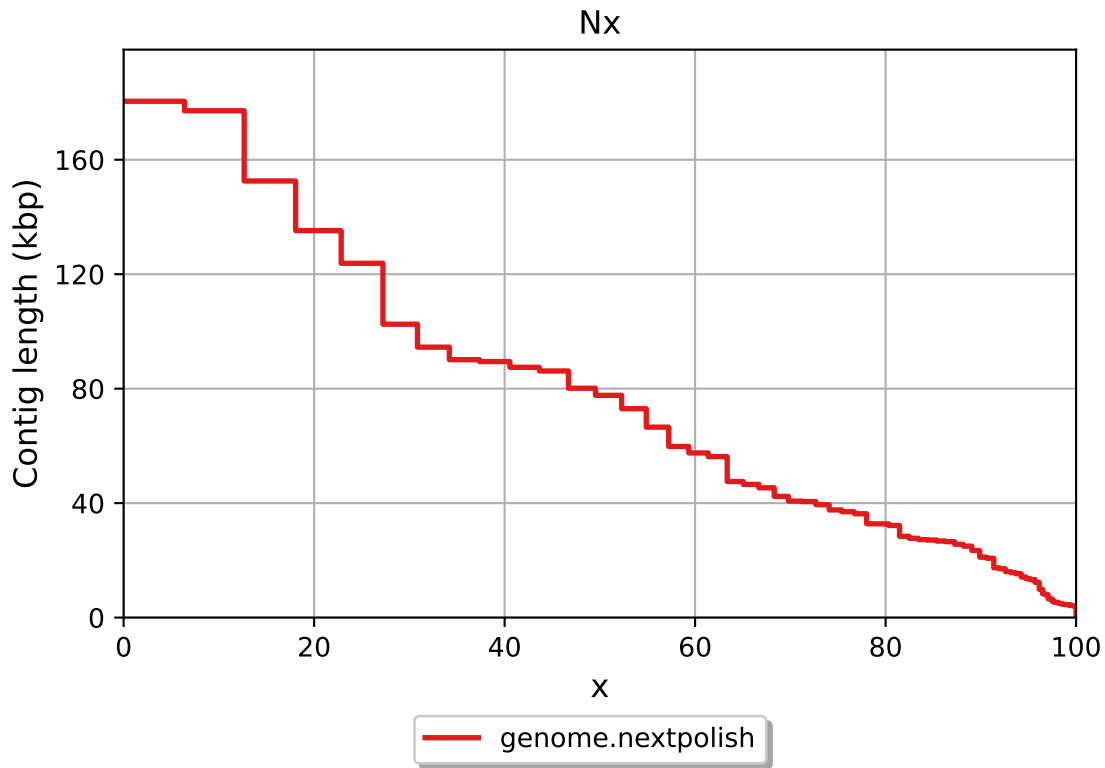
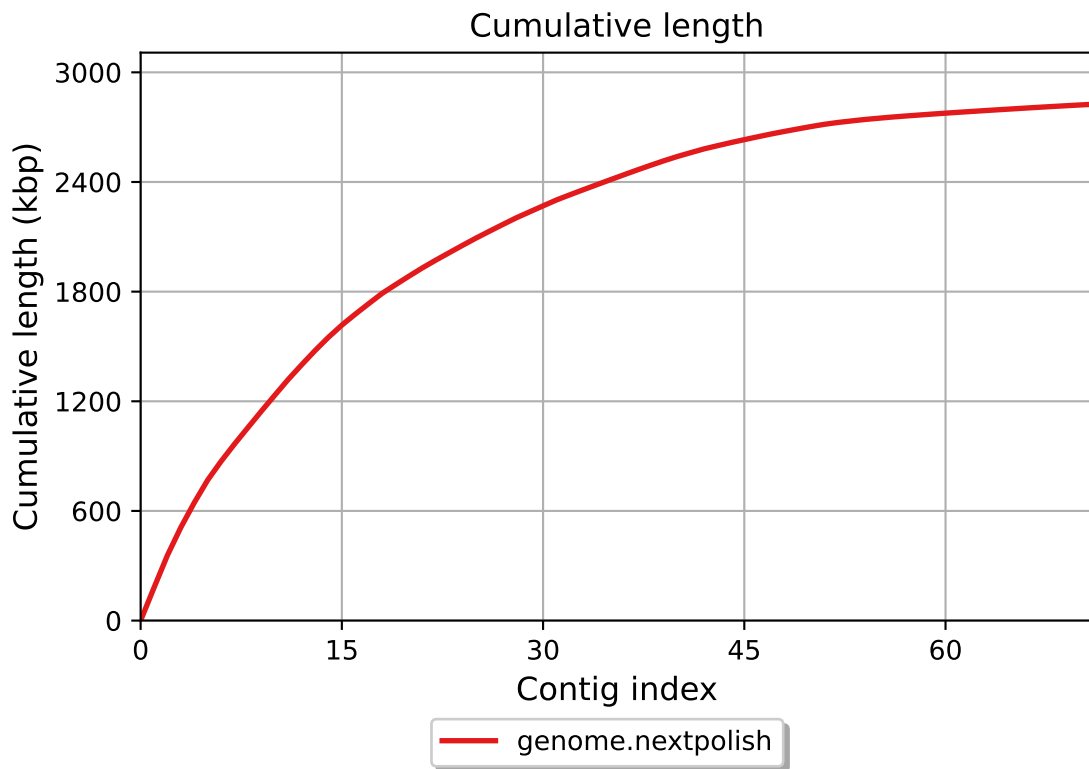


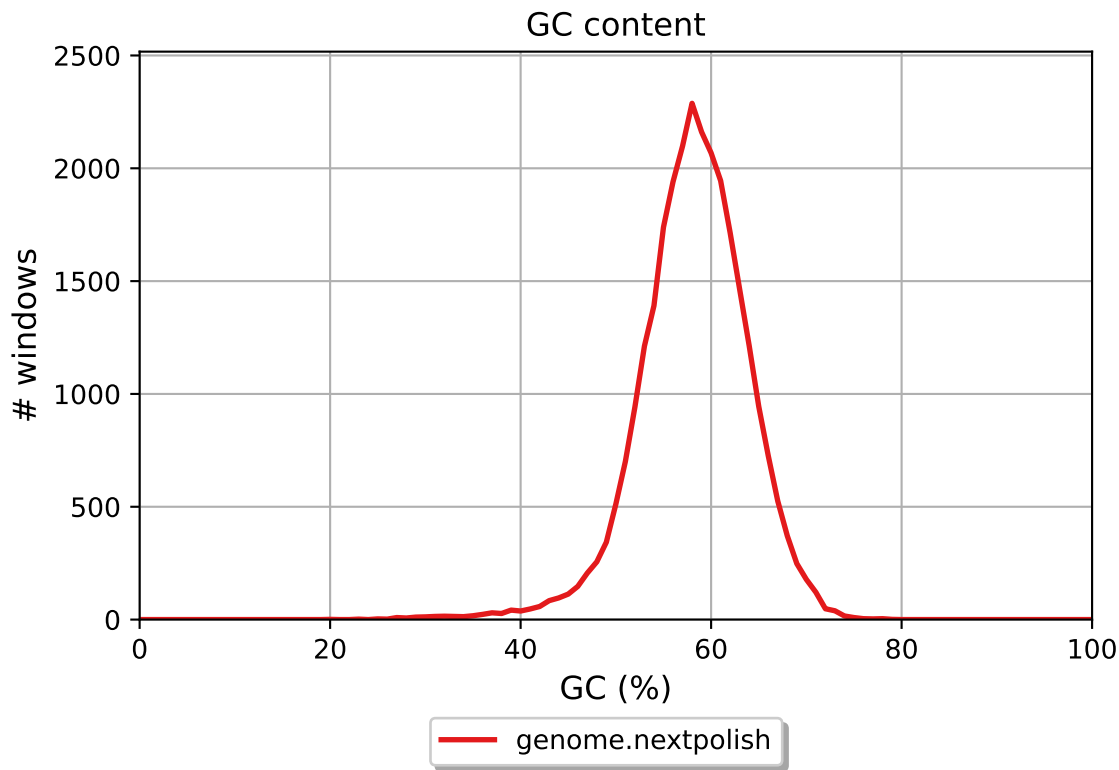
Report

	genome.nextpolish
# contigs (>= 0 bp)	71
# contigs (>= 1000 bp)	71
# contigs (>= 5000 bp)	59
# contigs (>= 10000 bp)	51
# contigs (>= 25000 bp)	38
# contigs (>= 50000 bp)	18
Total length (>= 0 bp)	2825265
Total length (>= 1000 bp)	2825265
Total length (>= 5000 bp)	2771565
Total length (>= 10000 bp)	2715988
Total length (>= 25000 bp)	2490882
Total length (>= 50000 bp)	1790378
# contigs	71
Largest contig	180433
Total length	2825265
GC (%)	58.12
N50	77670
N90	21097
auN	82831.4
L50	13
L90	41
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	12 + 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

