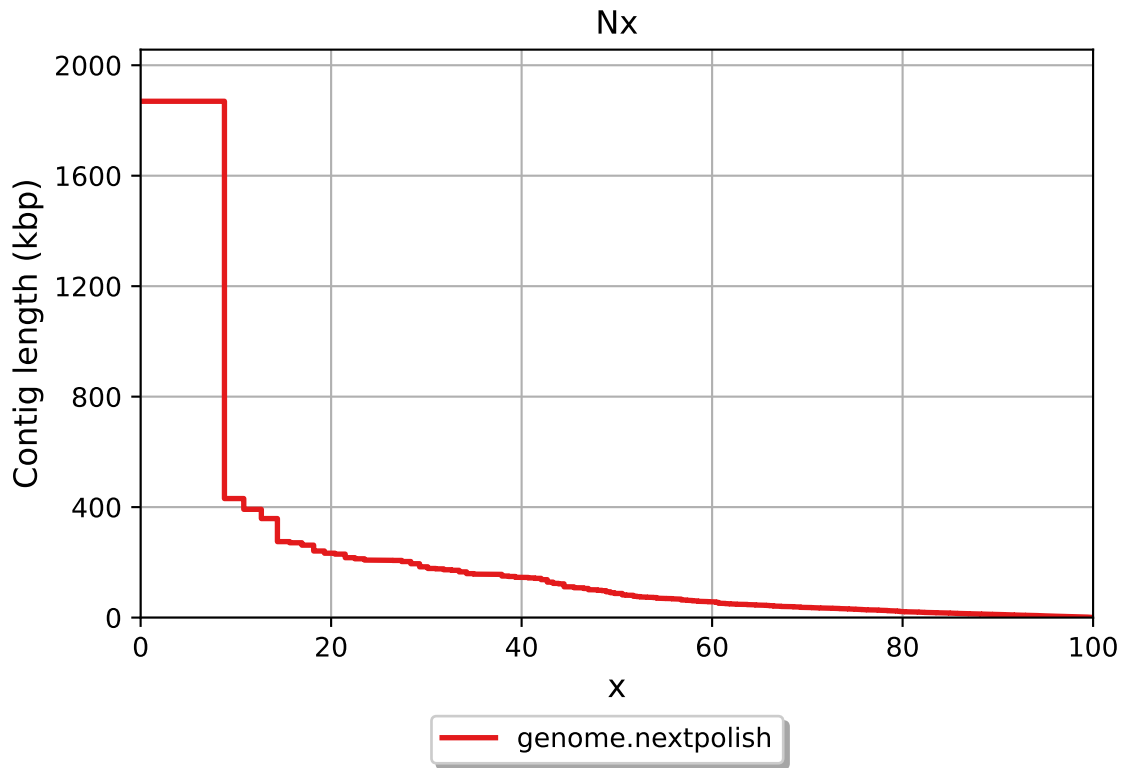


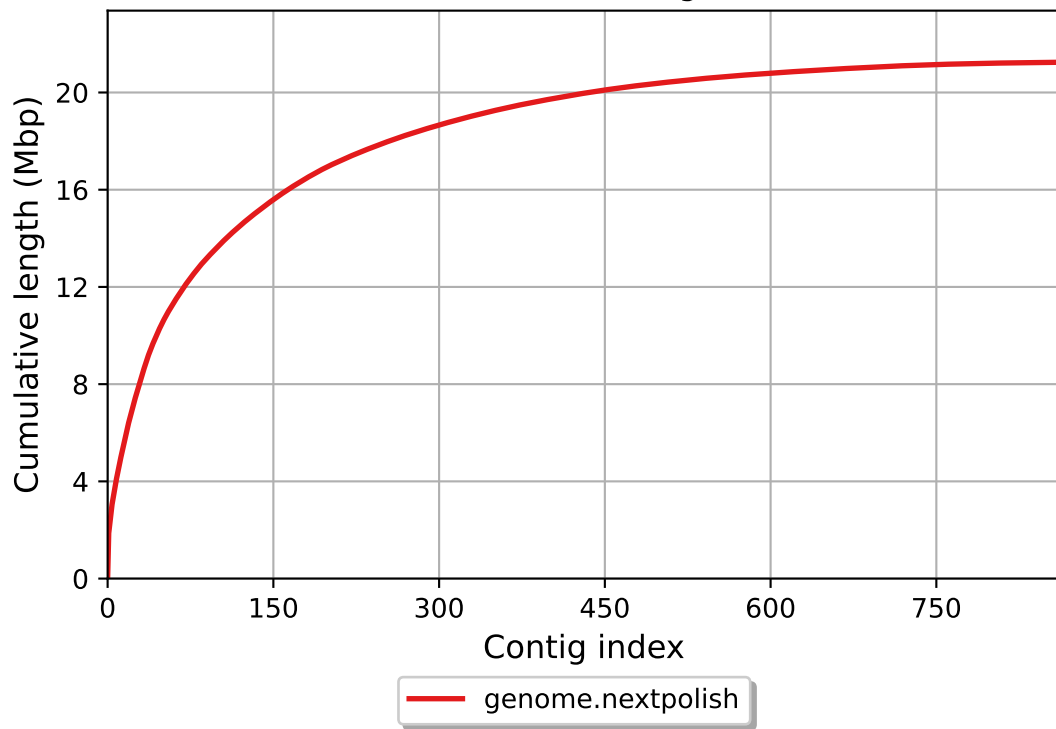
Report

	genome.nextpolish
# contigs (>= 0 bp)	874
# contigs (>= 1000 bp)	786
# contigs (>= 5000 bp)	504
# contigs (>= 10000 bp)	362
# contigs (>= 25000 bp)	187
# contigs (>= 50000 bp)	88
Total length (>= 0 bp)	21248044
Total length (>= 1000 bp)	21192435
Total length (>= 5000 bp)	20411174
