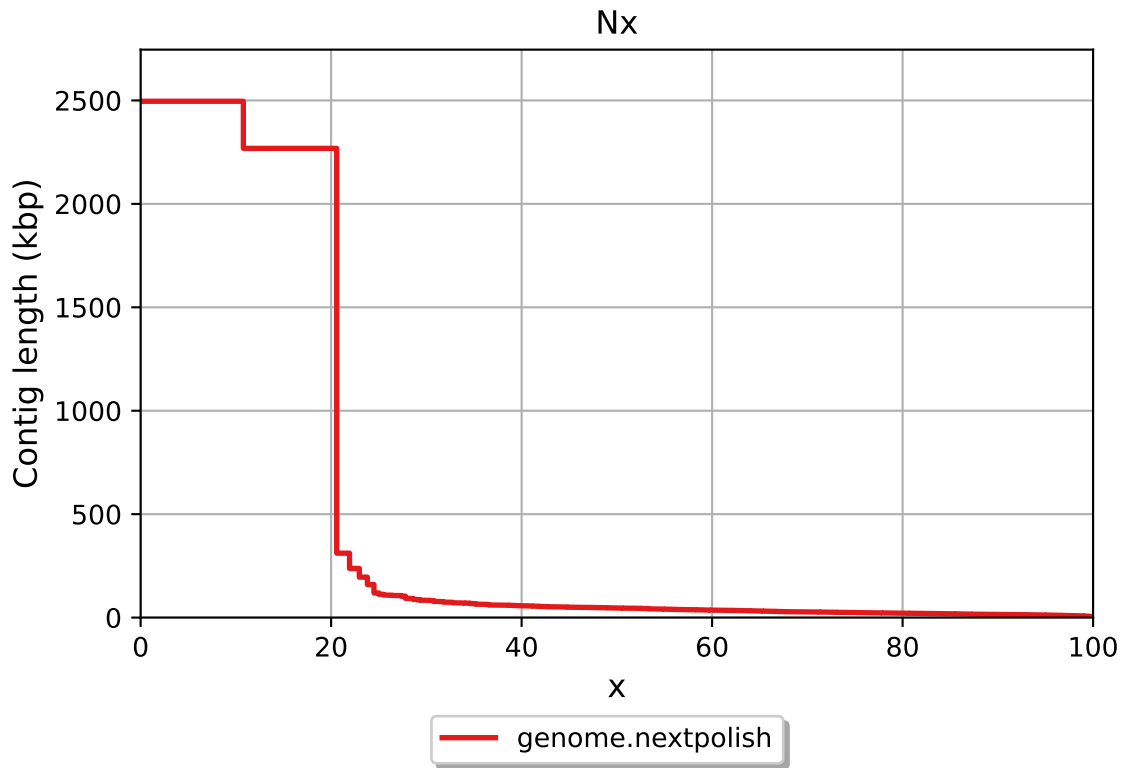


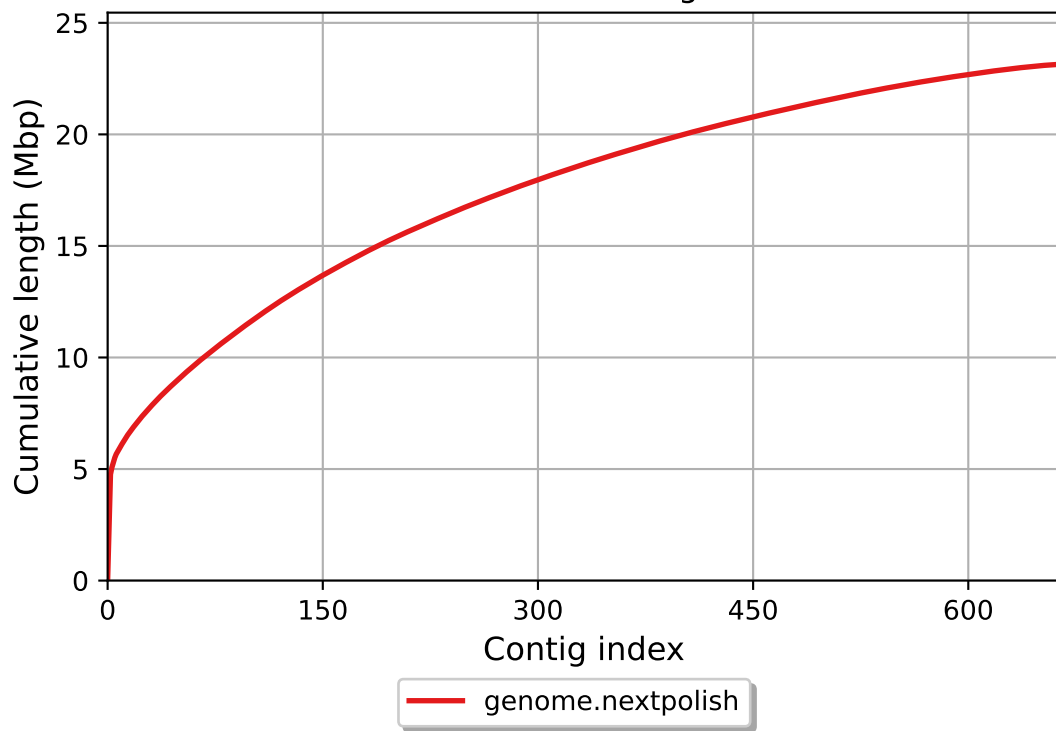
# Report

	genome.nextpolish
# contigs (>= 0 bp)	664
# contigs (>= 1000 bp)	663
# contigs (>= 5000 bp)	658
# contigs (>= 10000 bp)	583
# contigs (>= 25000 bp)	259
# contigs (>= 50000 bp)	77
Total length (>= 0 bp)	23142921
Total length (>= 1000 bp)	23142035
Total length (>= 5000 bp)	23119617
