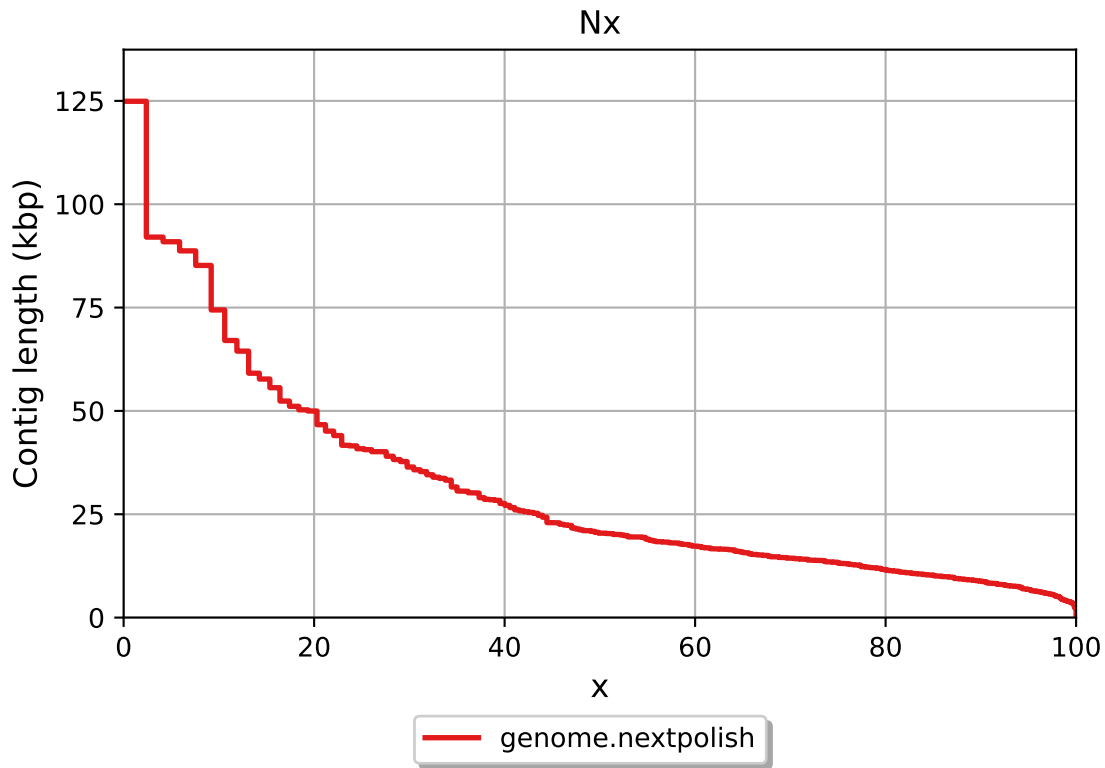


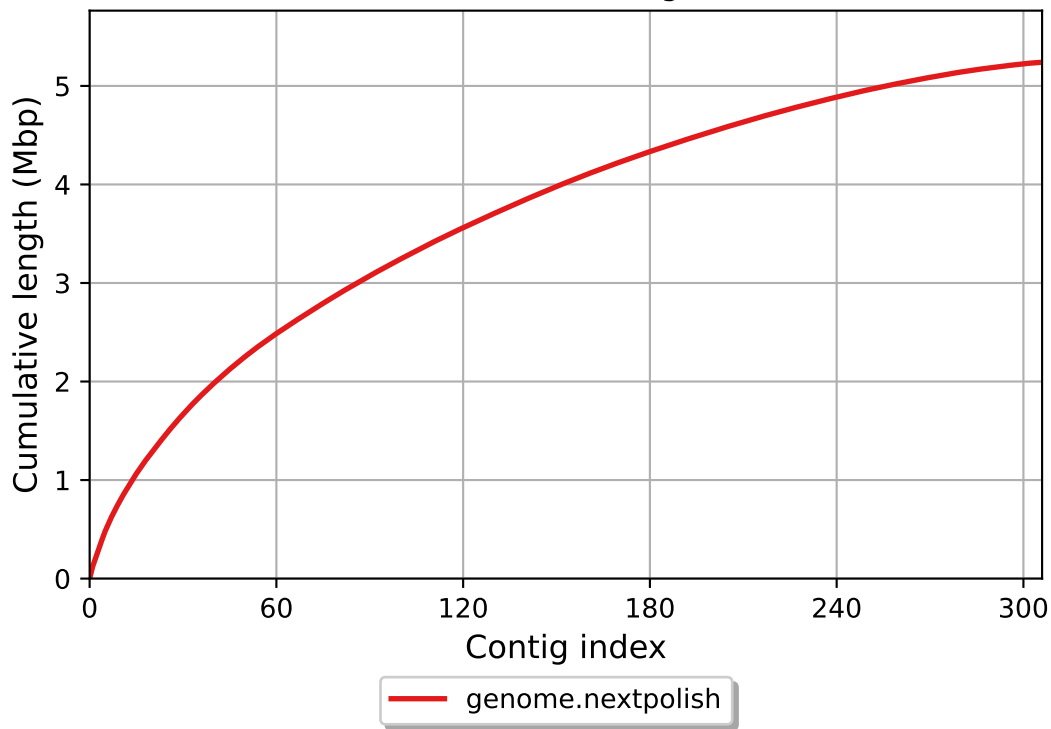
# Report

	genome.nextpolish
# contigs (>= 0 bp)	306
# contigs (>= 1000 bp)	306
# contigs (>= 5000 bp)	281
# contigs (>= 10000 bp)	196
# contigs (>= 25000 bp)	51
# contigs (>= 50000 bp)	14
Total length (>= 0 bp)	5240091
Total length (>= 1000 bp)	5240091
Total length (>= 5000 bp)	5146917
