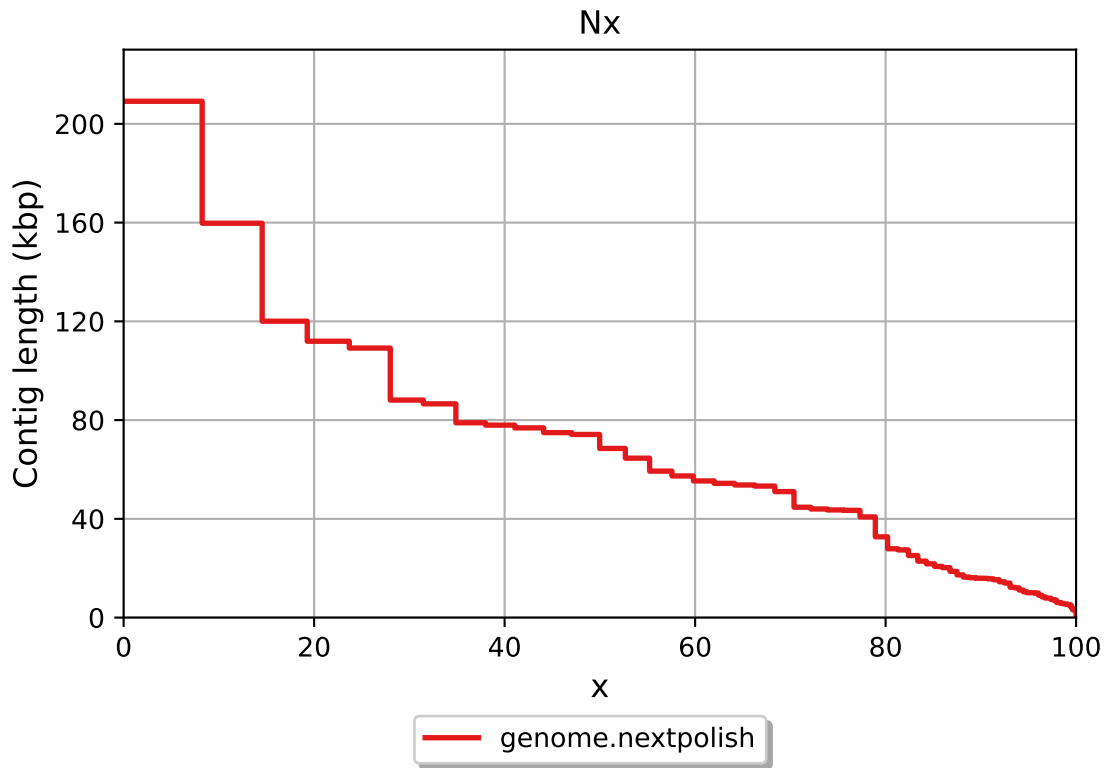
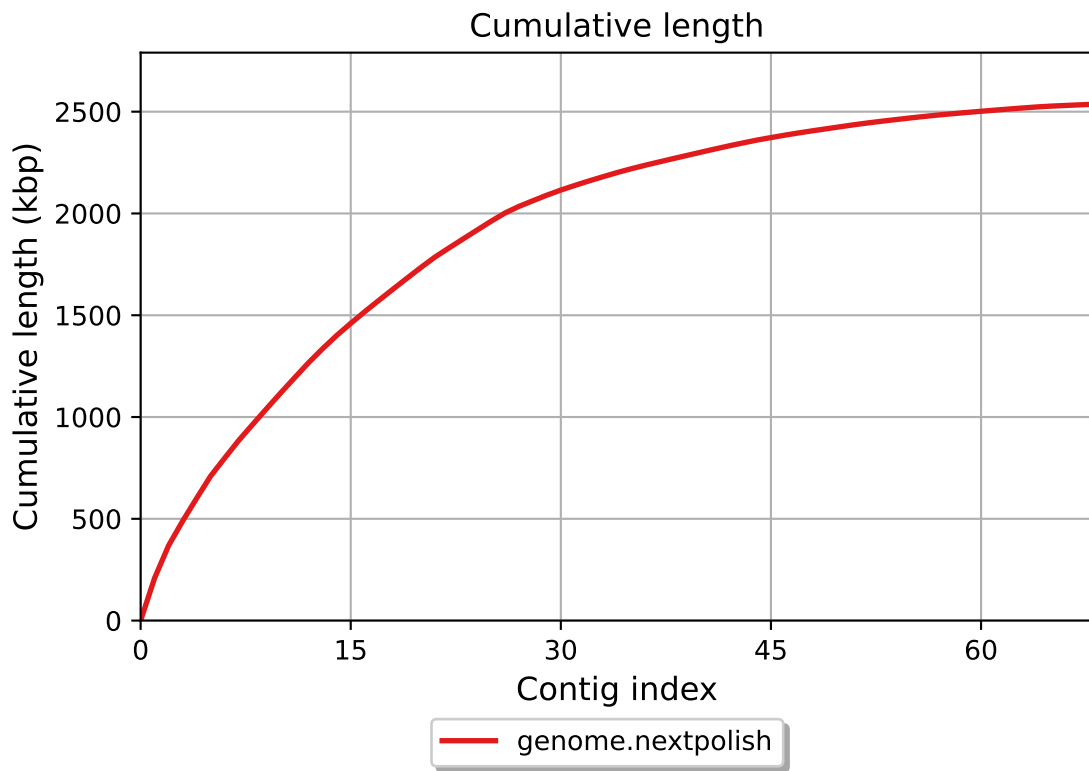


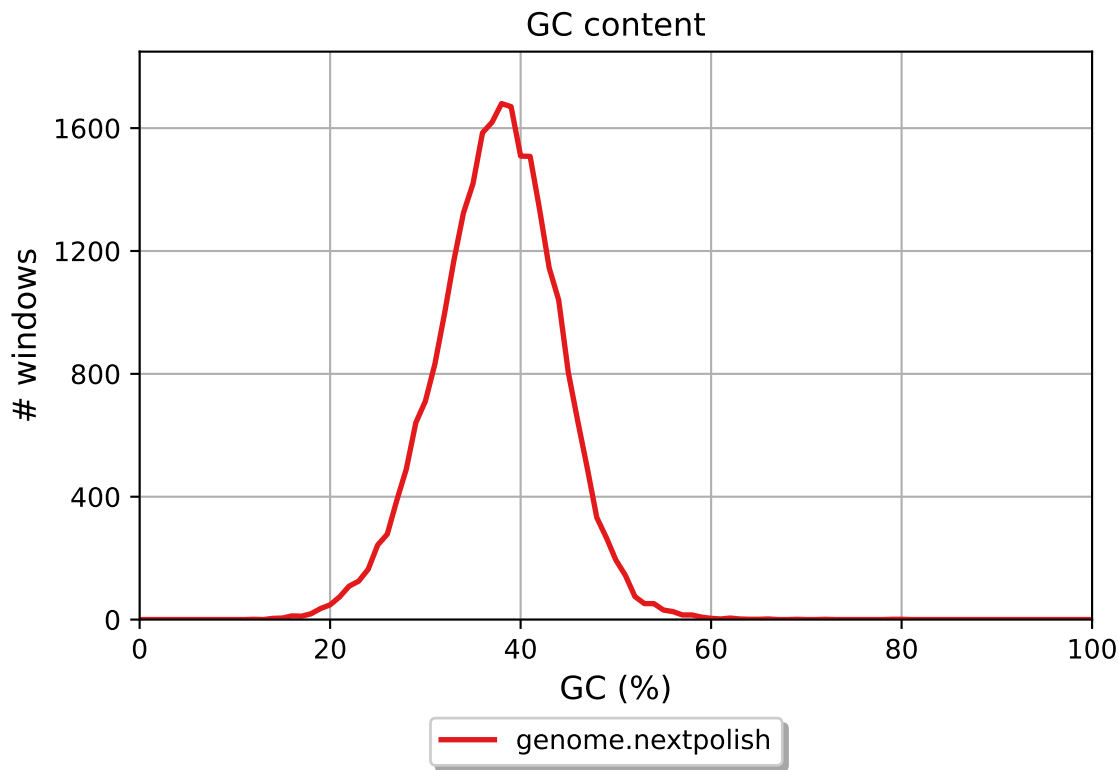
# Report

	genome.nextpolish
# contigs (>= 0 bp)	68
# contigs (>= 1000 bp)	68
# contigs (>= 5000 bp)	64
# contigs (>= 10000 bp)	50
# contigs (>= 25000 bp)	30
# contigs (>= 50000 bp)	21
Total length (>= 0 bp)	2536390
Total length (>= 1000 bp)	2536390
Total length (>= 5000 bp)	2523342
Total length (>= 10000 bp)	2426503
Total length (>= 25000 bp)	2114876
Total length (>= 50000 bp)	1785172
# contigs	68
Largest contig	209151
Total length	2536390
GC (%)	37.54
N50	68525
N90	15954
auN	79345.0
L50	13
L90	39
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 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

