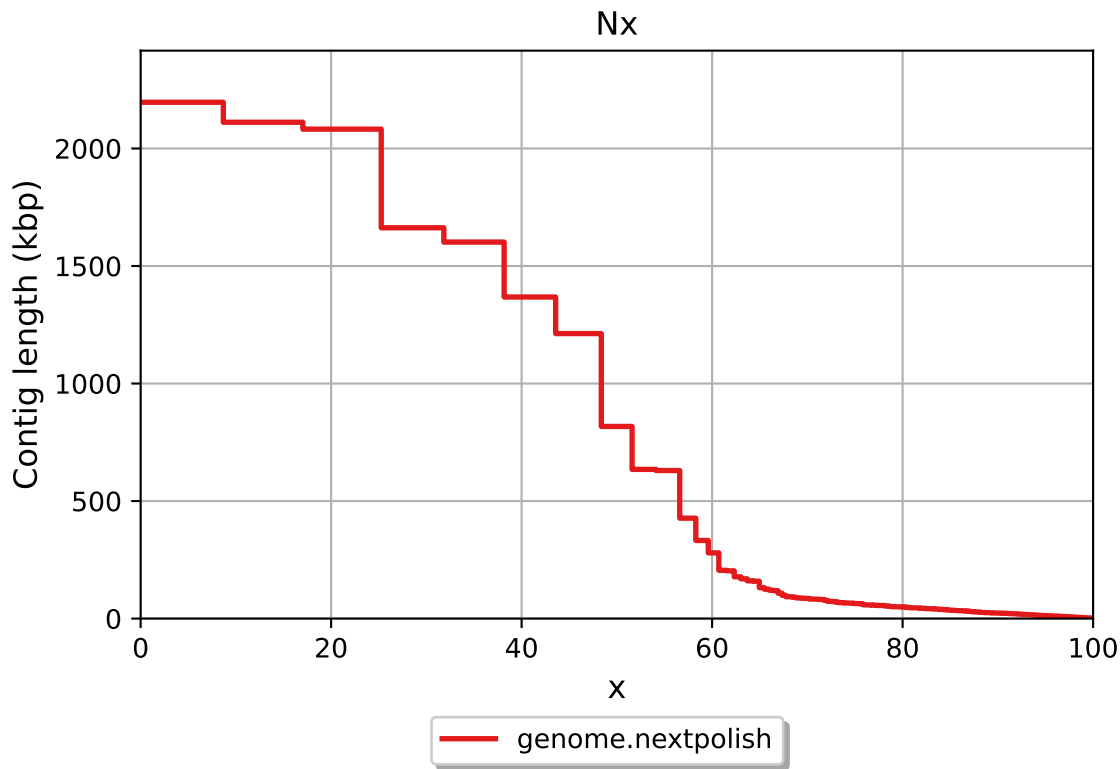


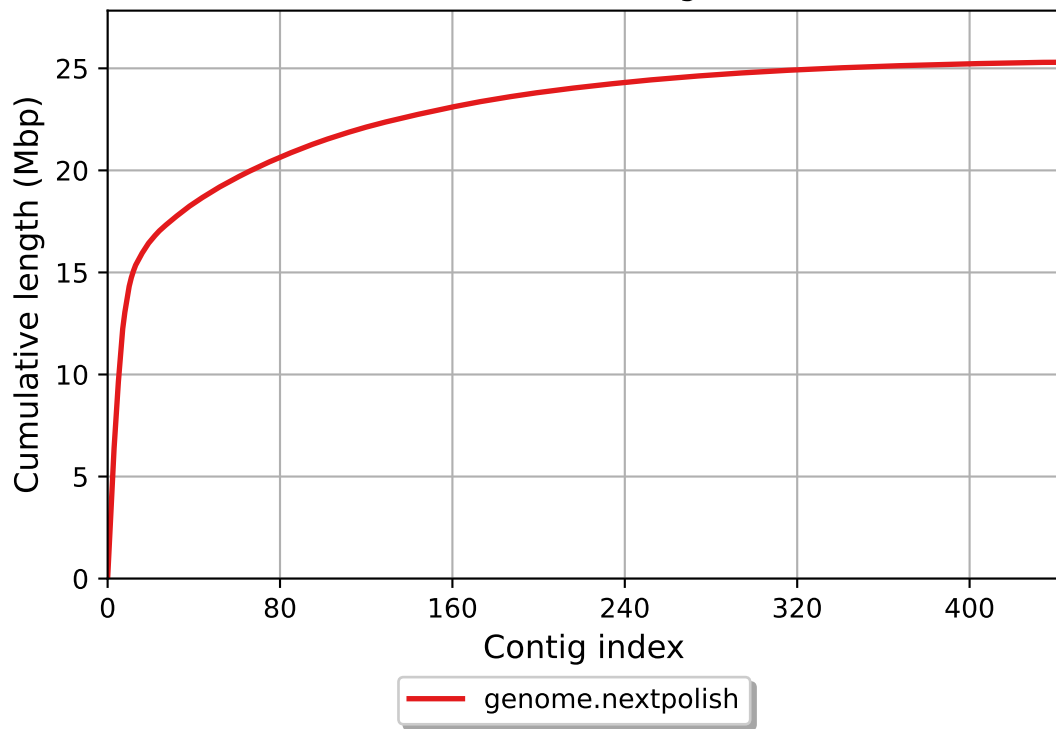
# Report

	genome.nextpolish
# contigs (>= 0 bp)	442
# contigs (>= 1000 bp)	432
# contigs (>= 5000 bp)	326
# contigs (>= 10000 bp)	249
# contigs (>= 25000 bp)	137
# contigs (>= 50000 bp)	69
Total length (>= 0 bp)	25297531
Total length (>= 1000 bp)	25290819
Total length (>= 5000 bp)	24959117
