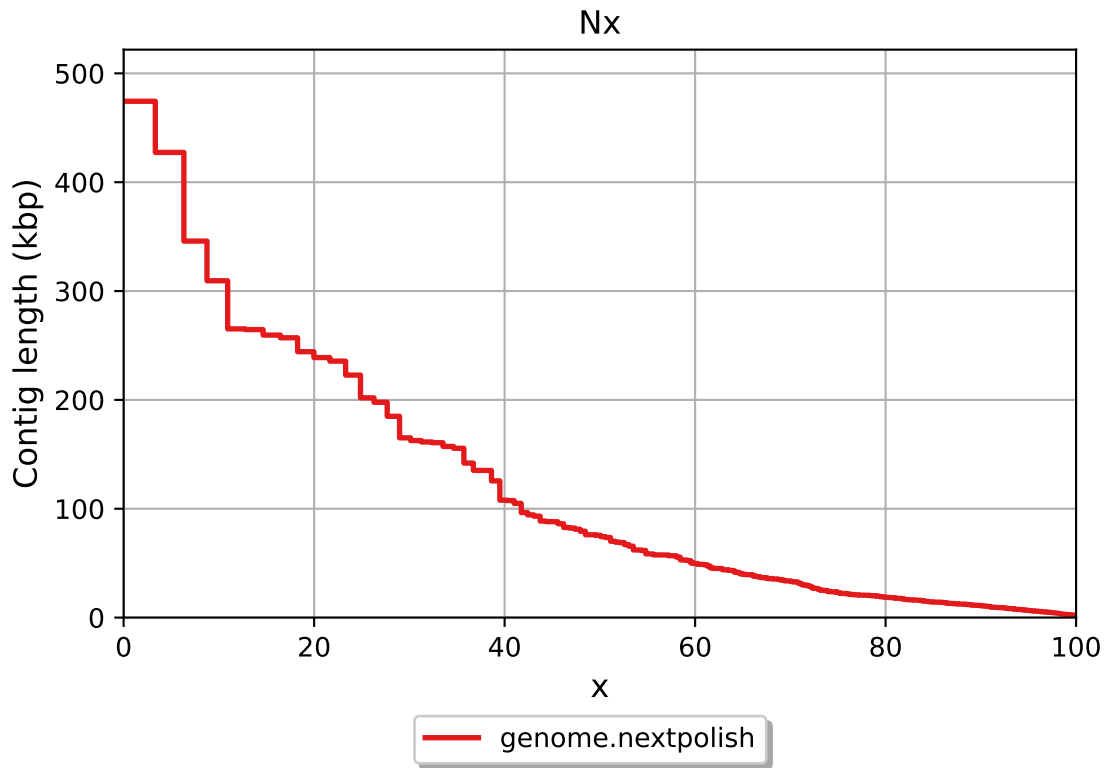


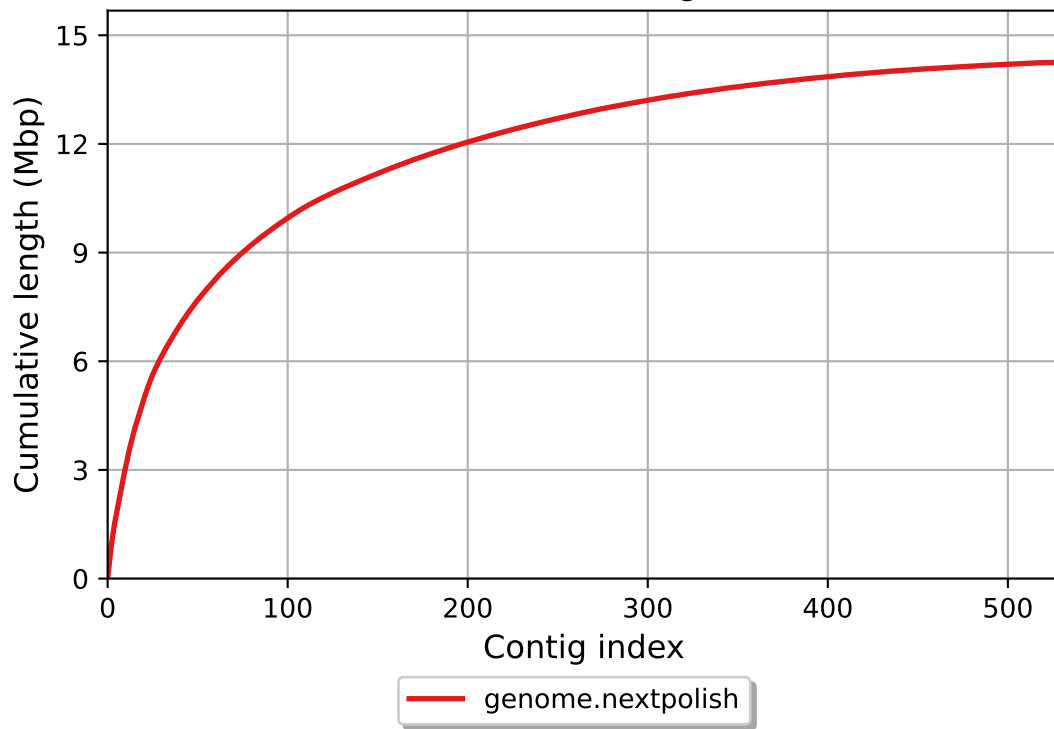
# Report

	genome.nextpolish
# contigs (>= 0 bp)	529
# contigs (>= 1000 bp)	526
# contigs (>= 5000 bp)	393
# contigs (>= 10000 bp)	275
# contigs (>= 25000 bp)	117
# contigs (>= 50000 bp)	65
Total length (>= 0 bp)	14254391
Total length (>= 1000 bp)	14252763
Total length (>= 5000 bp)	13822520
