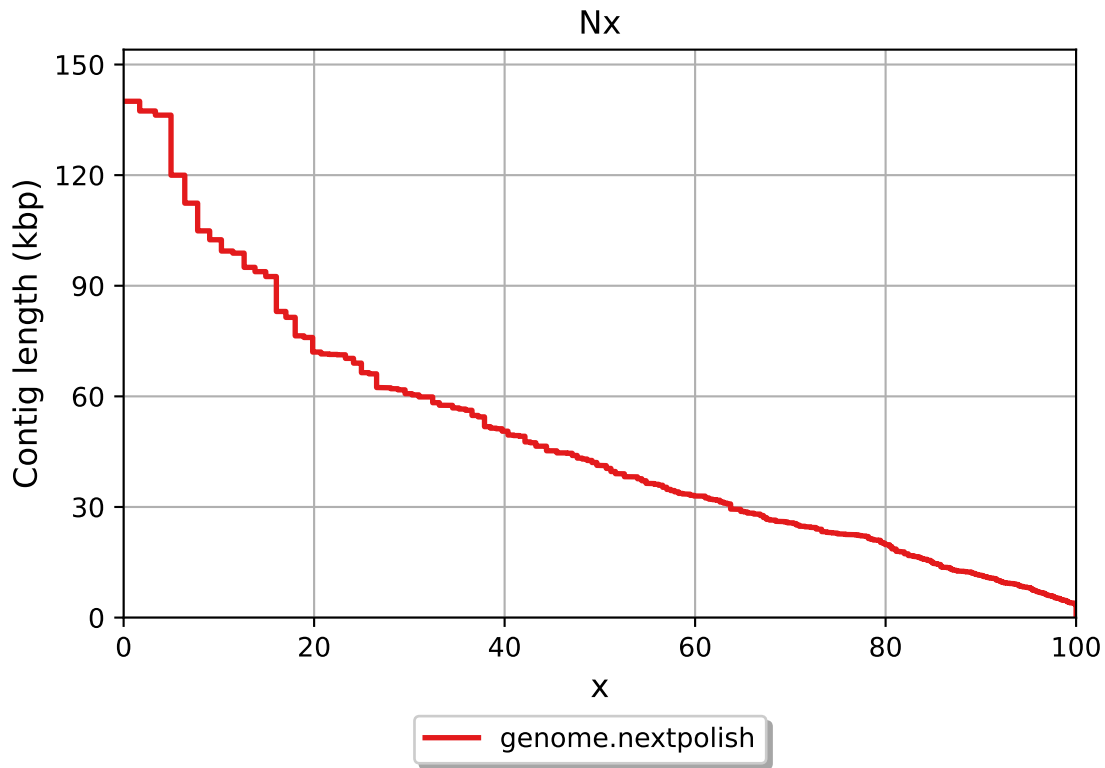


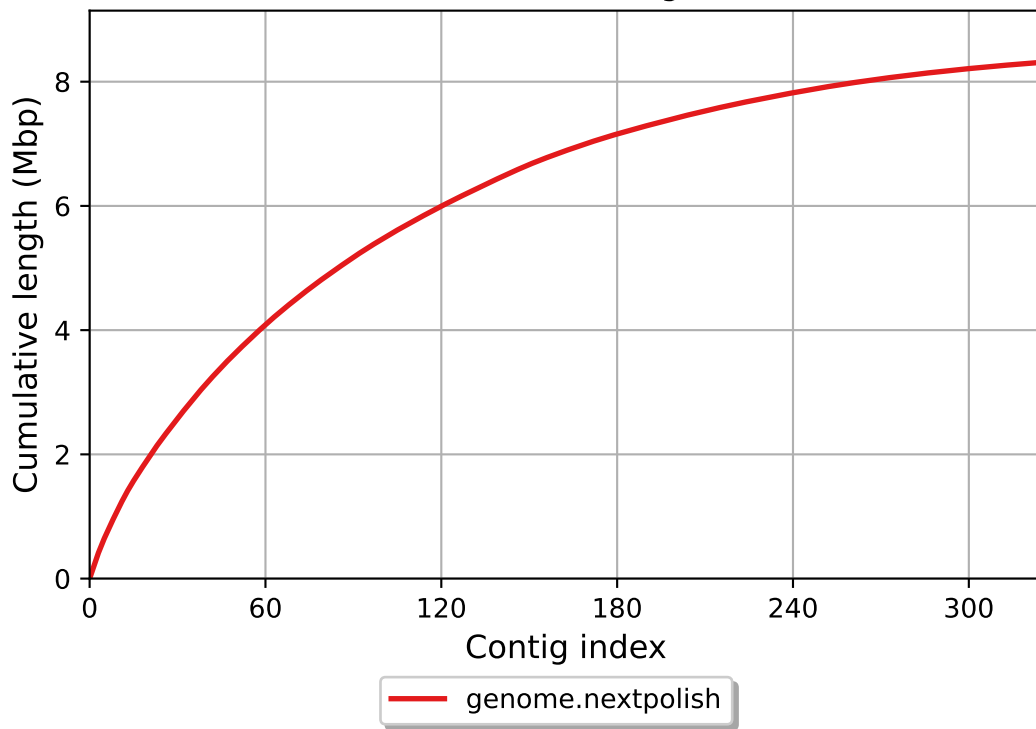
# Report

	genome.nextpolish
# contigs (>= 0 bp)	325
# contigs (>= 1000 bp)	325
# contigs (>= 5000 bp)	293
# contigs (>= 10000 bp)	221
# contigs (>= 25000 bp)	116
# contigs (>= 50000 bp)	44
Total length (>= 0 bp)	8312718
Total length (>= 1000 bp)	8312718
Total length (>= 5000 bp)	8176982
