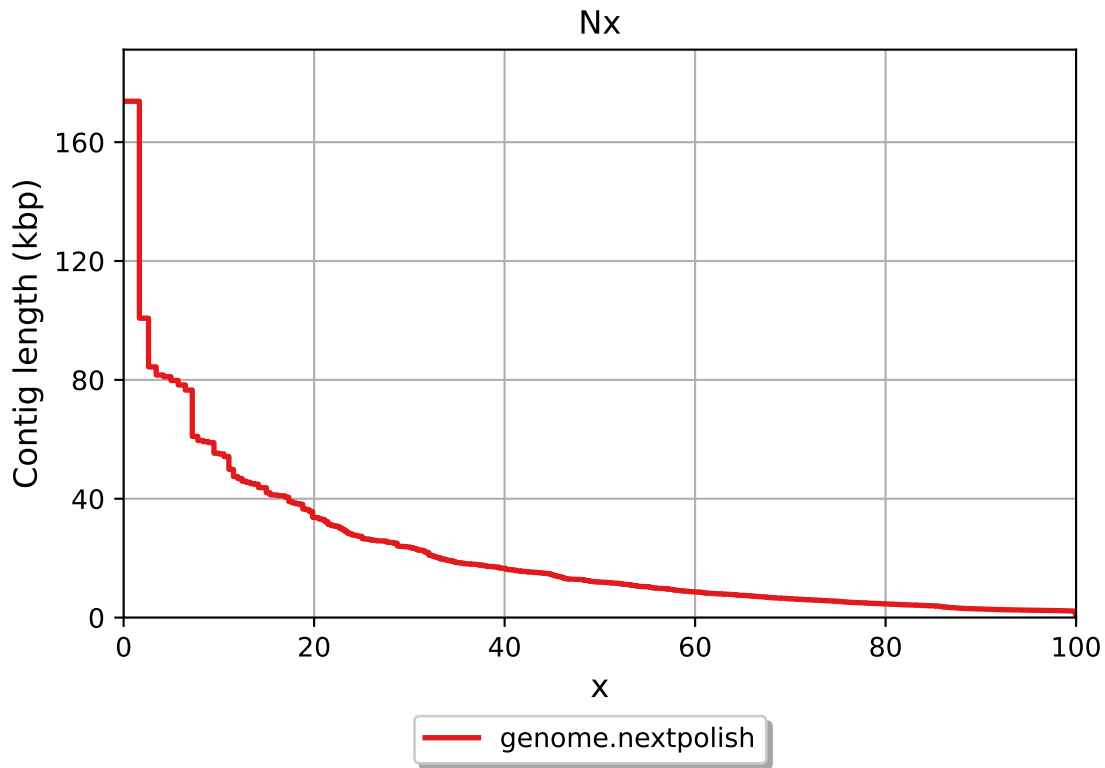


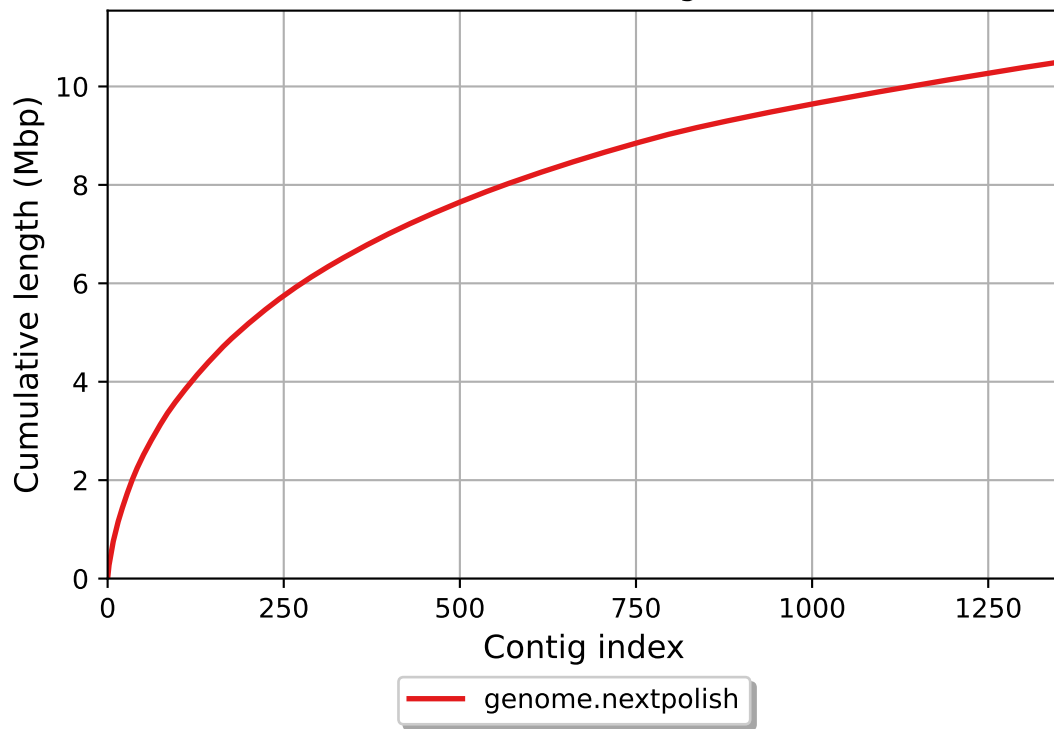
# Report

	genome.nextpolish
# contigs (>= 0 bp)	1352
# contigs (>= 1000 bp)	1348
# contigs (>= 5000 bp)	583
# contigs (>= 10000 bp)	259
# contigs (>= 25000 bp)	69
# contigs (>= 50000 bp)	15
Total length (>= 0 bp)	10491561
Total length (>= 1000 bp)	10488520
Total length (>= 5000 bp)	8099006
