
Total length (>= 0 bp)	7033138
Total length (>= 1000 bp)	7033138
Total length (>= 5000 bp)	6975686
Total length (>= 10000 bp)	6687217
Total length (>= 25000 bp)	5093112
Total length (>= 50000 bp)	3959668
# contigs	223
Largest contig	192445
Total length	7033138
GC (%)	41.62
N50	61967
N90	13811
auN	71019.5
L50	35
L90	141
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	19 + 4 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

