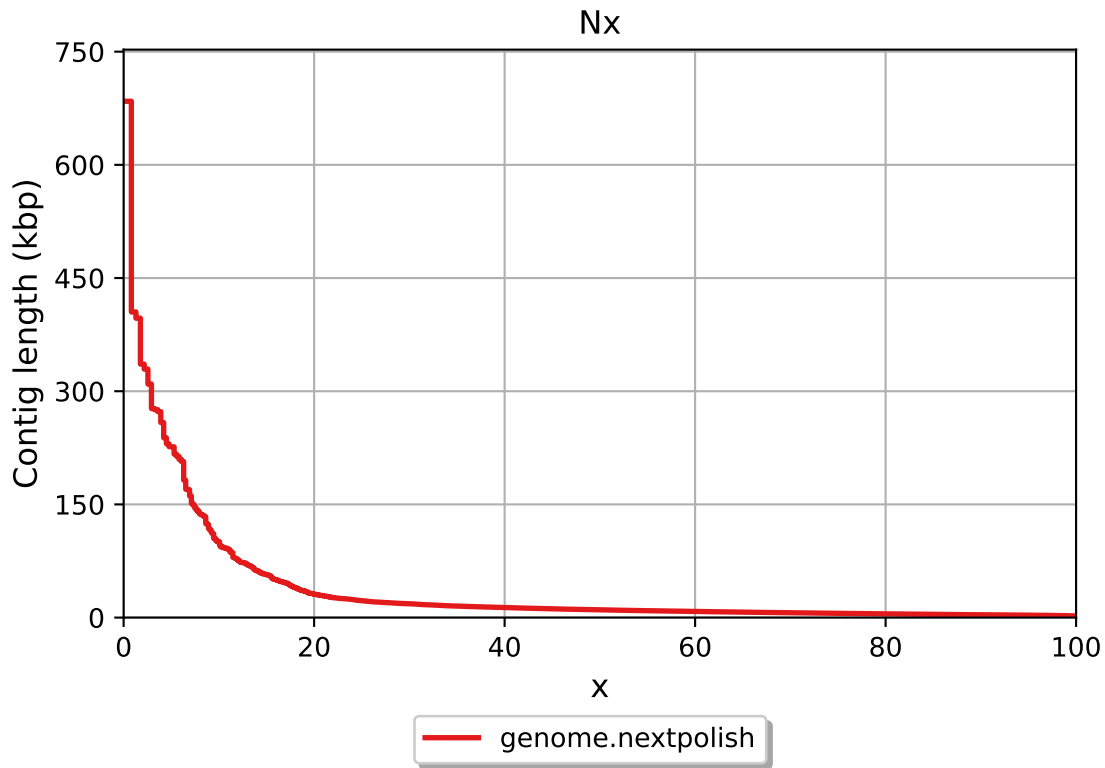


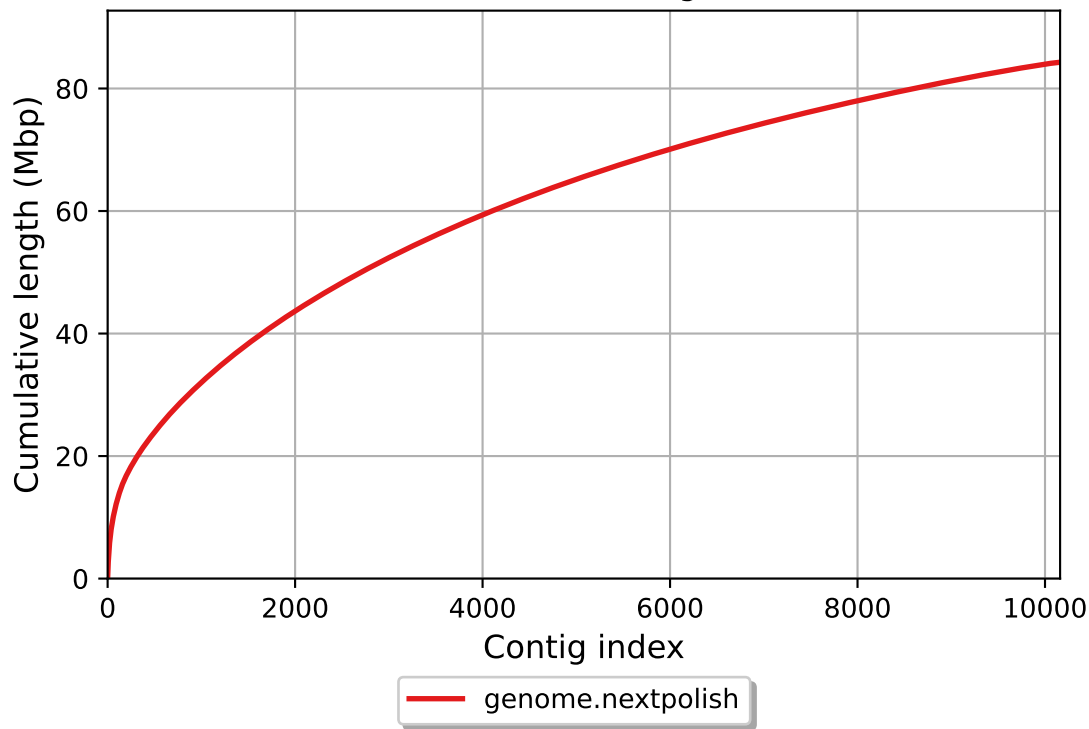
# Report

	genome.nextpolish
# contigs (>= 0 bp)	10161
# contigs (>= 1000 bp)	10144
# contigs (>= 5000 bp)	5381
# contigs (>= 10000 bp)	1946
# contigs (>= 25000 bp)	296
# contigs (>= 50000 bp)	116
Total length (>= 0 bp)	84281452
Total length (>= 1000 bp)	84269670
Total length (>= 5000 bp)	67142464
Total length (>= 10000 bp)	43135988
