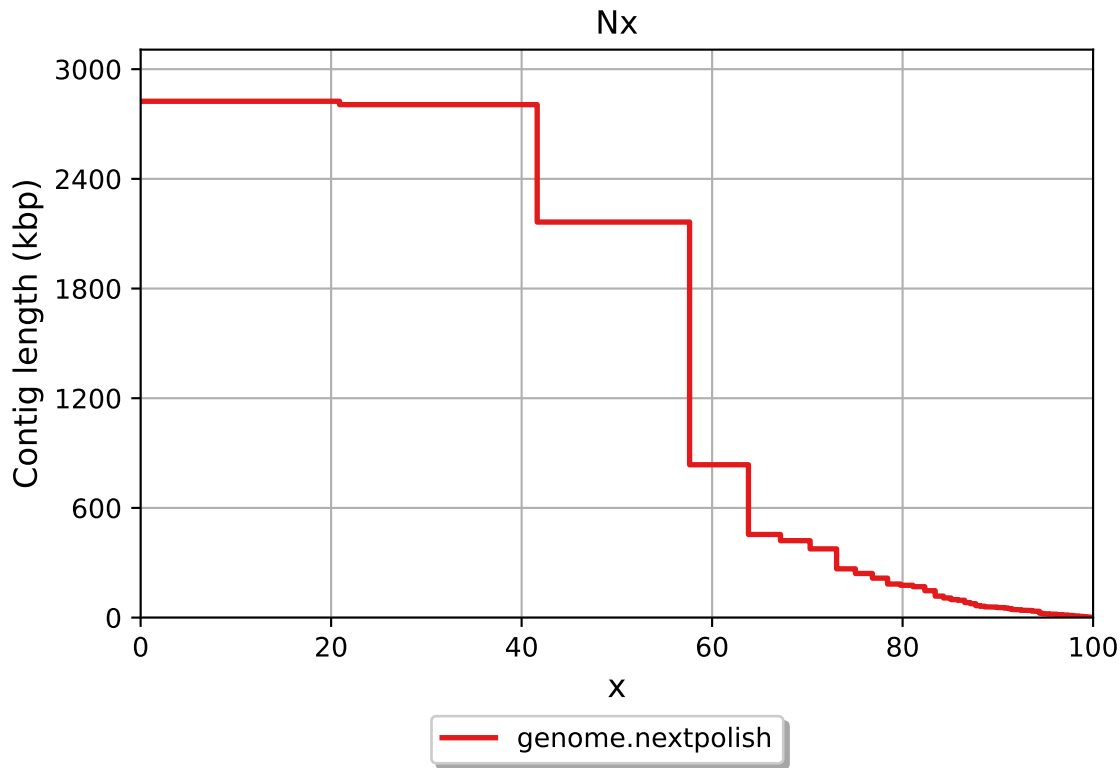


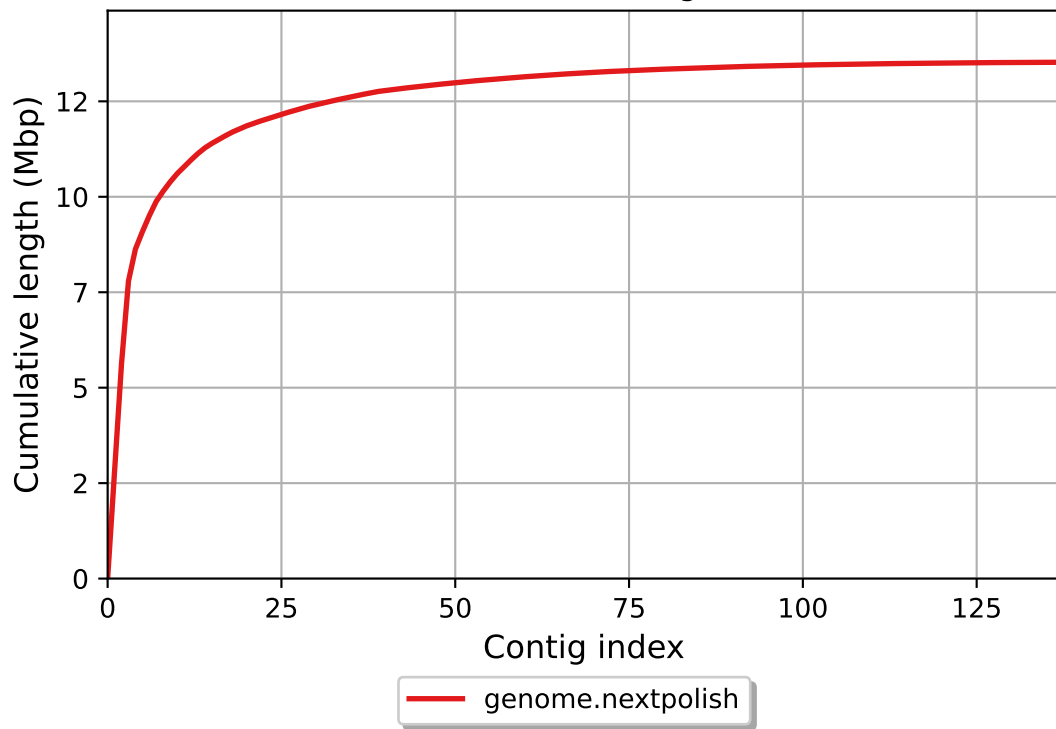
Report

	genome.nextpolish
# contigs (>= 0 bp)	137
# contigs (>= 1000 bp)	131
# contigs (>= 5000 bp)	92
# contigs (>= 10000 bp)	68
# contigs (>= 25000 bp)	39
# contigs (>= 50000 bp)	28
Total length (>= 0 bp)	13523145
Total length (>= 1000 bp)	13518403
Total length (>= 5000 bp)	13415276
