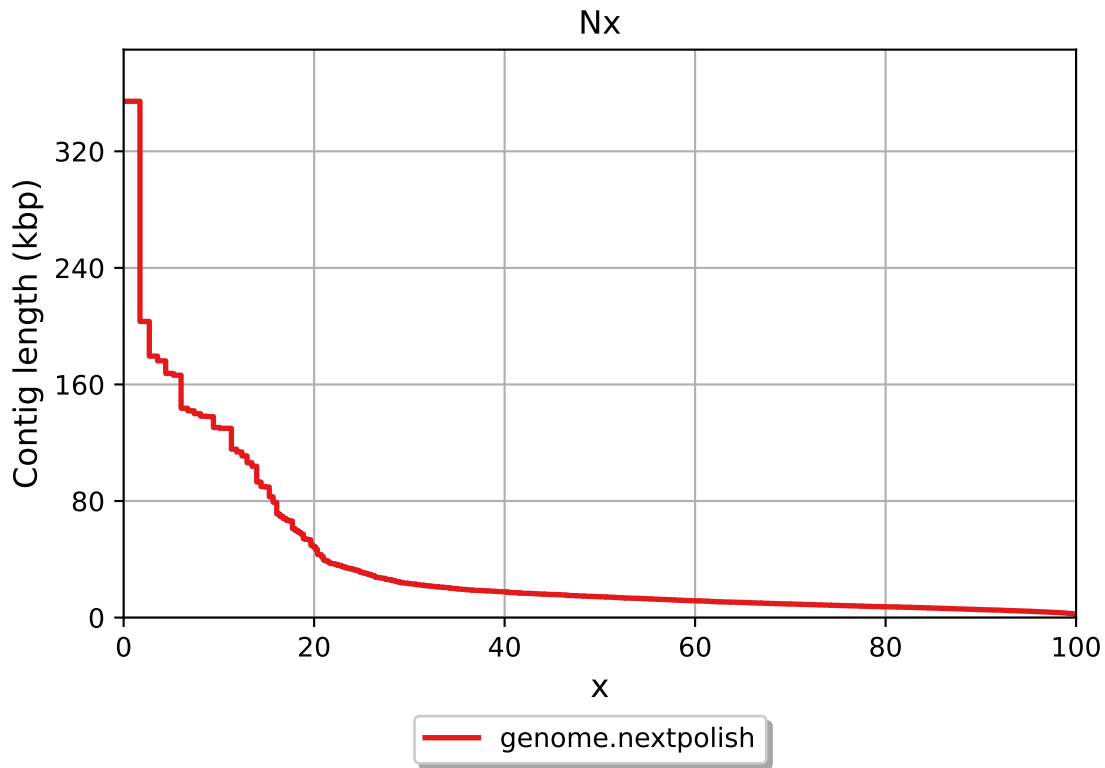


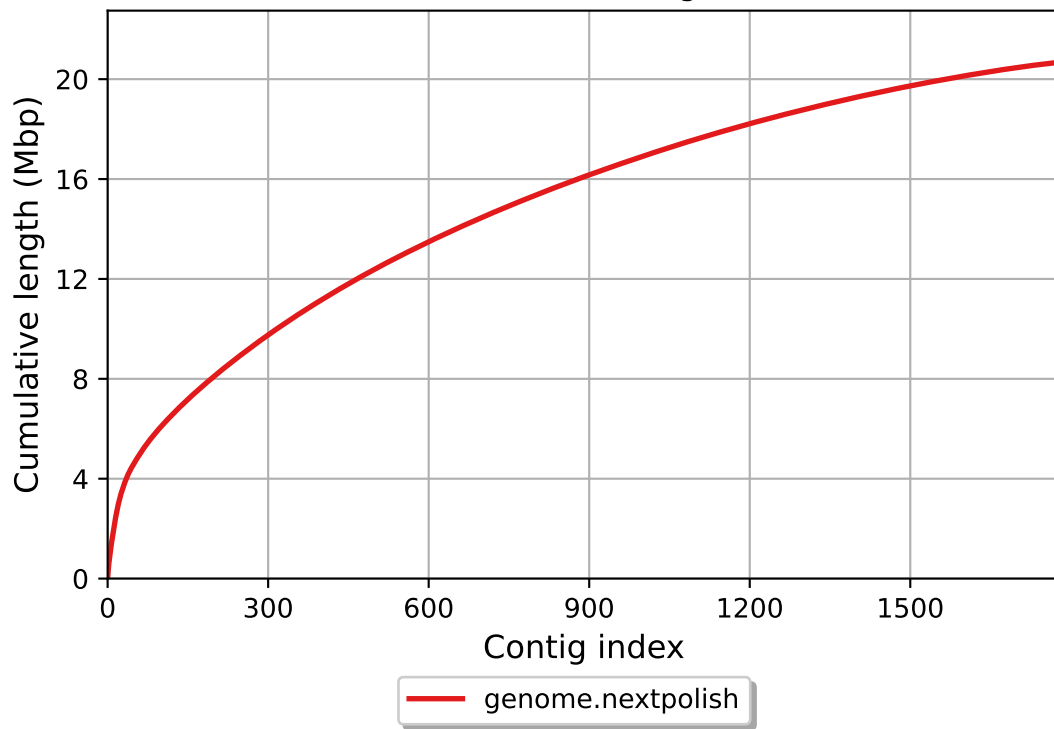
# Report

	genome.nextpolish
# contigs (>= 0 bp)	1780
# contigs (>= 1000 bp)	1778
# contigs (>= 5000 bp)	1363
# contigs (>= 10000 bp)	631
# contigs (>= 25000 bp)	92
# contigs (>= 50000 bp)	36
Total length (>= 0 bp)	20676716
Total length (>= 1000 bp)	20675020
Total length (>= 5000 bp)	19100309
