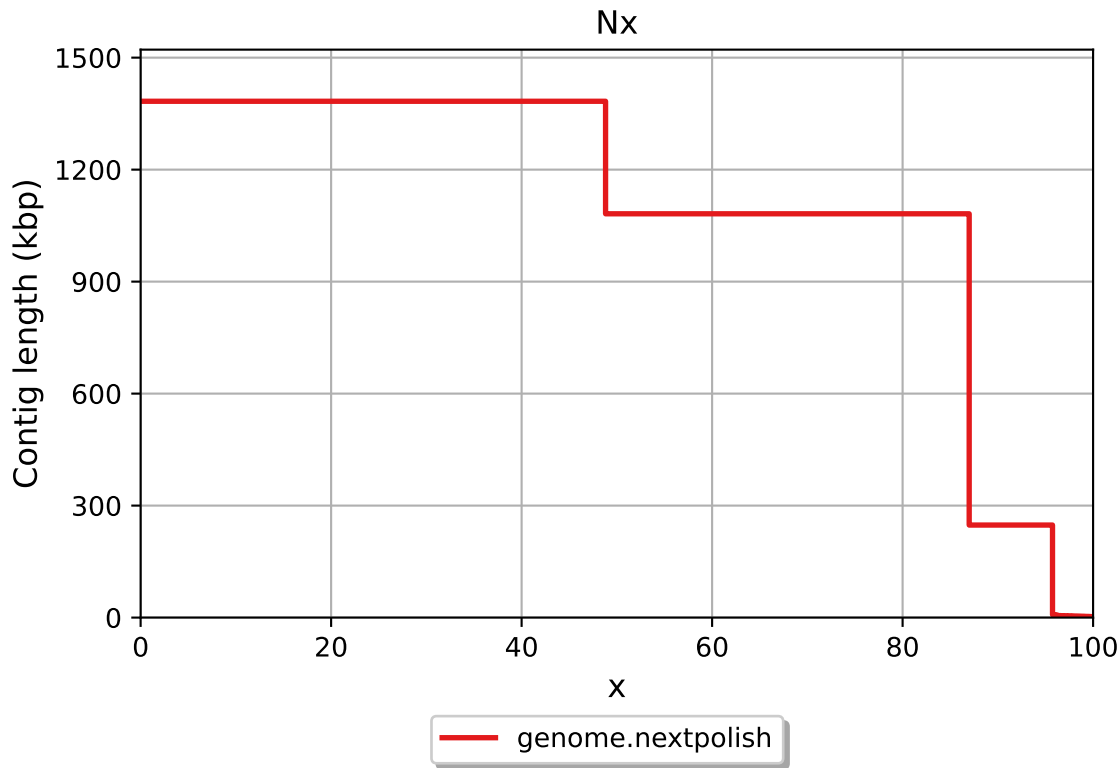
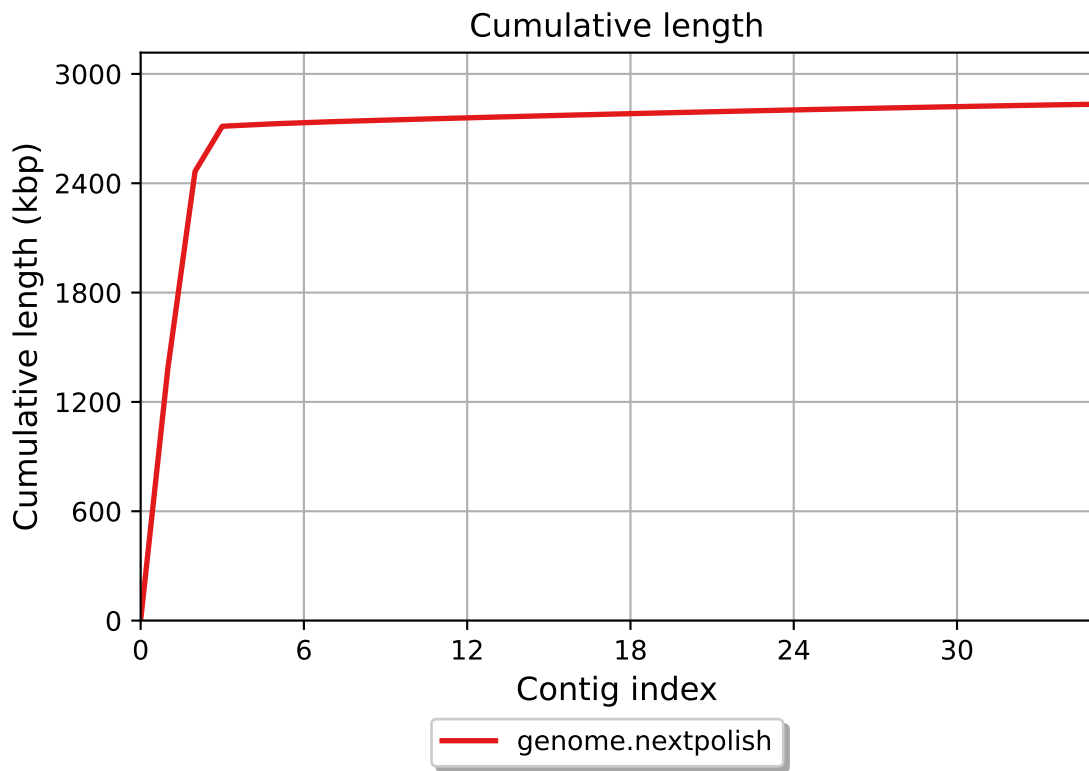