Total length (>= 10000 bp)	7645474
Total length (>= 25000 bp)	5896397
Total length (>= 50000 bp)	3354491
# contigs	325
Largest contig	140034
Total length	8312718
GC (%)	57.53
N50	41283
N90	11447
auN	49478.3
L50	62
L90	206
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	11 + 0 part

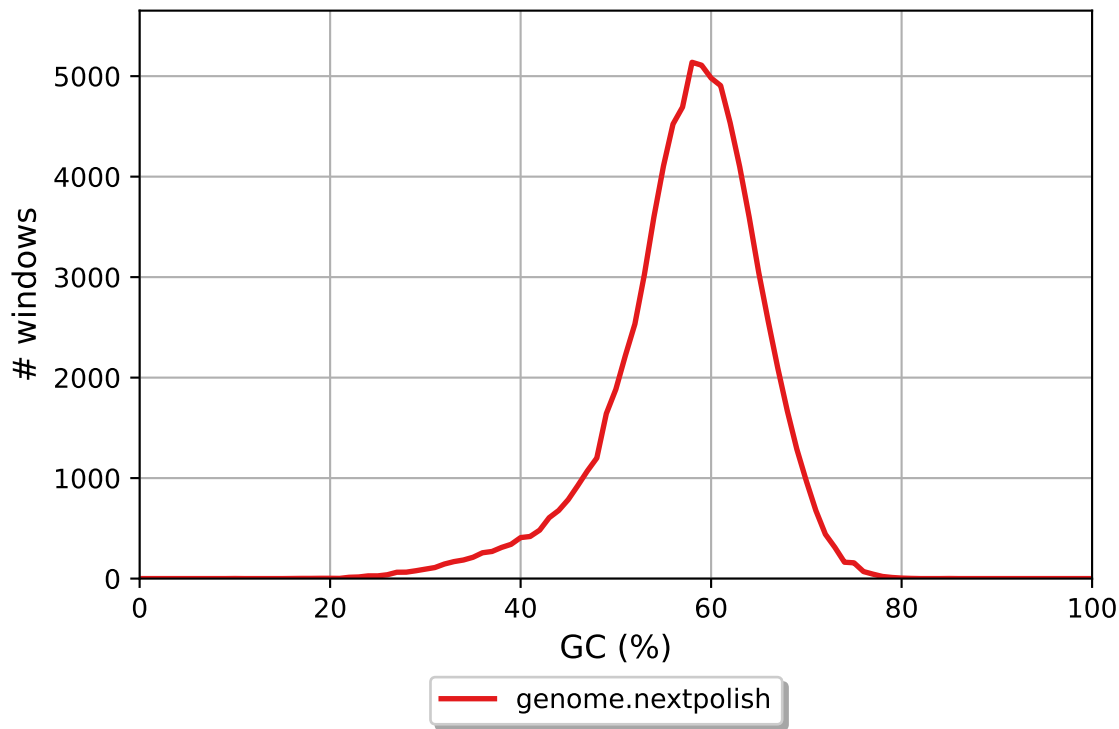
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

