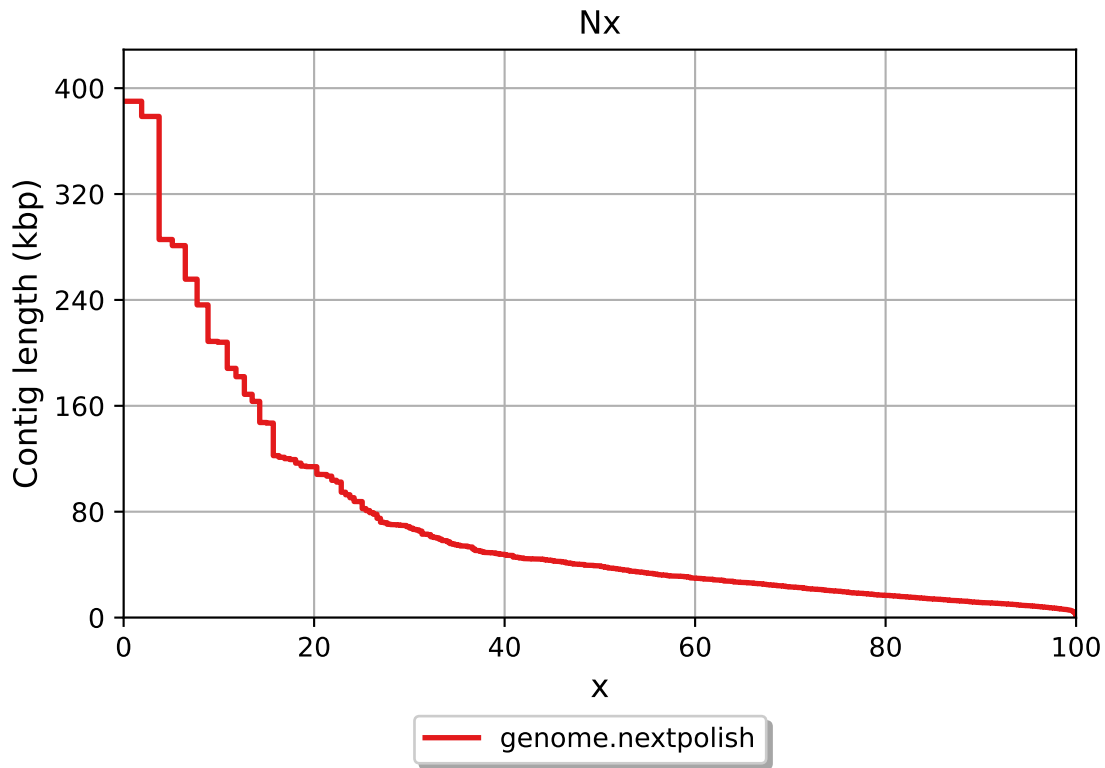


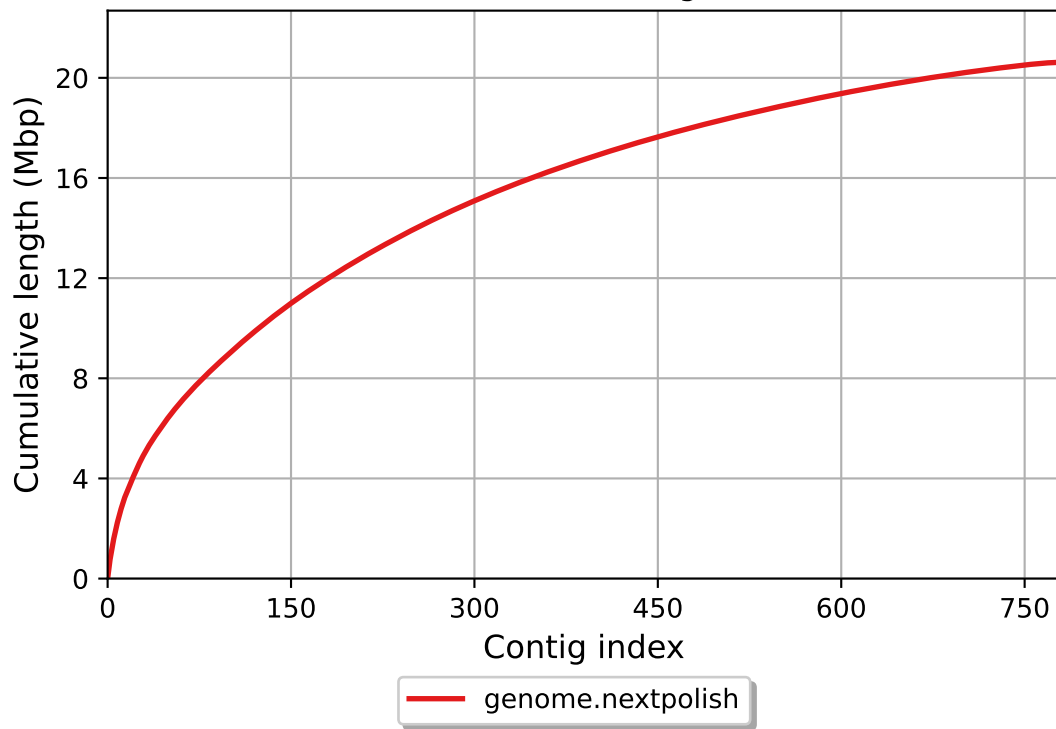
Report

	genome.nextpolish
# contigs (>= 0 bp)	779
# contigs (>= 1000 bp)	778
# contigs (>= 5000 bp)	756
# contigs (>= 10000 bp)	586
# contigs (>= 25000 bp)	249
# contigs (>= 50000 bp)	73
Total length (>= 0 bp)	20622121
Total length (>= 1000 bp)	20621572
Total length (>= 5000 bp)	20543892