Total length (>= 10000 bp)	13803628
Total length (>= 25000 bp)	5898481
Total length (>= 50000 bp)	4063007
# contigs	1780
Largest contig	354340
Total length	20676716
GC (%)	49.30
N50	14333
N90	5455
auN	39223.0
L50	340
L90	1270
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	41 + 7 part

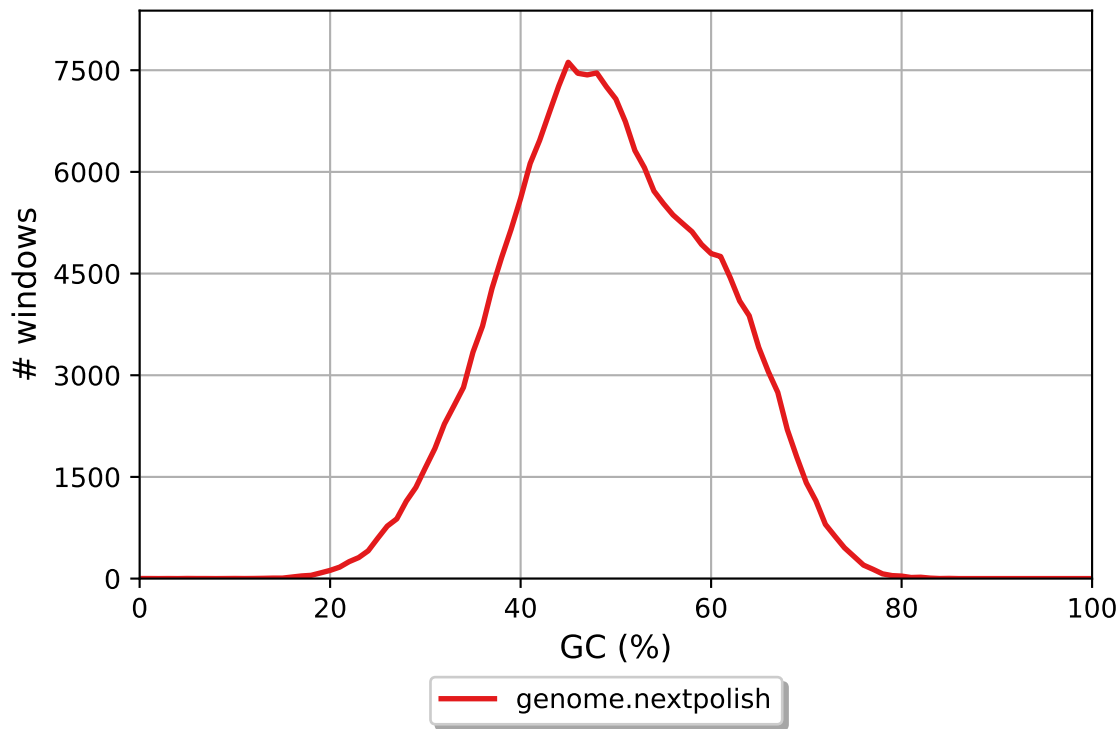
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