# Report

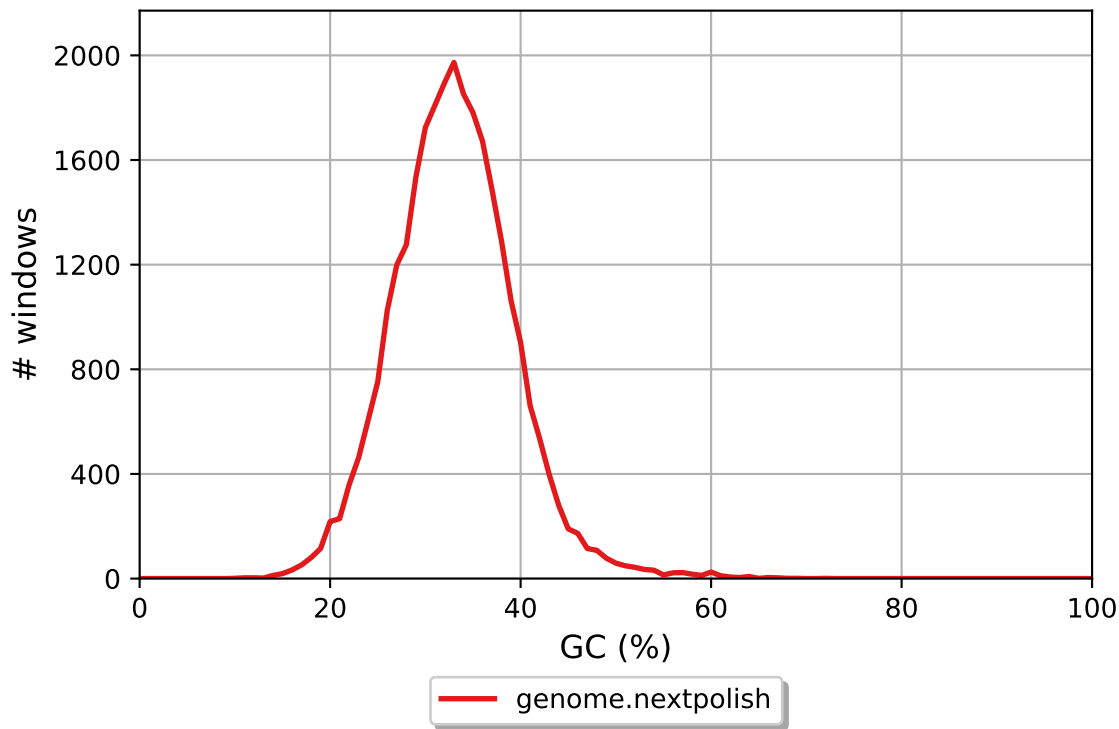
	genome.nextpolish
# contigs (>= 0 bp)	35
# contigs (>= 1000 bp)	35
# contigs (>= 5000 bp)	7
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	3
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	2833434
Total length (>= 1000 bp)	2833434
Total length (>= 5000 bp)	2737639
Total length (>= 10000 bp)	2712700
Total length (>= 25000 bp)	2712700
Total length (>= 50000 bp)	2712700
# contigs	35
Largest contig	1383127
Total length	2833434
GC (%)	33.06
N50	1081699
N90	247874
auN	1109978.4
L50	2
L90	3
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	10 + 8 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).





GC content



genome.nextpolish GC content

