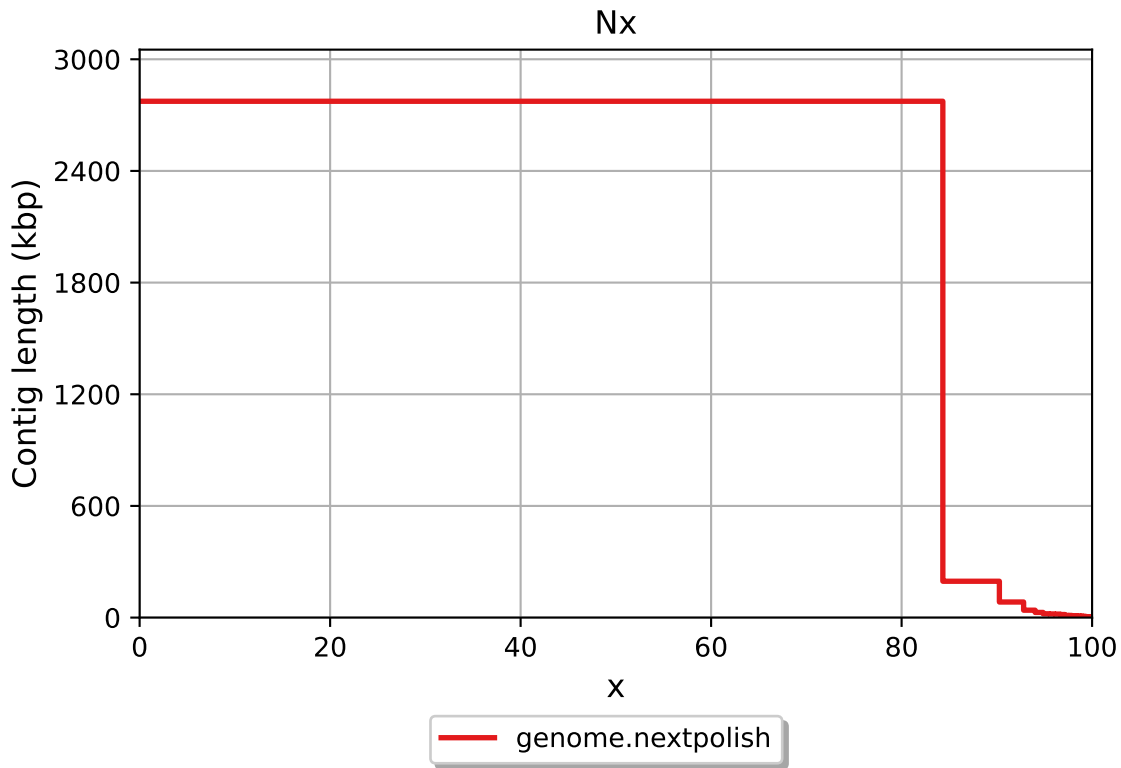
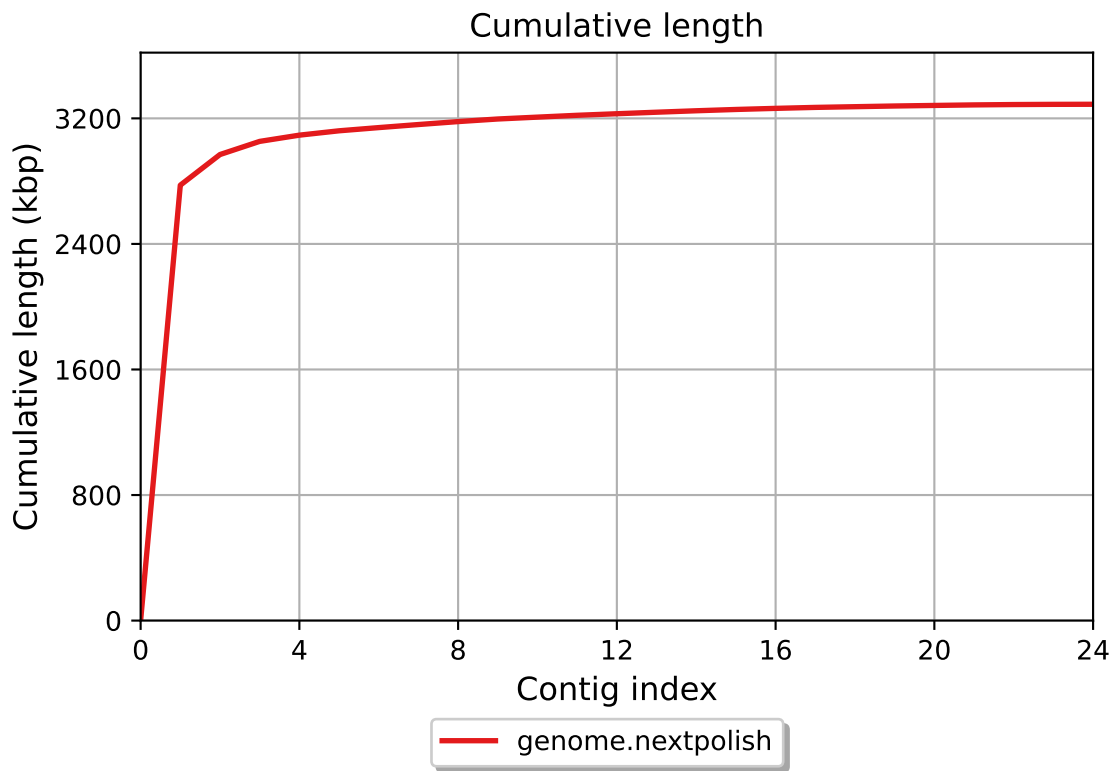


