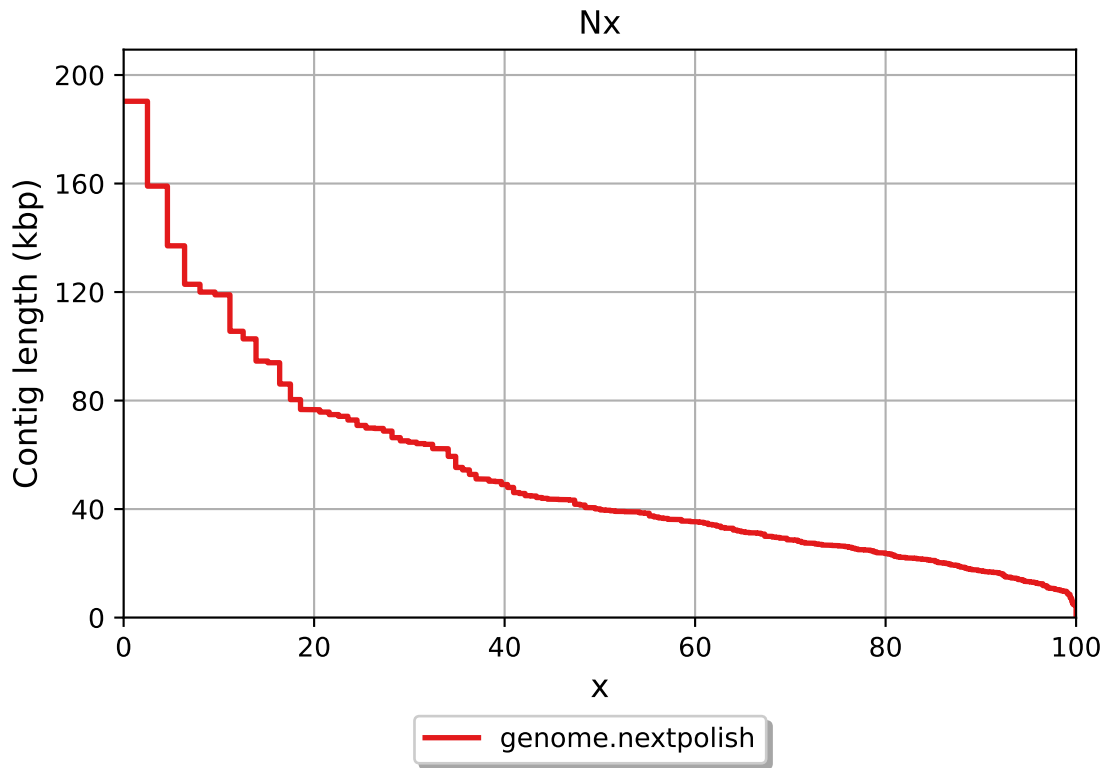


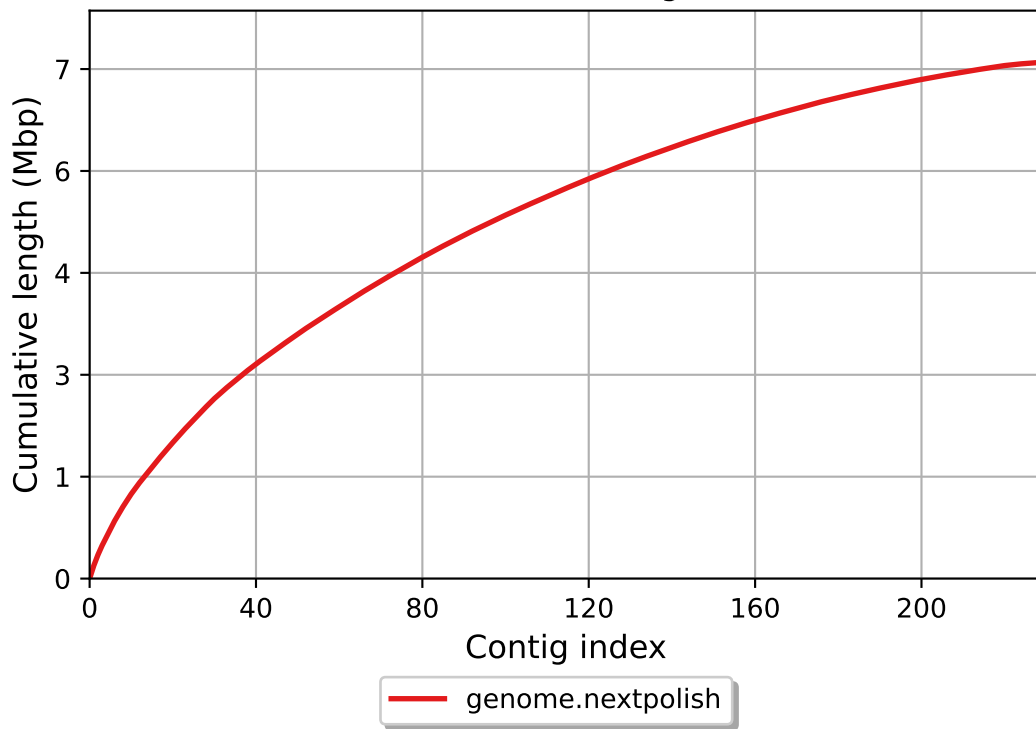
# Report

	genome.nextpolish
# contigs (>= 0 bp)	229
# contigs (>= 1000 bp)	229
# contigs (>= 5000 bp)	224
# contigs (>= 10000 bp)	212
# contigs (>= 25000 bp)	121
# contigs (>= 50000 bp)	37
Total length (>= 0 bp)	7599960
Total length (>= 1000 bp)	7599960
Total length (>= 5000 bp)	7576950