Total length (>= 25000 bp)	19484645
Total length (>= 50000 bp)	13537758
# contigs	10161
Largest contig	684256
Total length	84281452
GC (%)	38.50
N50	10316
N90	3715
auN	40198.9
L50	1849
L90	7401
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	71 + 14 part

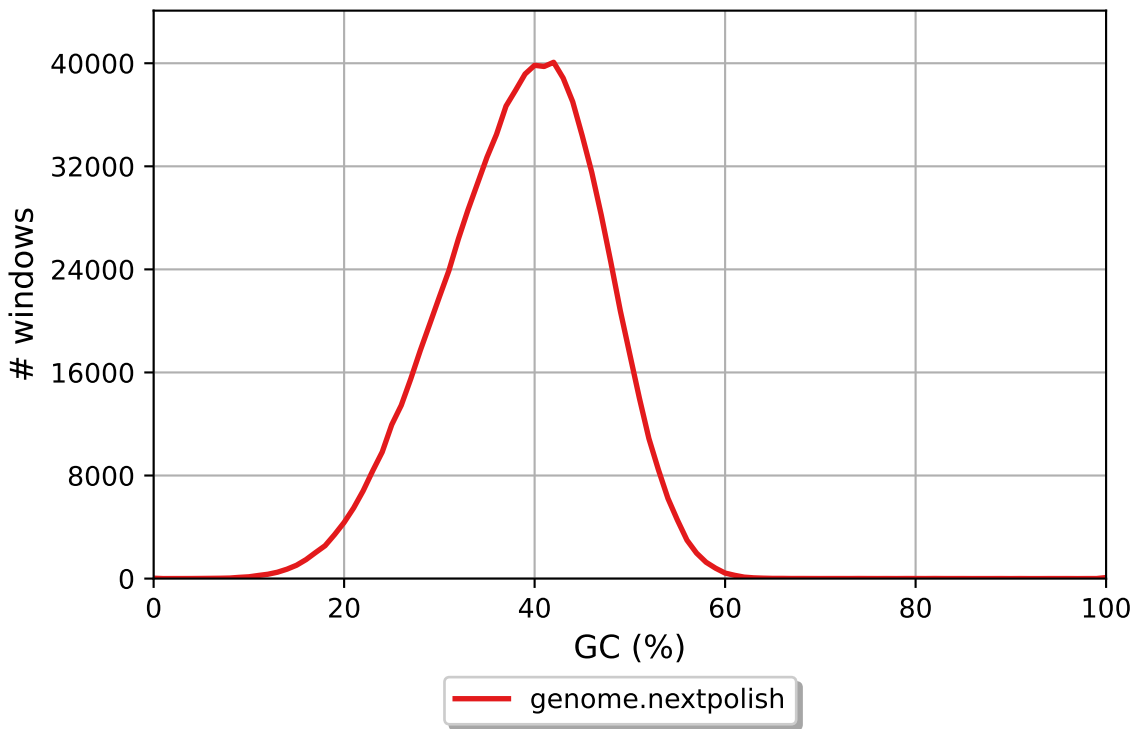
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



# GC content



genome.nextpolish GC content

