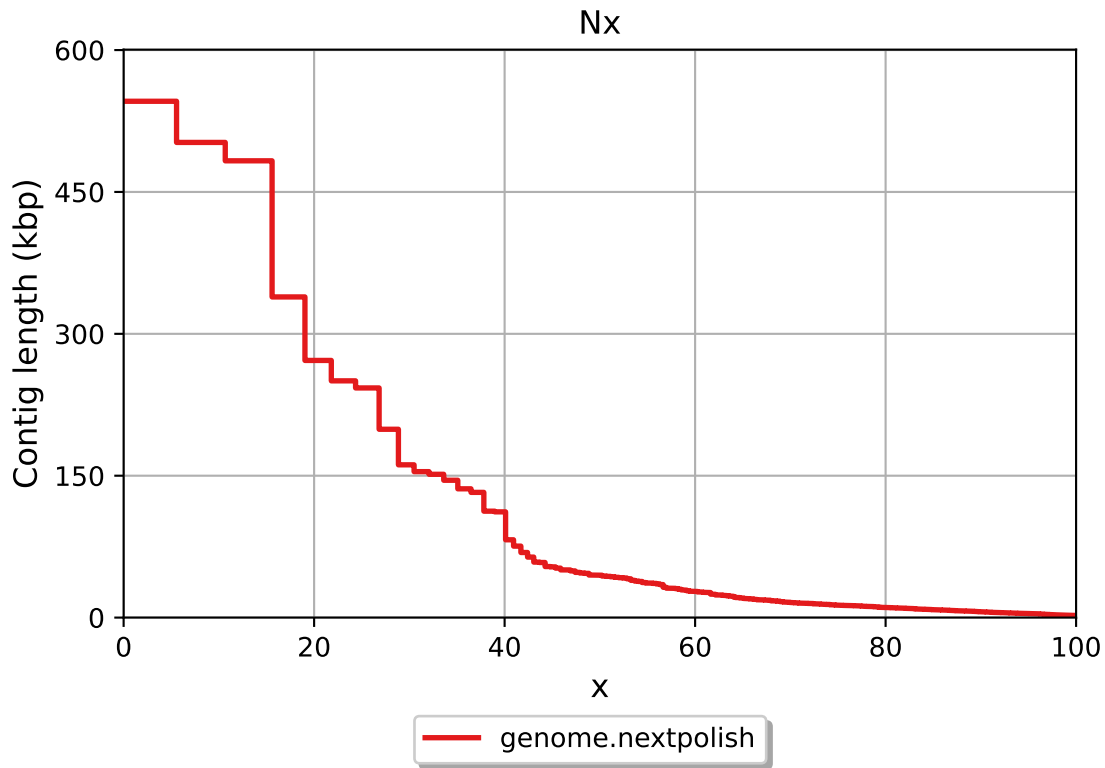


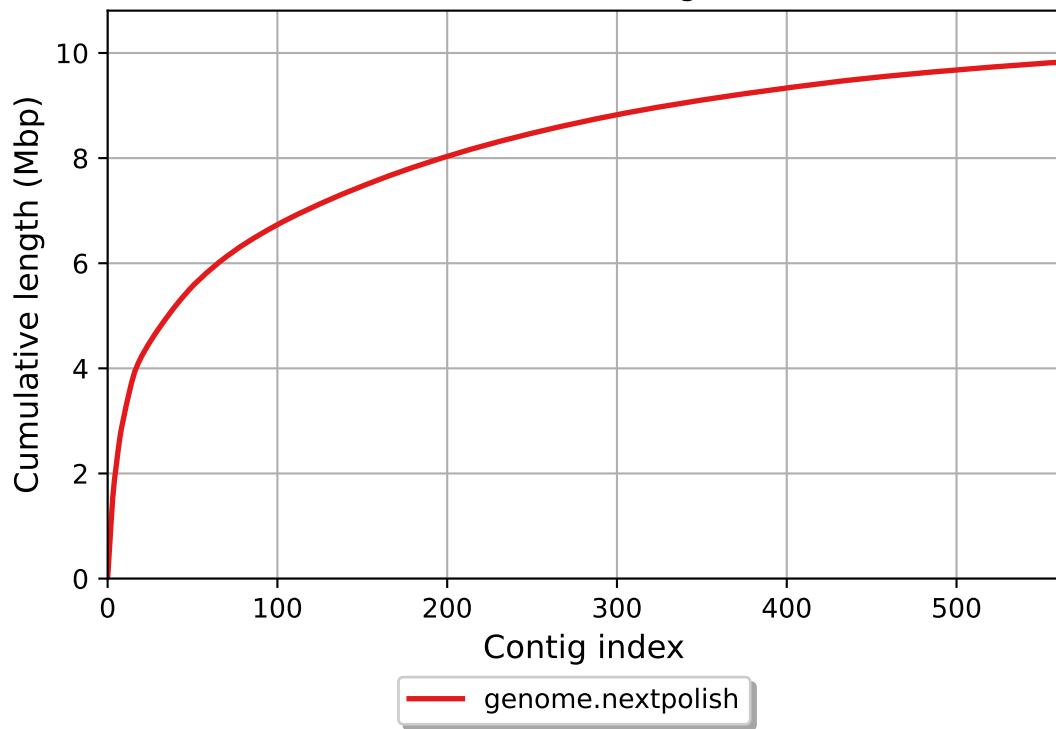
Report

	genome.nextpolish
# contigs (>= 0 bp)	561
# contigs (>= 1000 bp)	559
# contigs (>= 5000 bp)	351
# contigs (>= 10000 bp)	201
# contigs (>= 25000 bp)	67
# contigs (>= 50000 bp)	27
Total length (>= 0 bp)	9823570
Total length (>= 1000 bp)	9822144
Total length (>= 5000 bp)	9109778
