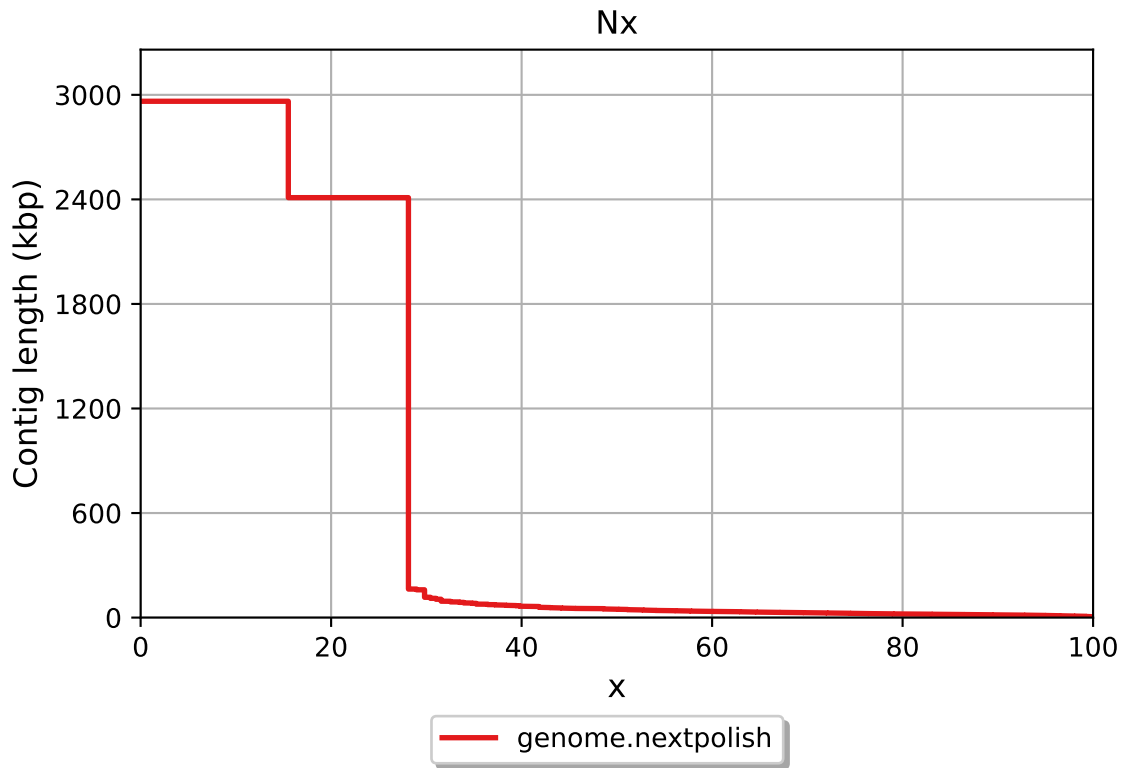


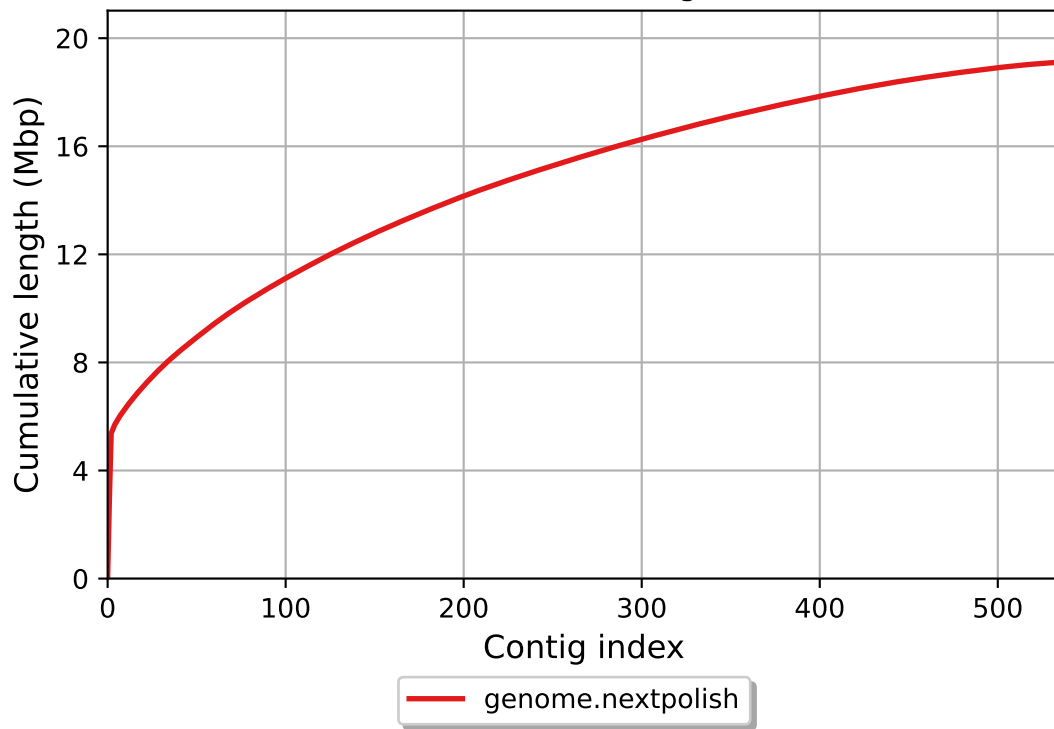
# Report

	genome.nextpolish
# contigs (>= 0 bp)	535
# contigs (>= 1000 bp)	535
# contigs (>= 5000 bp)	526
# contigs (>= 10000 bp)	458
# contigs (>= 25000 bp)	200
# contigs (>= 50000 bp)	57
Total length (>= 0 bp)	19109037
Total length (>= 1000 bp)	19109037
Total length (>= 5000 bp)	19074961