Total length (>= 10000 bp)	4499438
Total length (>= 25000 bp)	2279667
Total length (>= 50000 bp)	1013881
# contigs	306
Largest contig	124905
Total length	5240091
GC (%)	52.50
N50	20422
N90	8818
auN	31652.6
L50	67
L90	220
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 0 part

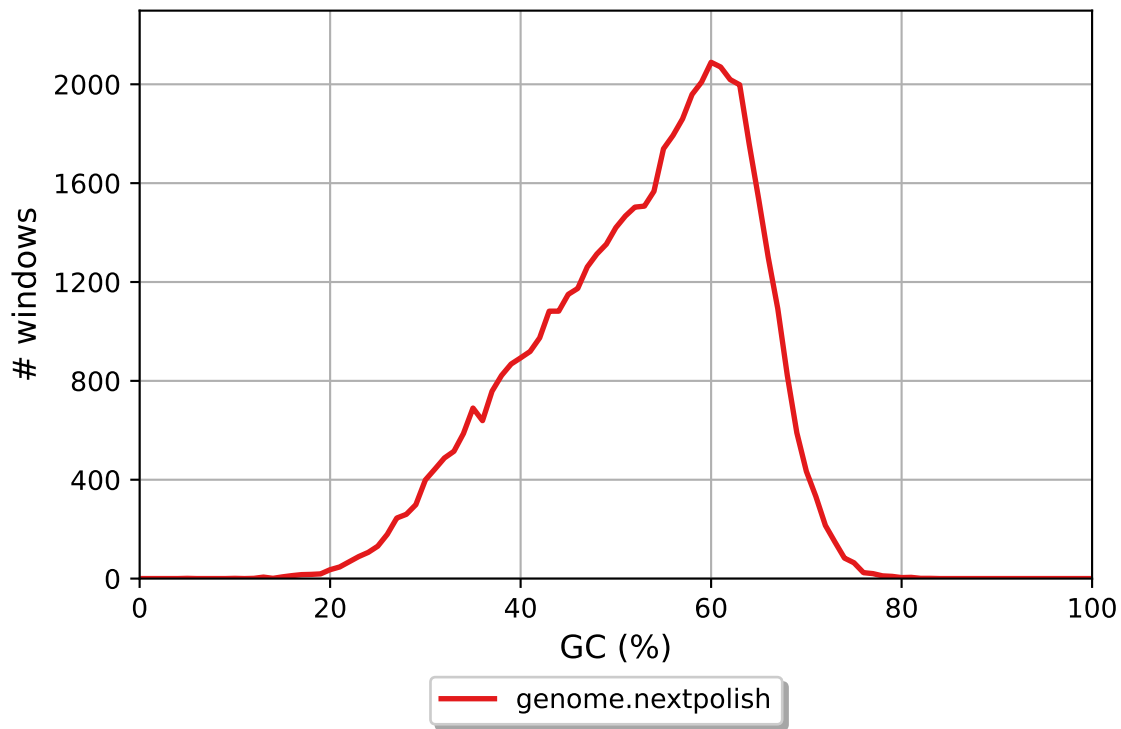
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