Total length (>= 10000 bp)	22519564
Total length (>= 25000 bp)	16988075
Total length (>= 50000 bp)	10515413
# contigs	664
Largest contig	2496014
Total length	23142921
GC (%)	42.81
N50	46549
N90	15178
auN	531064.3
L50	99
L90	453
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	73 + 4 part

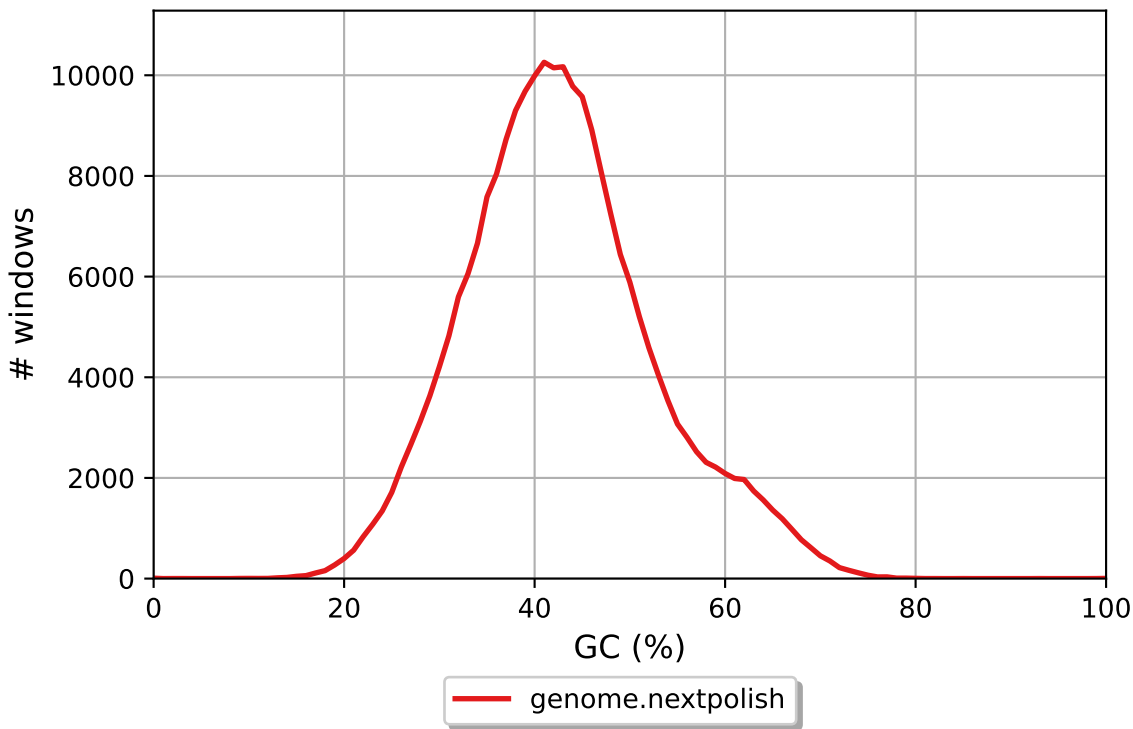
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

