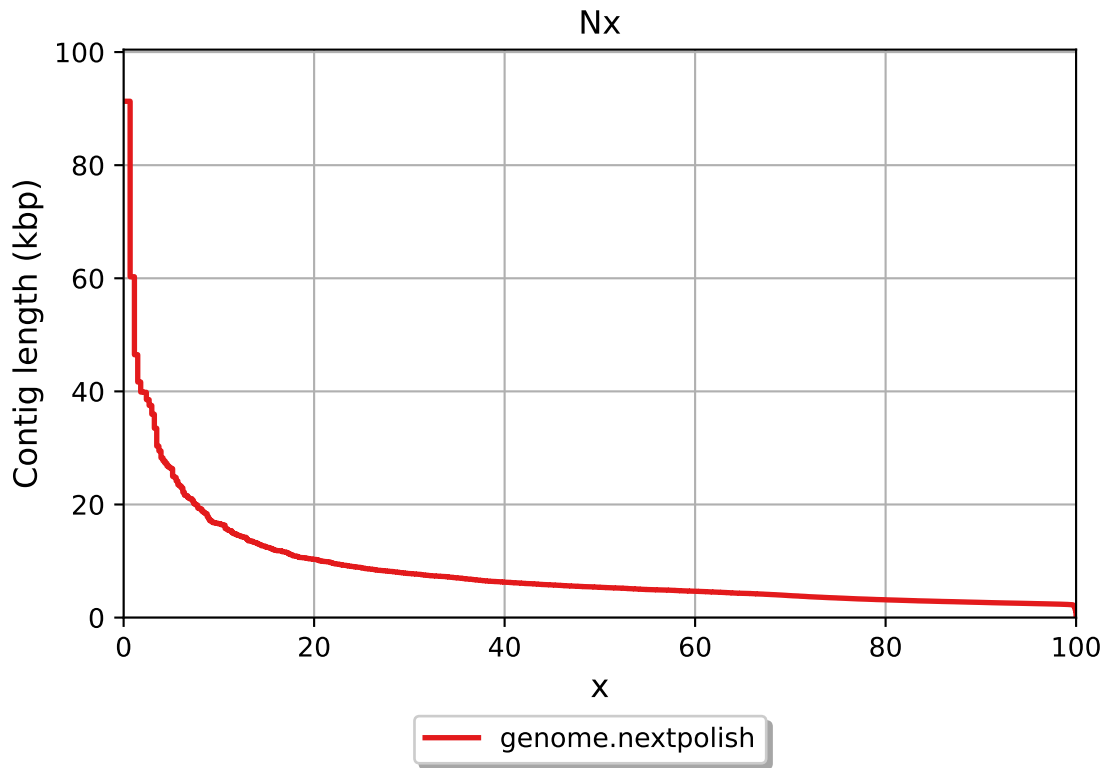


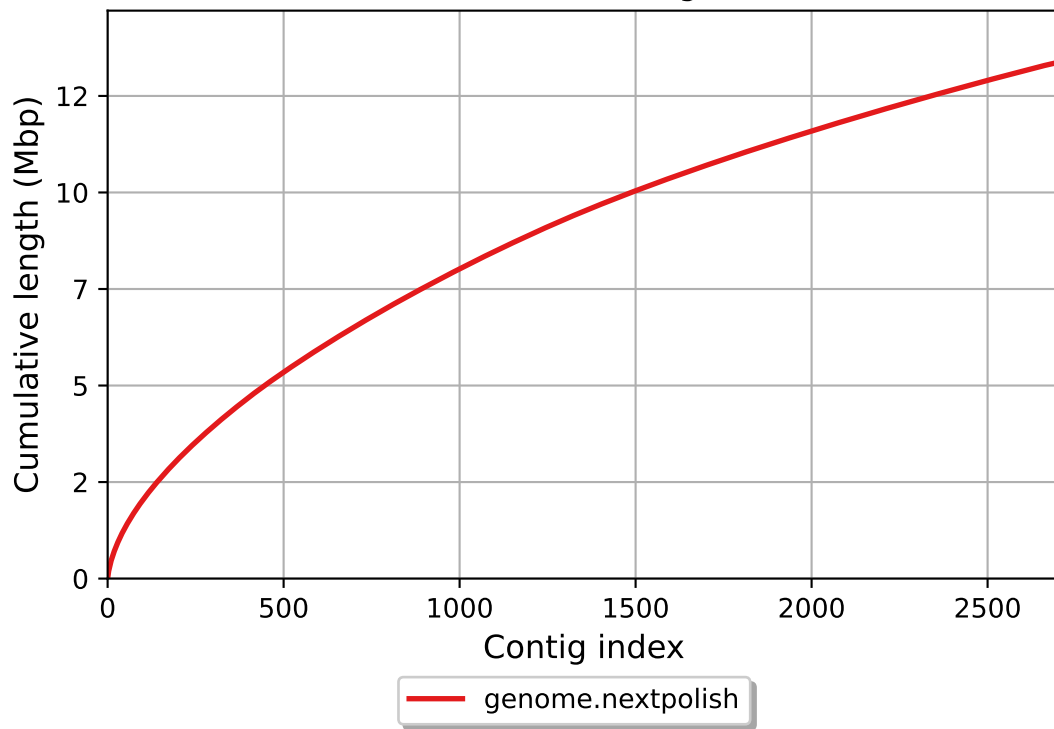
# Report

	genome.nextpolish
# contigs (>= 0 bp)	2706
# contigs (>= 1000 bp)	2704
# contigs (>= 5000 bp)	851
# contigs (>= 10000 bp)	168
# contigs (>= 25000 bp)	18
# contigs (>= 50000 bp)	2
Total length (>= 0 bp)	13372068
Total length (>= 1000 bp)	13371012
Total length (>= 5000 bp)	7302274