Report

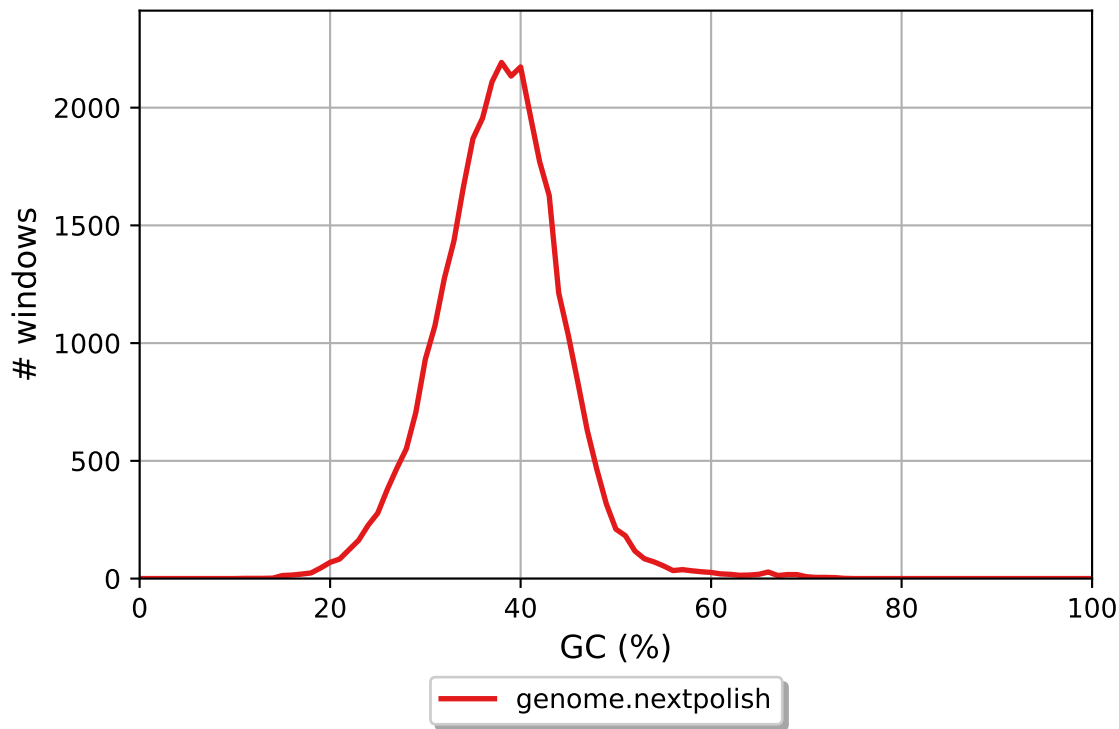
	genome.nextpolish
# contigs (>= 0 bp)	24
# contigs (>= 1000 bp)	23
# contigs (>= 5000 bp)	17
# contigs (>= 10000 bp)	11
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	3289849
Total length (>= 1000 bp)	3289337
Total length (>= 5000 bp)	3269902
Total length (>= 10000 bp)	3219579
Total length (>= 25000 bp)	3120779
Total length (>= 50000 bp)	3053319
# contigs	24
Largest contig	2774332
Total length	3289849
GC (%)	37.84
N50	2774332
N90	195275
auN	2354701.4
L50	1
L90	2
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	18 + 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).





GC content



genome.nextpolish GC content