Total length (>= 10000 bp)	19380021
Total length (>= 25000 bp)	16660331
Total length (>= 50000 bp)	13095005
# contigs	862
Largest contig	1869532
Total length	21243573
GC (%)	43.56
N50	87333
N90	11205
auN	266746.8
L50	51
L90	338
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	39 + 1 part

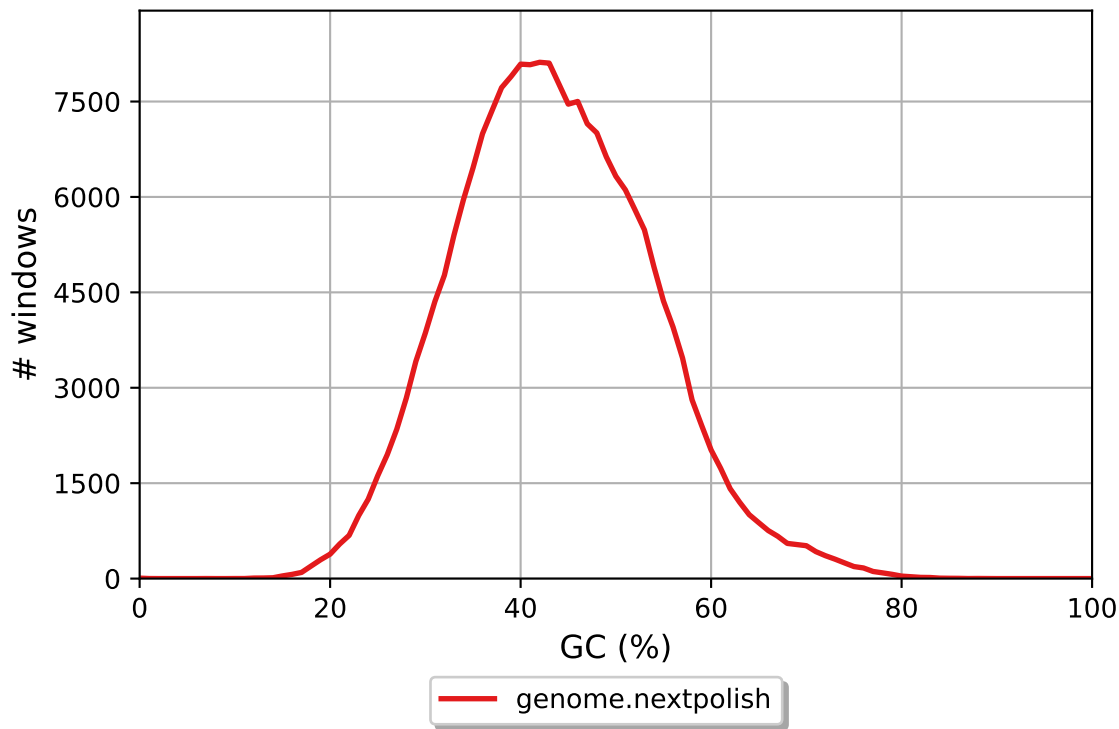
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

