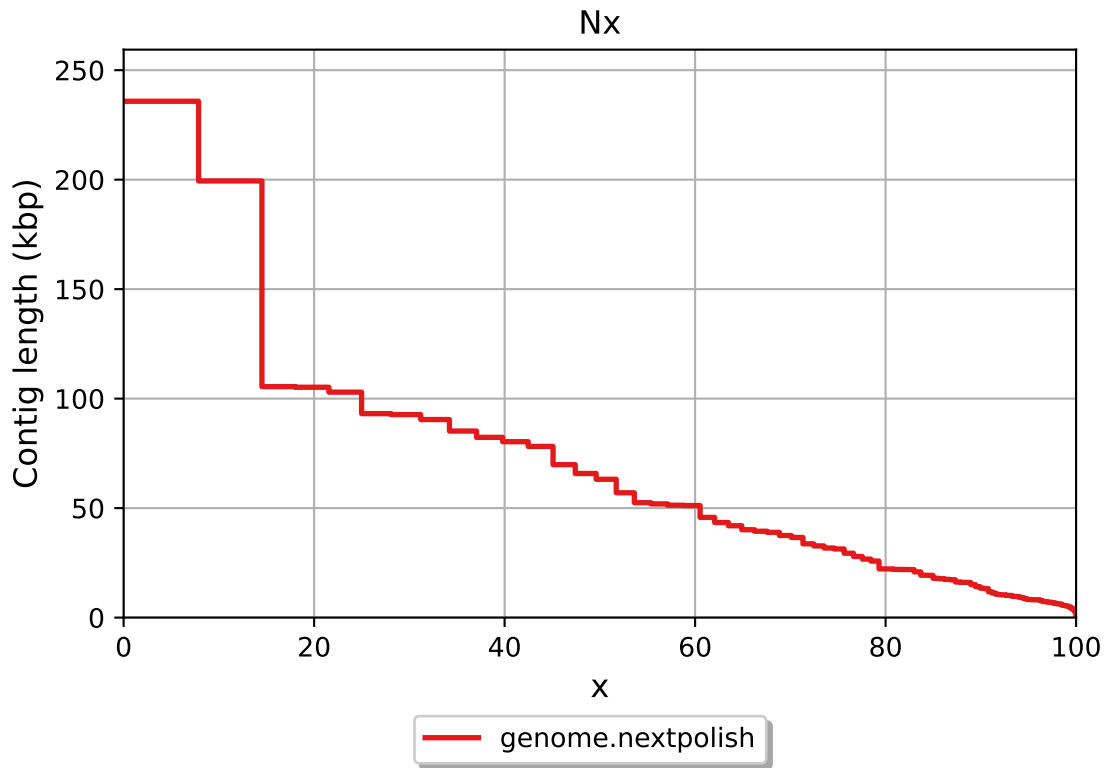


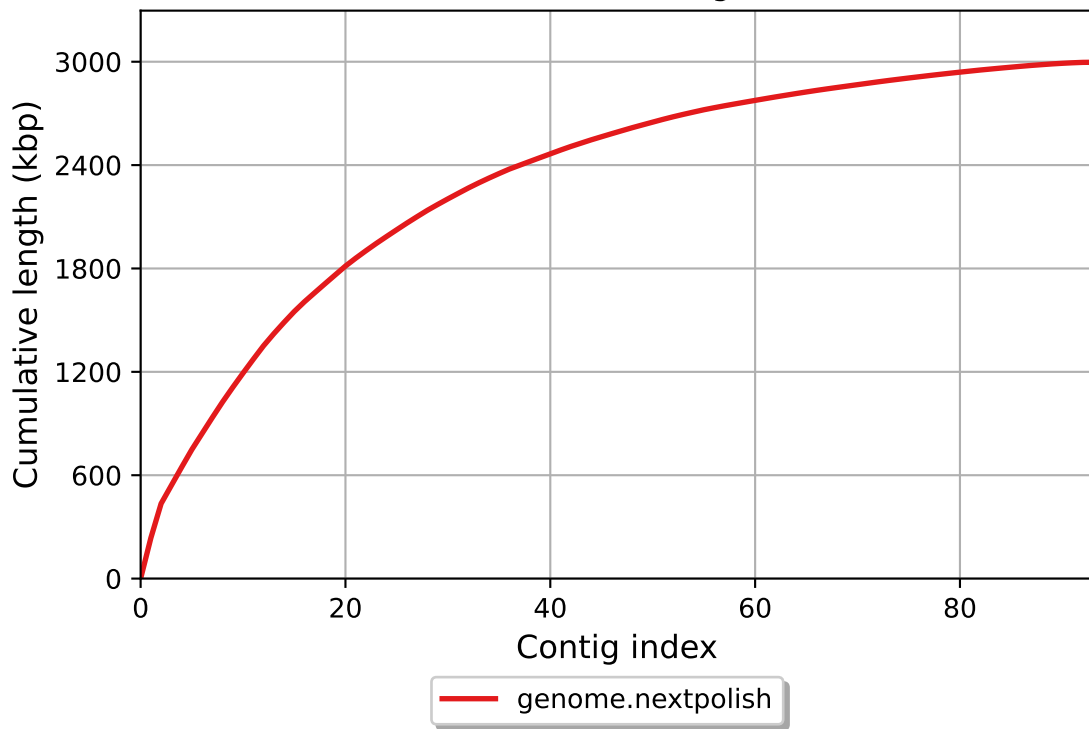
Report

	genome.nextpolish
# contigs (>= 0 bp)	93
# contigs (>= 1000 bp)	92
# contigs (>= 5000 bp)	86
# contigs (>= 10000 bp)	62
# contigs (>= 25000 bp)	36
# contigs (>= 50000 bp)	20
Total length (>= 0 bp)	2996893
Total length (>= 1000 bp)	2996306
Total length (>= 5000 bp)	2973801
