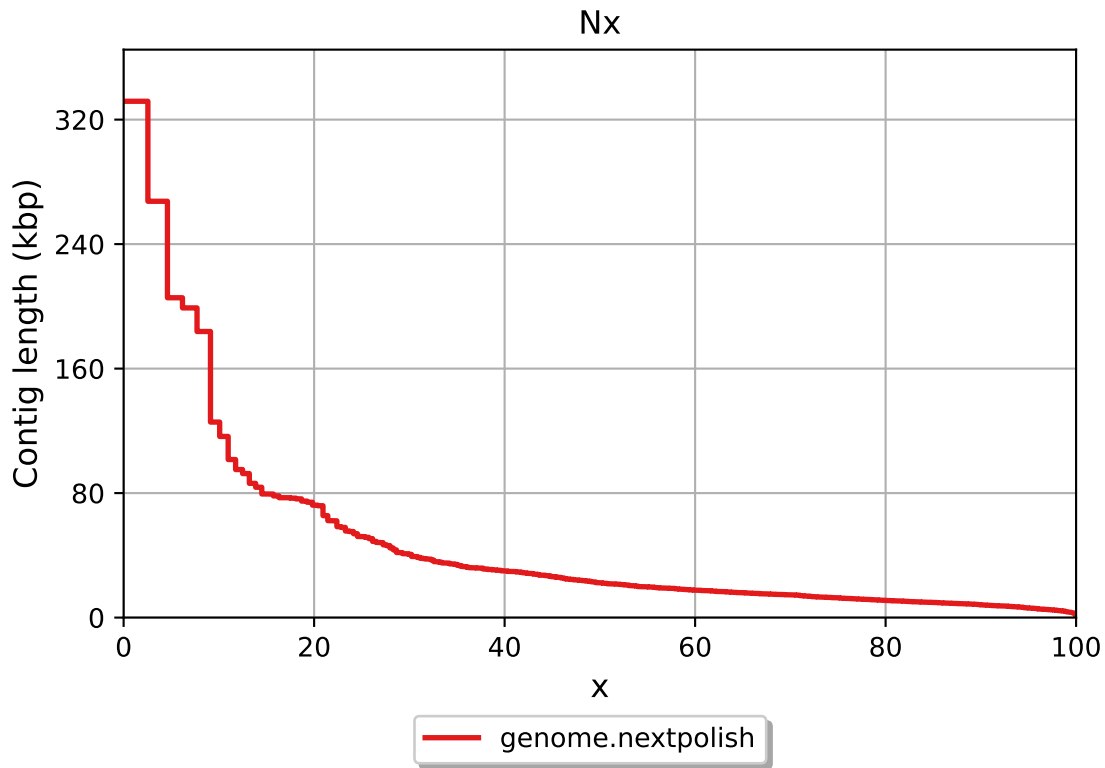


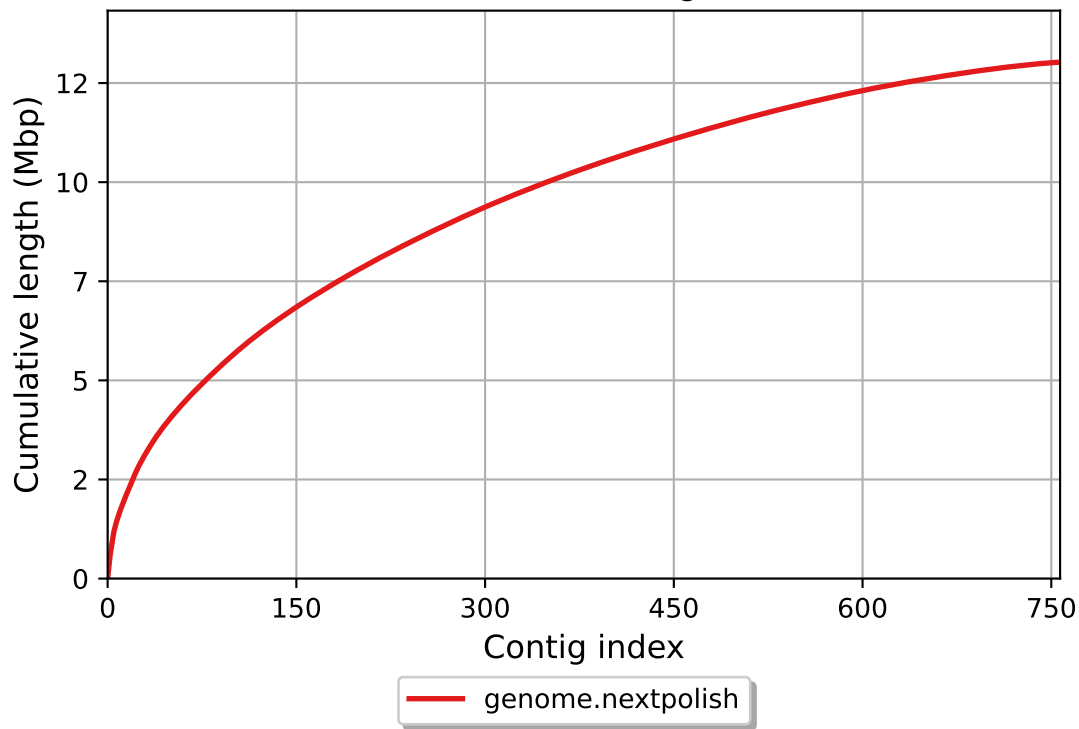
Report

	genome.nextpolish
# contigs (>= 0 bp)	757
# contigs (>= 1000 bp)	755
# contigs (>= 5000 bp)	668
# contigs (>= 10000 bp)	431
# contigs (>= 25000 bp)	115
# contigs (>= 50000 bp)	35
Total length (>= 0 bp)	13024239
Total length (>= 1000 bp)	13023102
Total length (>= 5000 bp)	12695389