Total length (>= 10000 bp)	7476242
Total length (>= 25000 bp)	5909779
Total length (>= 50000 bp)	3014671
# contigs	229
Largest contig	190322
Total length	7599960
GC (%)	57.42
N50	40071
N90	17226
auN	54600.2
L50	55
L90	166
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	27 + 1 part

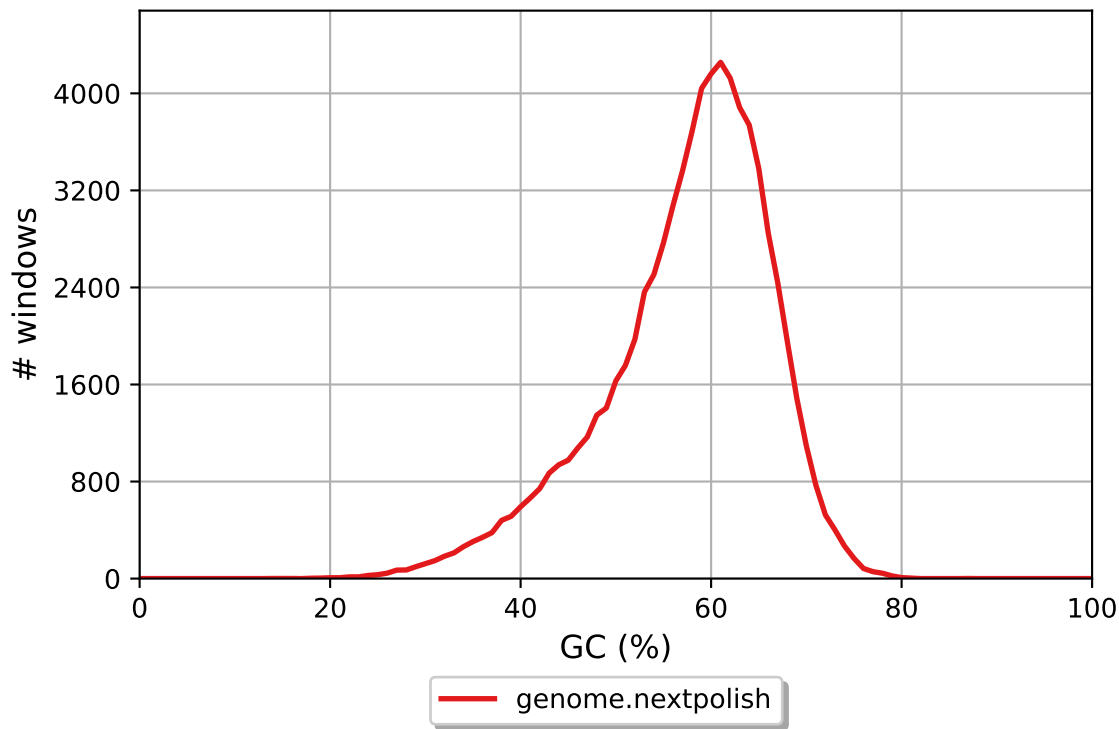
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