Total length (>= 10000 bp)	12976627
Total length (>= 25000 bp)	10456457
Total length (>= 50000 bp)	8538075
# contigs	529
Largest contig	474352
Total length	14254391
GC (%)	41.80
N50	75644
N90	11100
auN	126659.5
L50	42
L90	261
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	35 + 2 part

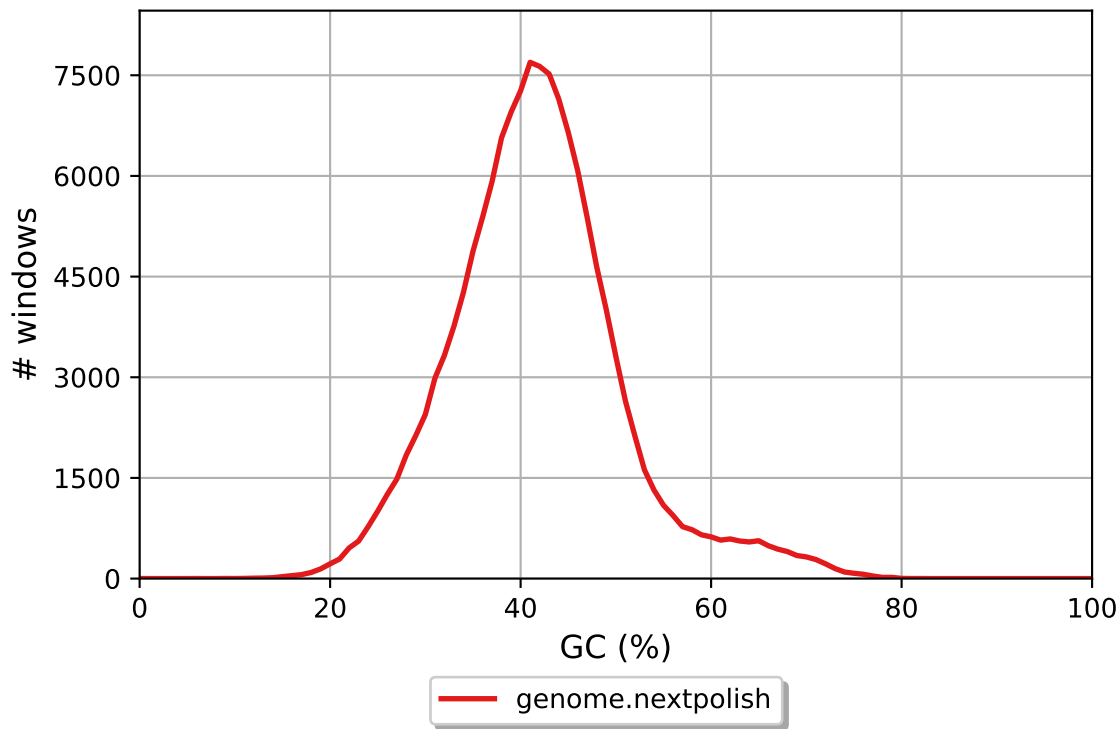
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