Total length (>= 10000 bp)	24403952
Total length (>= 25000 bp)	22574883
Total length (>= 50000 bp)	20119667
# contigs	442
Largest contig	2196639
Total length	25297531
GC (%)	44.24
N50	817472
N90	23325
auN	978329.1
L50	8
L90	146
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	79 + 3 part

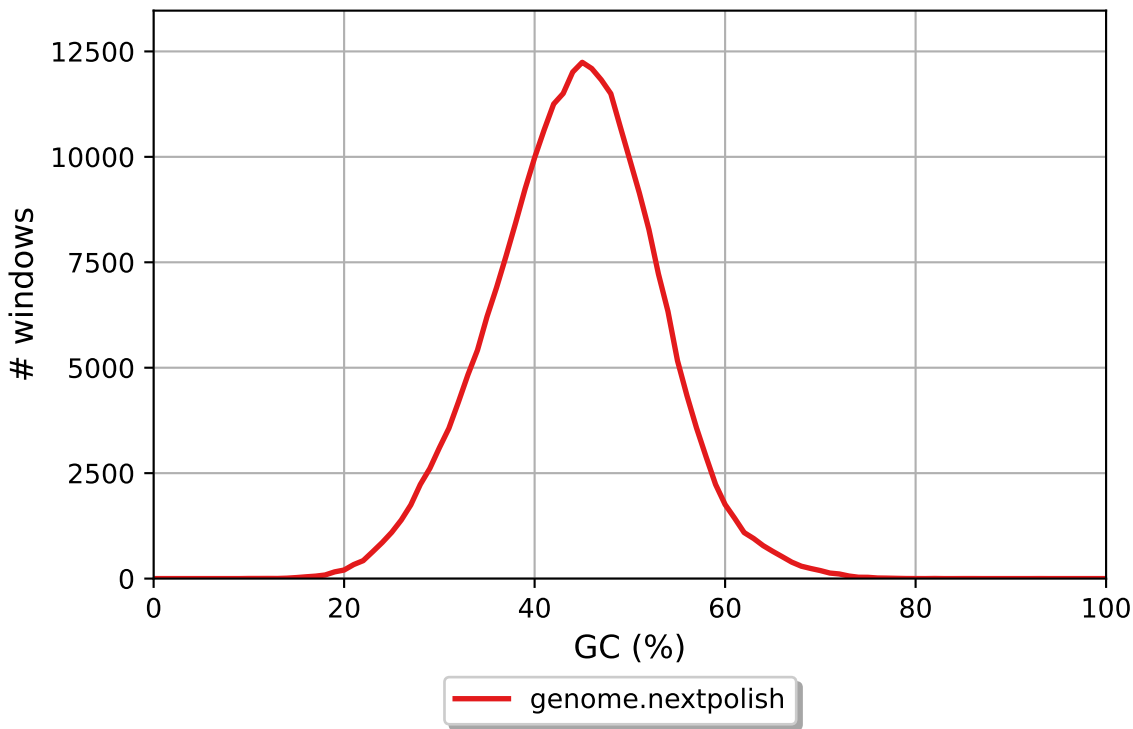
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