Total length (>= 10000 bp)	8043718
Total length (>= 25000 bp)	6053603
Total length (>= 50000 bp)	4607592
# contigs	561
Largest contig	545865
Total length	9823570
GC (%)	54.41
N50	44794
N90	6194
auN	145250.5
L50	34
L90	303
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	21 + 8 part

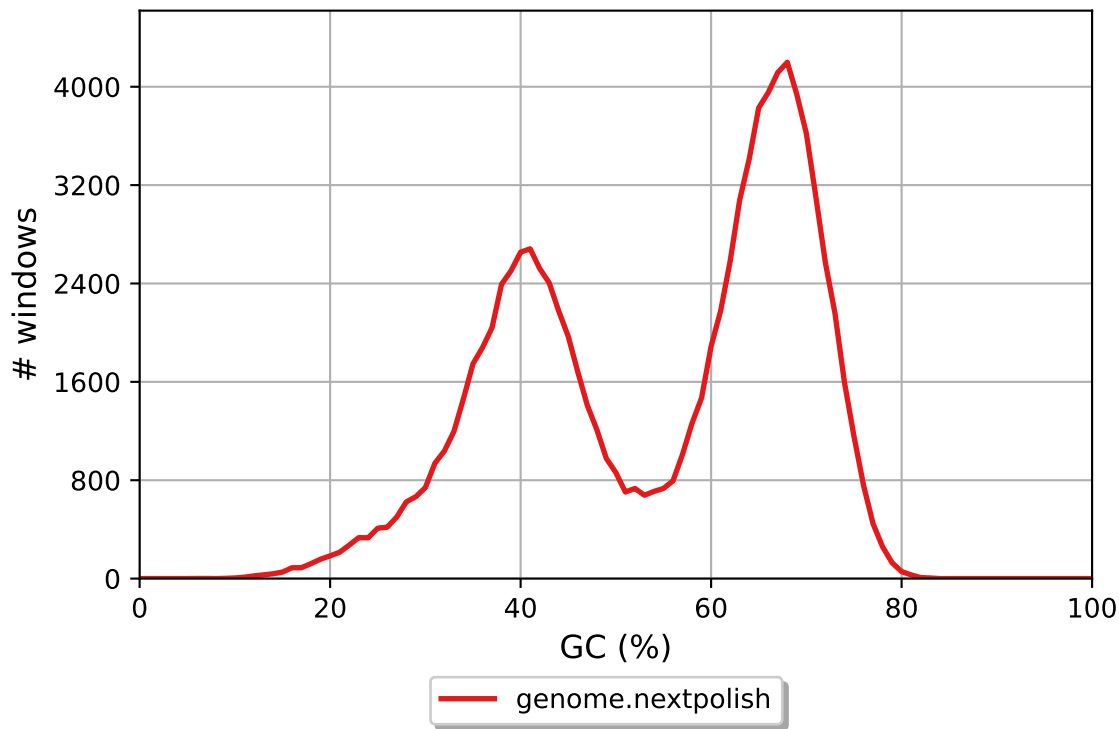
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

