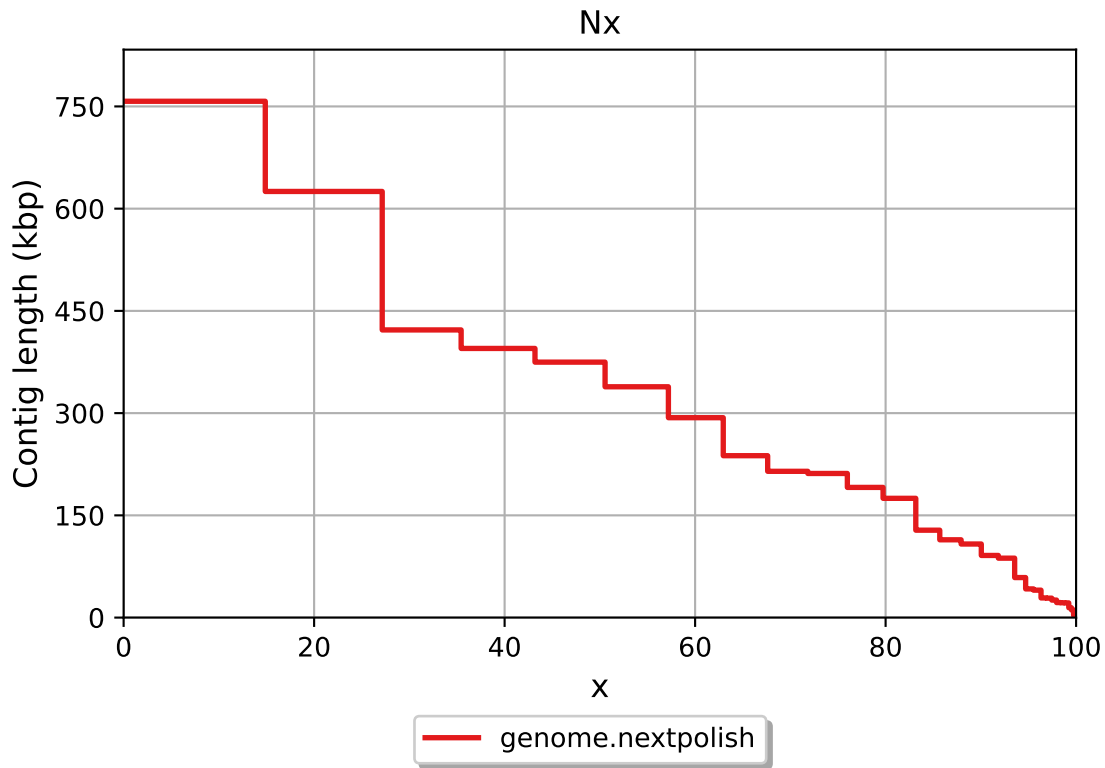


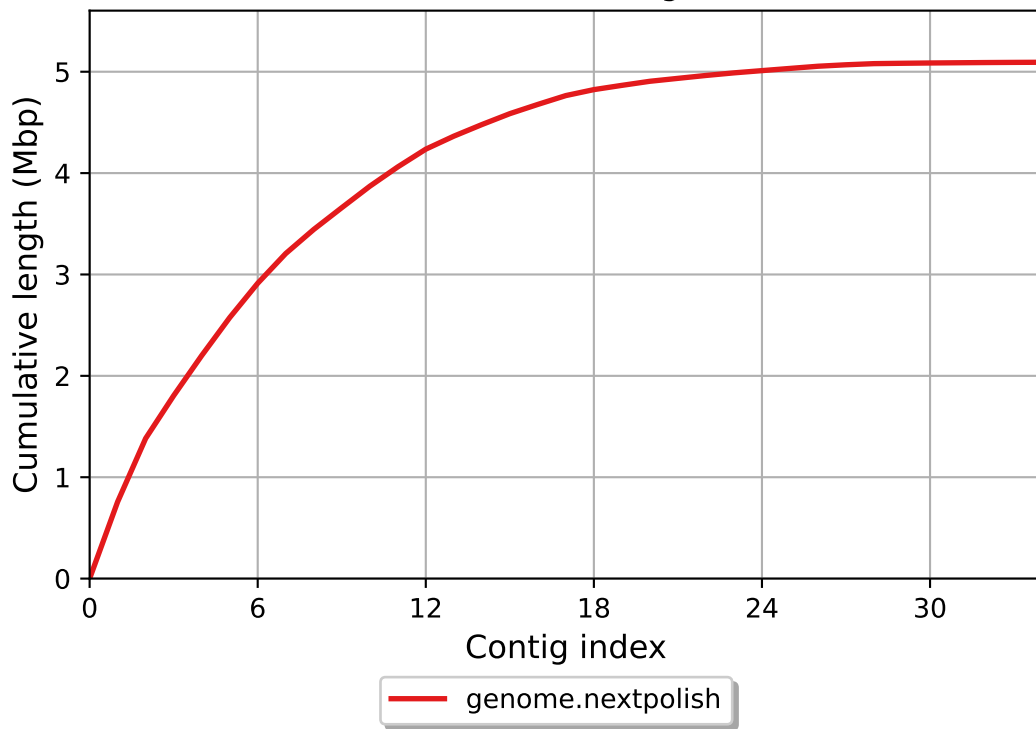
# Report

	genome.nextpolish
# contigs (>= 0 bp)	34
# contigs (>= 1000 bp)	34
# contigs (>= 5000 bp)	28
# contigs (>= 10000 bp)	28
# contigs (>= 25000 bp)	23
# contigs (>= 50000 bp)	18
Total length (>= 0 bp)	5093390
Total length (>= 1000 bp)	5093390
Total length (>= 5000 bp)	5079908