Total length (>= 10000 bp)	2783786
Total length (>= 25000 bp)	687663
Total length (>= 50000 bp)	151572
# contigs	2706
Largest contig	91300
Total length	13372068
GC (%)	48.32
N50	5359
N90	2688
auN	8528.0
L50	733
L90	2161
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	17 + 3 part

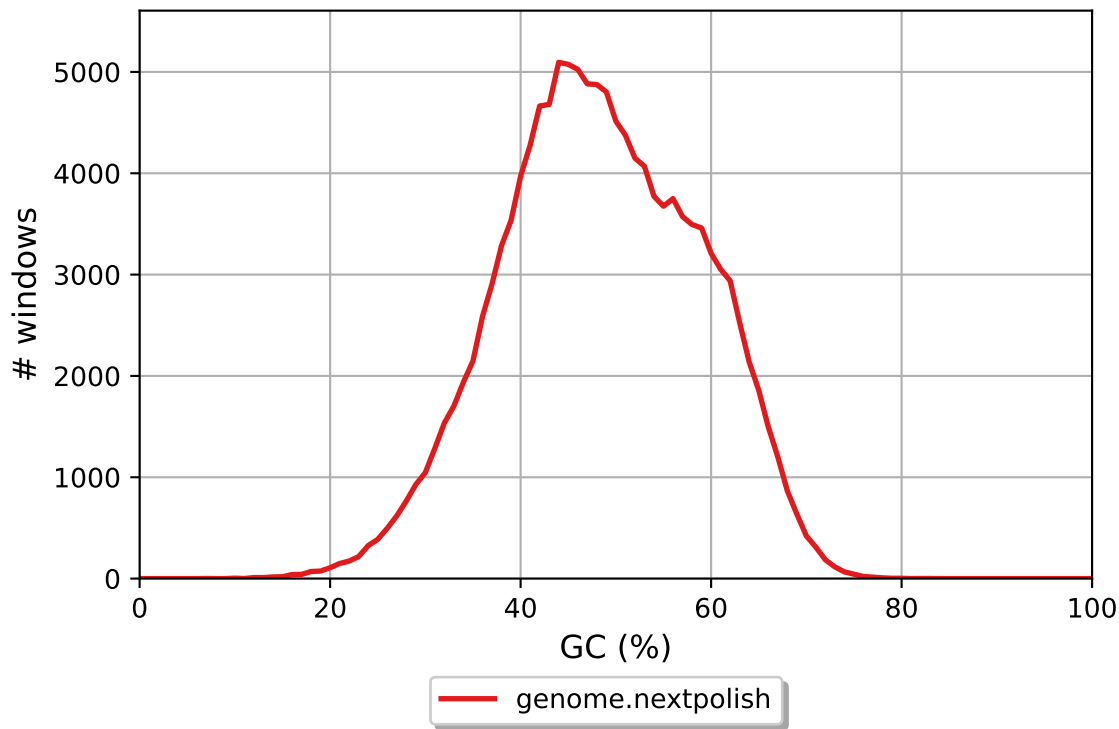
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