Total length (>= 10000 bp)	10901742
Total length (>= 25000 bp)	6051454
Total length (>= 50000 bp)	3403854
# contigs	757
Largest contig	331789
Total length	13024239
GC (%)	53.88
N50	22421
N90	8186
auN	50408.1
L50	135
L90	521
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	36 + 3 part

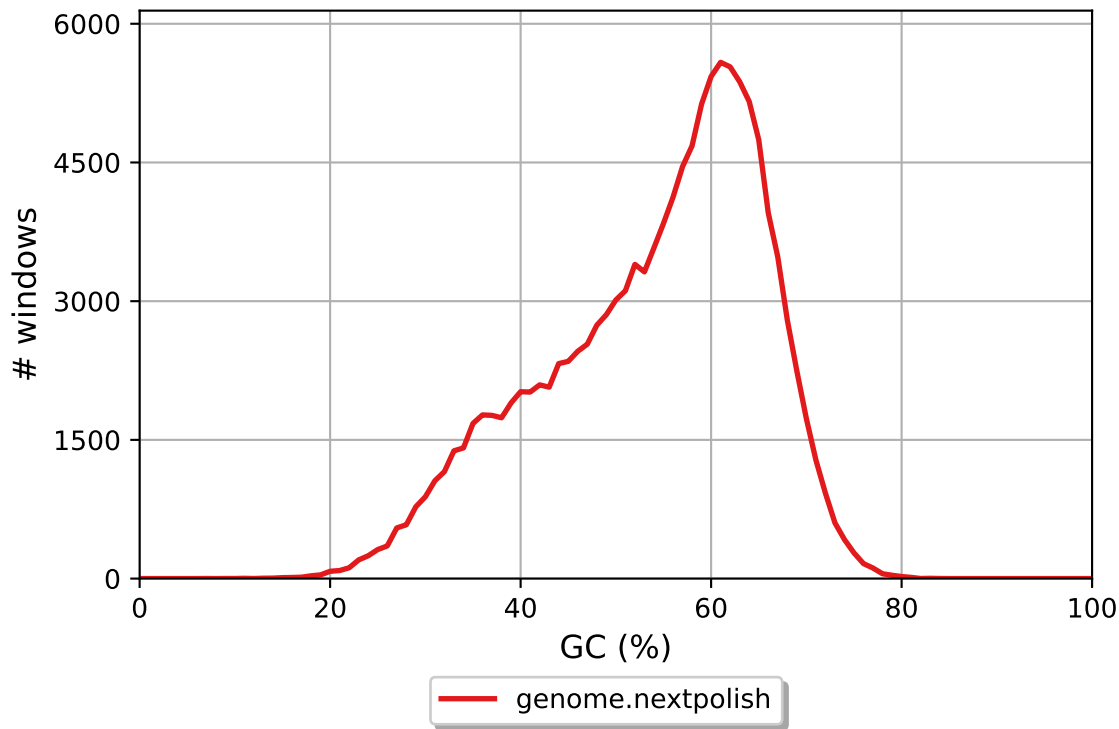
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