Total length (>= 10000 bp)	2795929
Total length (>= 25000 bp)	2376854
Total length (>= 50000 bp)	1813863
# contigs	93
Largest contig	235799
Total length	2996893
GC (%)	59.35
N50	63170
N90	13445
auN	78882.3
L50	15
L90	54
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 1 part

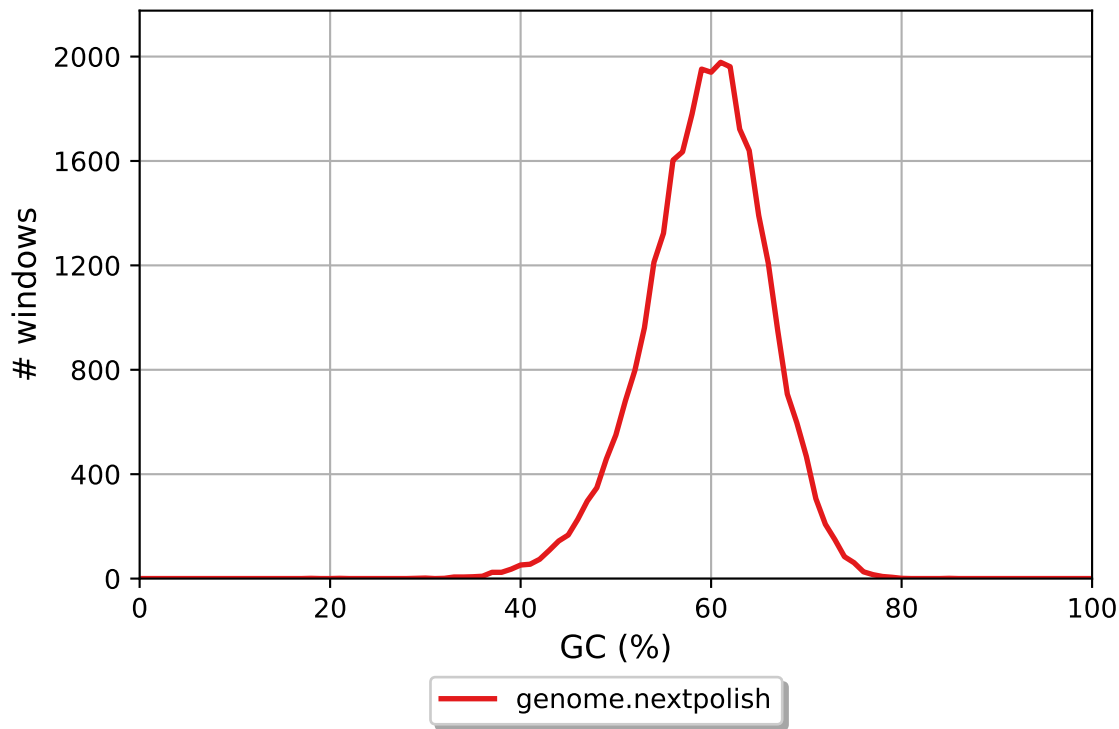
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