Total length (>= 10000 bp)	18537286
Total length (>= 25000 bp)	14158789
Total length (>= 50000 bp)	9289306
# contigs	535
Largest contig	2963128
Total length	19109037
GC (%)	42.41
N50	47844
N90	15439
auN	792966.4
L50	63
L90	356
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	69 + 5 part

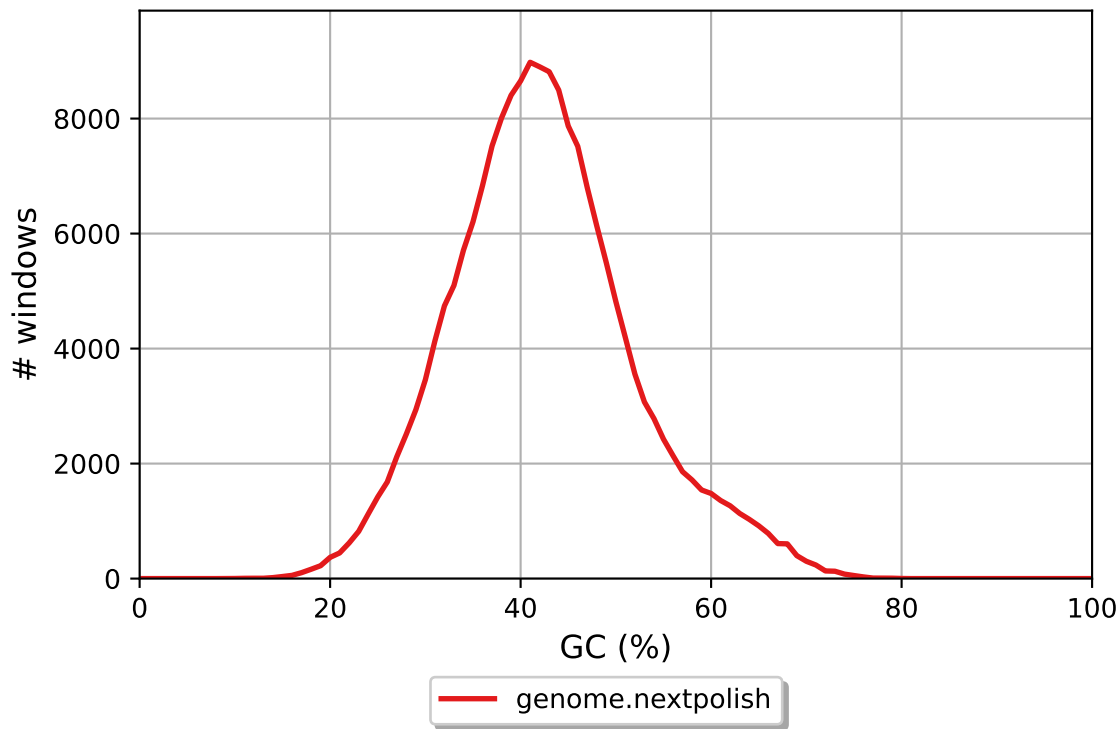
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

