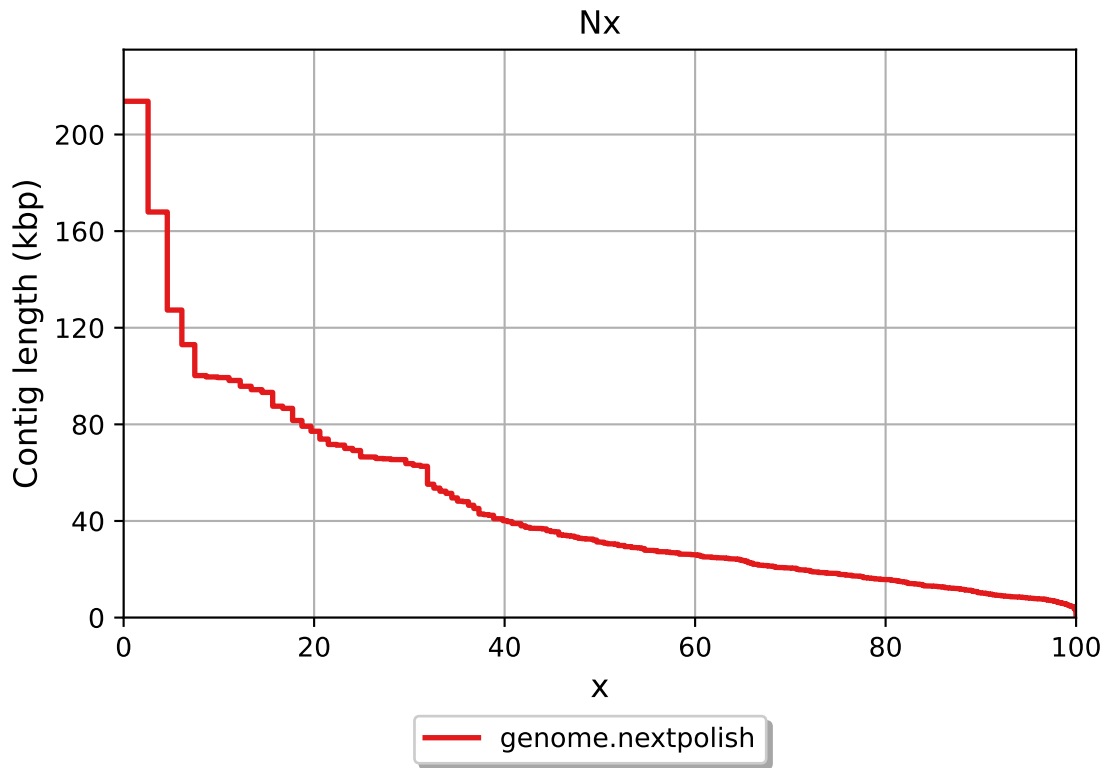


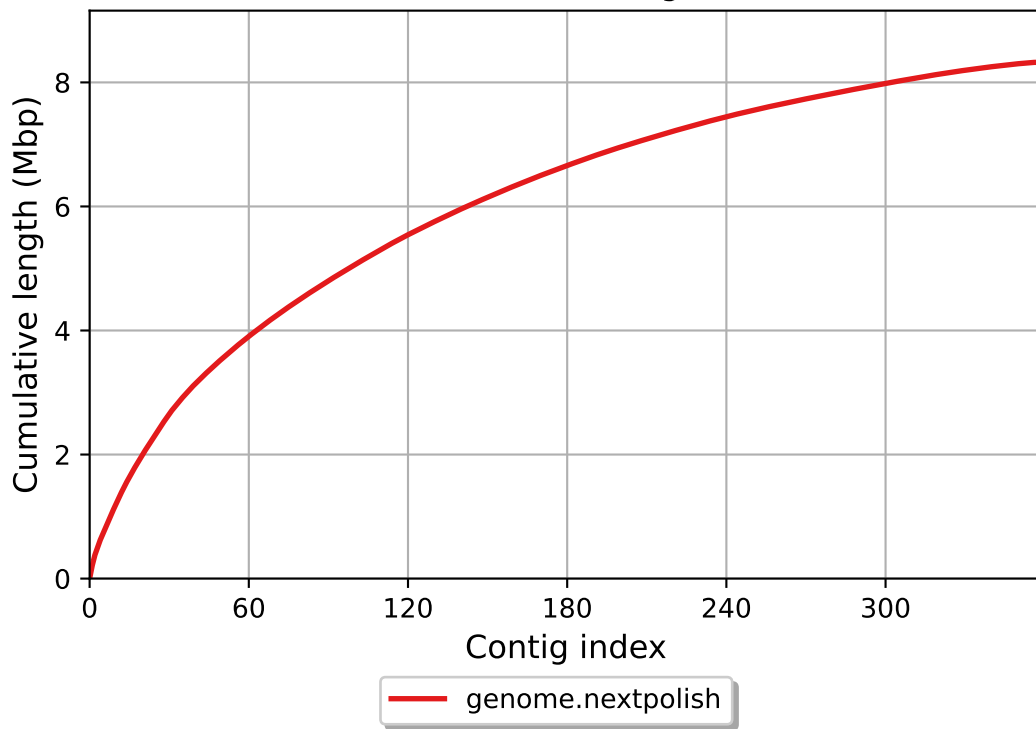
# Report

	genome.nextpolish
# contigs (>= 0 bp)	359
# contigs (>= 1000 bp)	356
# contigs (>= 5000 bp)	342
# contigs (>= 10000 bp)	249
# contigs (>= 25000 bp)	103
# contigs (>= 50000 bp)	34
Total length (>= 0 bp)	8324673
Total length (>= 1000 bp)	8322886
Total length (>= 5000 bp)	8263506
