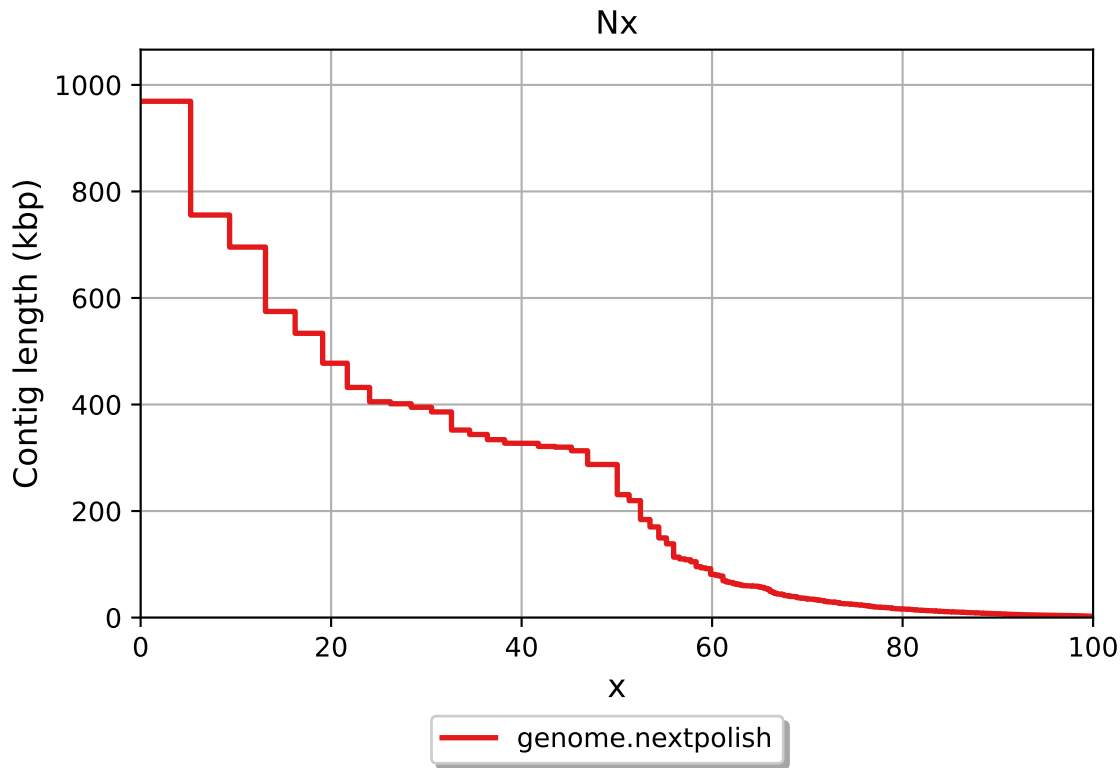


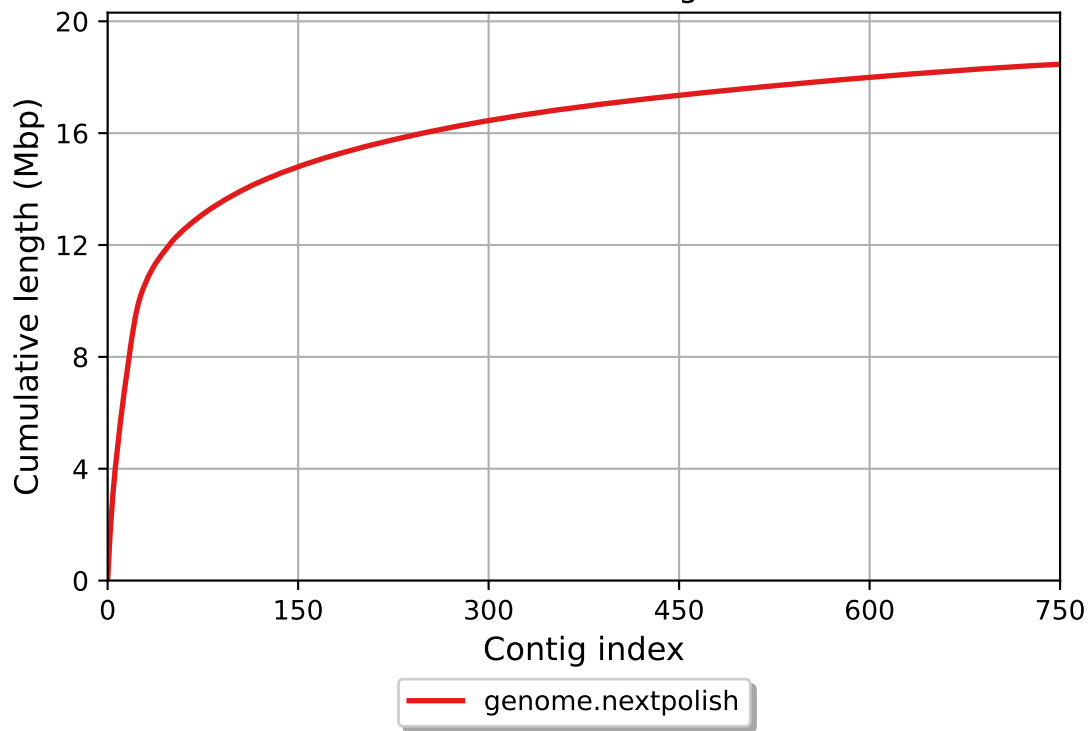
# Report

	genome.nextpolish
# contigs (>= 0 bp)	750
# contigs (>= 1000 bp)	749
# contigs (>= 5000 bp)	435
# contigs (>= 10000 bp)	242
# contigs (>= 25000 bp)	100
# contigs (>= 50000 bp)	52
Total length (>= 0 bp)	18463556
Total length (>= 1000 bp)	18462772
Total length (>= 5000 bp)	17281198