Total length (>= 10000 bp)	13241051
Total length (>= 25000 bp)	12760805
Total length (>= 50000 bp)	12320925
# contigs	137
Largest contig	2824428
Total length	13523145
GC (%)	41.41
N50	2163706
N90	55052
auN	1638569.5
L50	3
L90	26
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	58 + 0 part

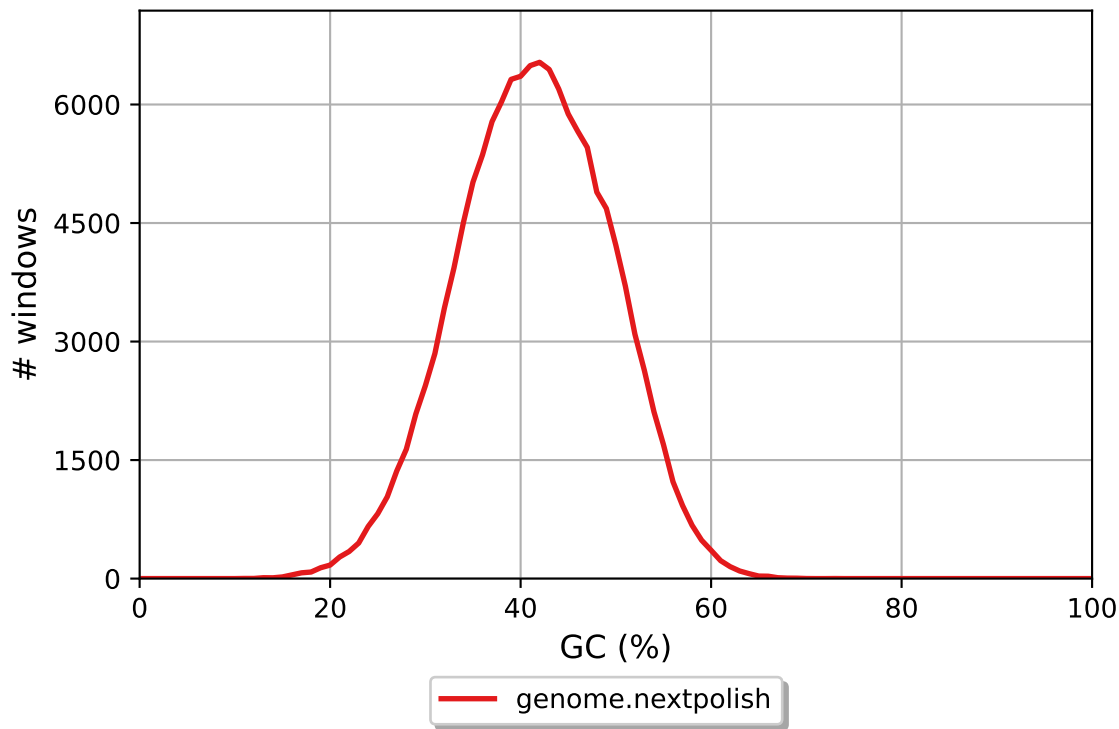
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