Total length (>= 10000 bp)	7539561
Total length (>= 25000 bp)	5137825
Total length (>= 50000 bp)	2868327
# contigs	359
Largest contig	213762
Total length	8324673
GC (%)	46.05
N50	31321
N90	10240
auN	48462.7
L50	68
L90	245
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	20 + 3 part

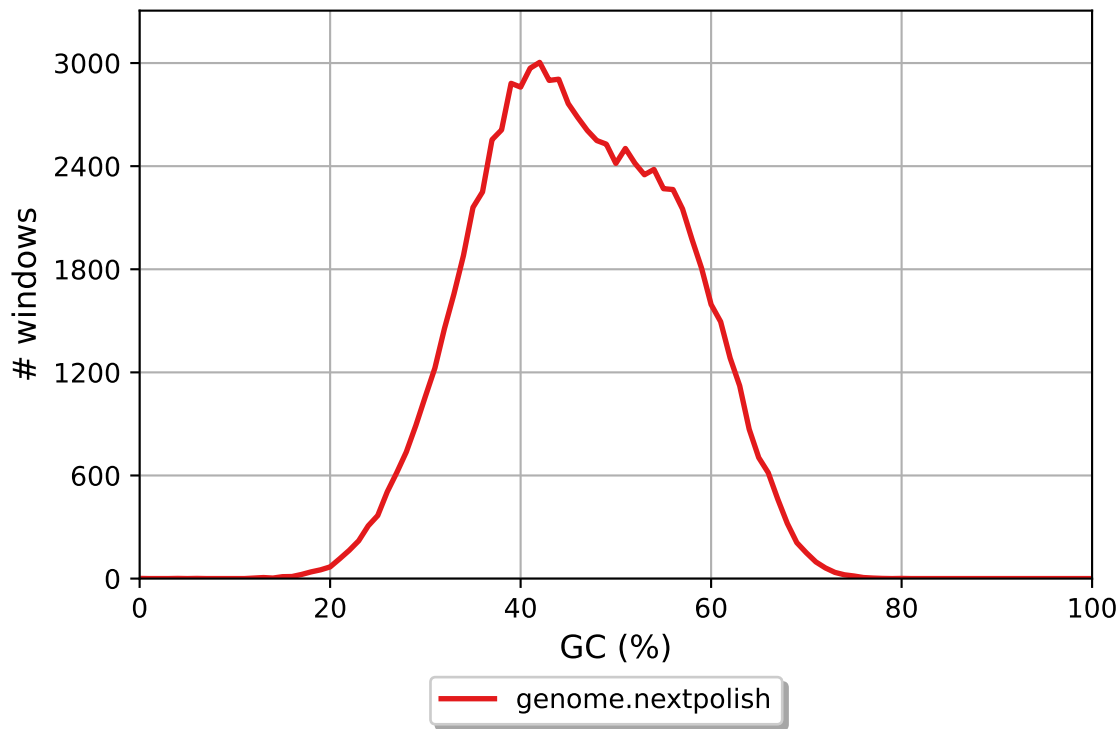
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