Total length (>= 10000 bp)	15941593
Total length (>= 25000 bp)	13813158
Total length (>= 50000 bp)	12196036
# contigs	750
Largest contig	969279
Total length	18463556
GC (%)	42.17
N50	287299
N90	7199
auN	278937.5
L50	21
L90	323
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	47 + 6 part

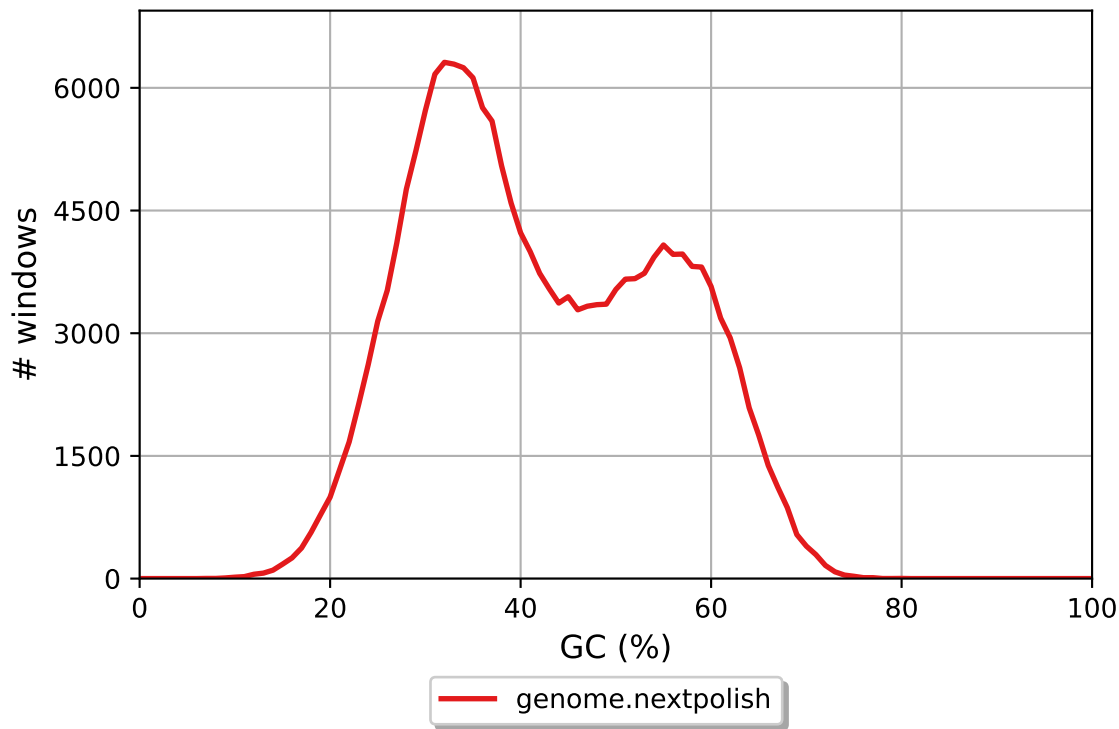
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