Total length (>= 10000 bp)	5079908
Total length (>= 25000 bp)	4988893
Total length (>= 50000 bp)	4823556
# contigs	34
Largest contig	757567
Total length	5093390
GC (%)	50.90
N50	374804
N90	107959
auN	377429.7
L50	5
L90	15
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 7 part

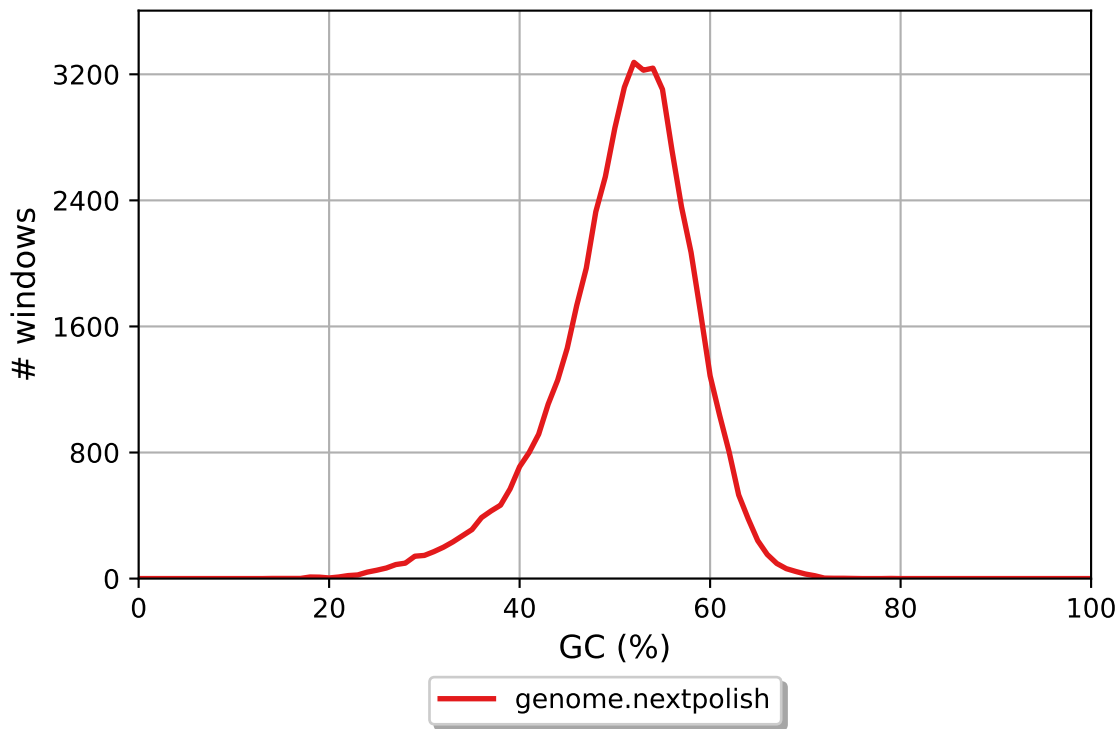
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