Total length (>= 10000 bp)	5840635
Total length (>= 25000 bp)	2989972
Total length (>= 50000 bp)	1159483
# contigs	1352
Largest contig	173761
Total length	10491561
GC (%)	66.35
N50	11961
N90	2890
auN	22611.6
L50	206
L90	928
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 0 part

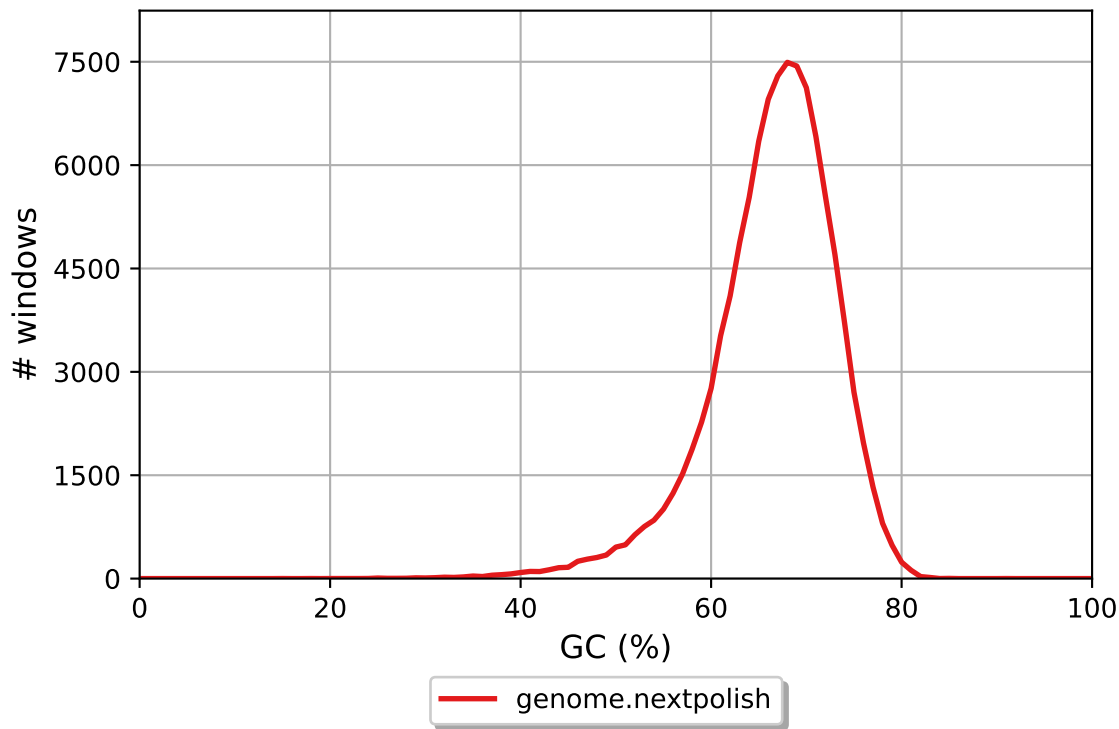
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