Total length (>= 10000 bp)	19237012
Total length (>= 25000 bp)	13916829
Total length (>= 50000 bp)	7756181
# contigs	779
Largest contig	390091
Total length	20622121
GC (%)	49.30
N50	39058
N90	11400
auN	74263.1
L50	132
L90	524
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	63 + 3 part

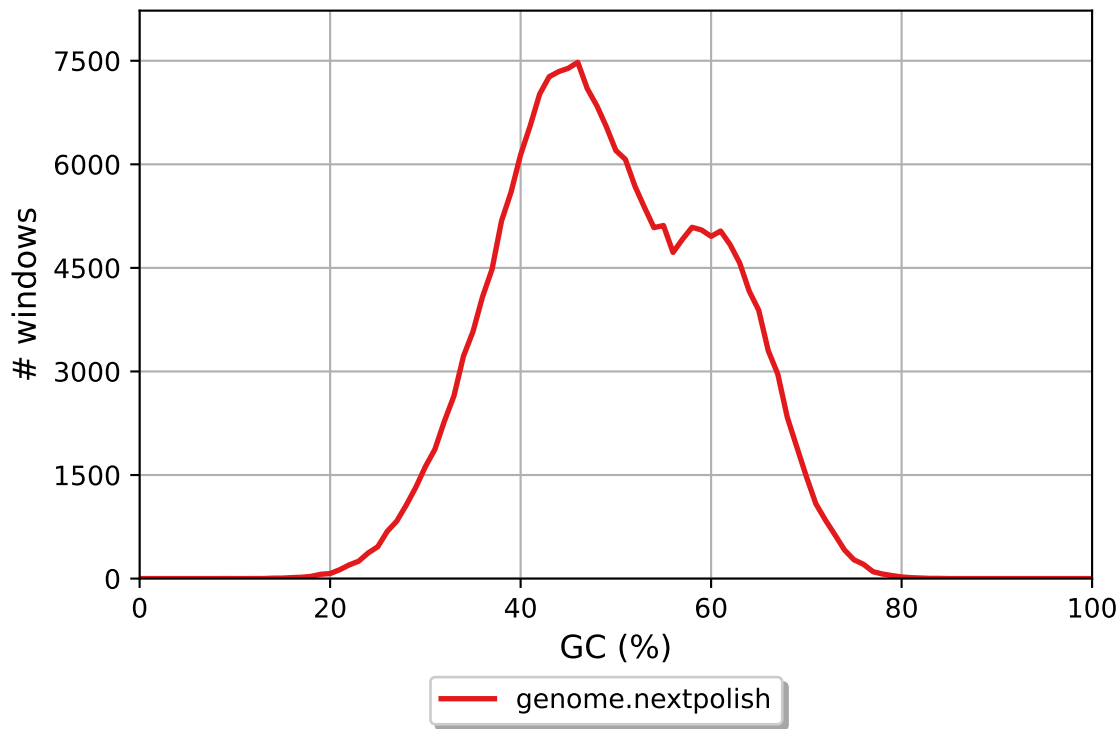
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

