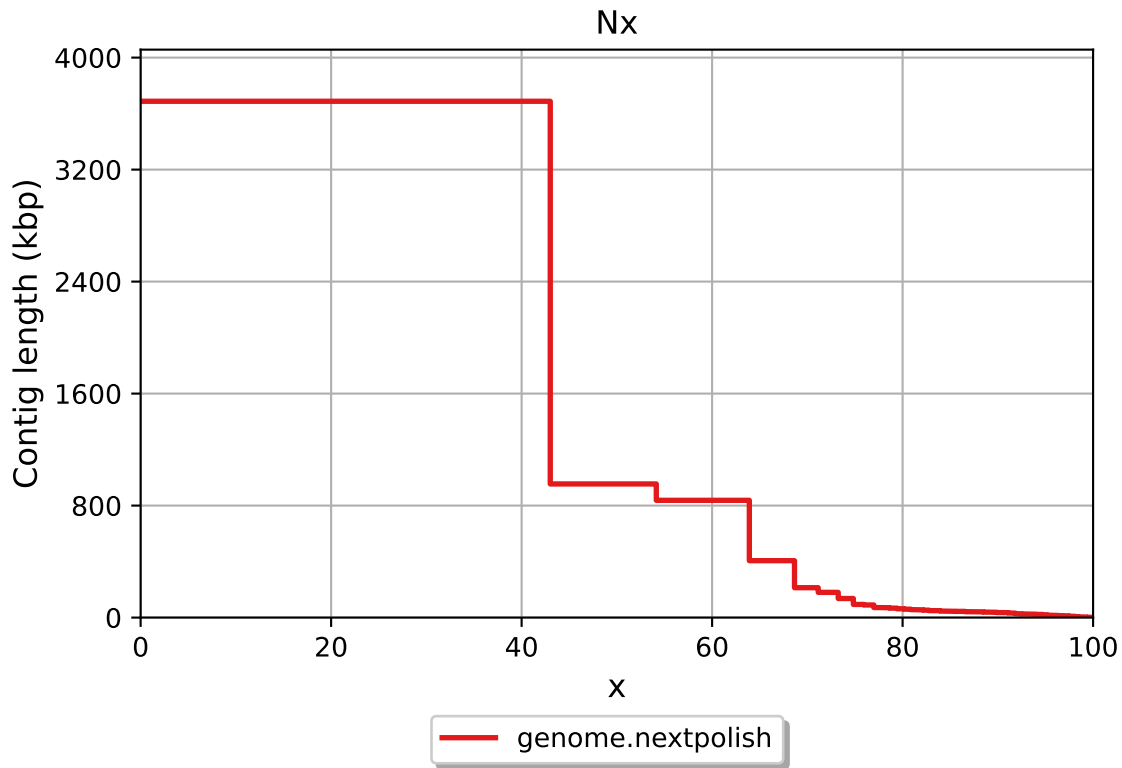


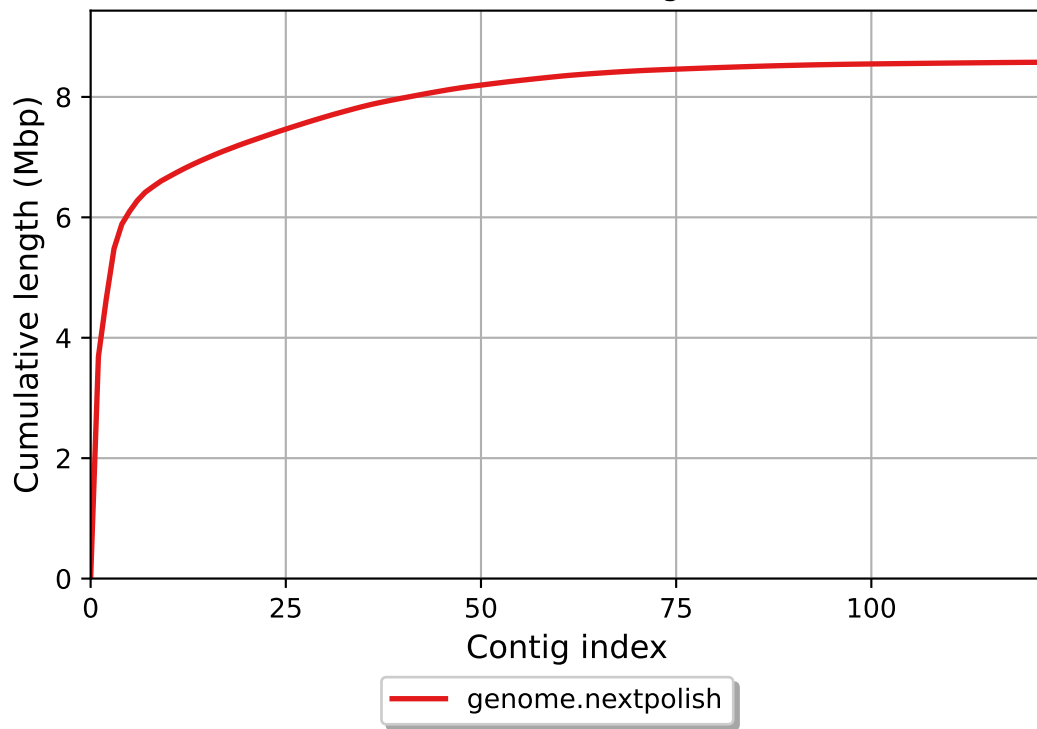
Report

	genome.nextpolish
# contigs (>= 0 bp)	122
# contigs (>= 1000 bp)	120
# contigs (>= 5000 bp)	75
# contigs (>= 10000 bp)	63
# contigs (>= 25000 bp)	39
# contigs (>= 50000 bp)	17
Total length (>= 0 bp)	8575413
Total length (>= 1000 bp)	8574225
Total length (>= 5000 bp)	8461082
Total length (>= 10000 bp)	8375235
Total length (>= 25000 bp)	7957664
Total length (>= 50000 bp)	7098897
# contigs	122
Largest contig	3688105
Total length	8575413
GC (%)	33.85
N50	954634
N90	35998
auN	1815505.9
L50	2
L90	32
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	30 + 7 part

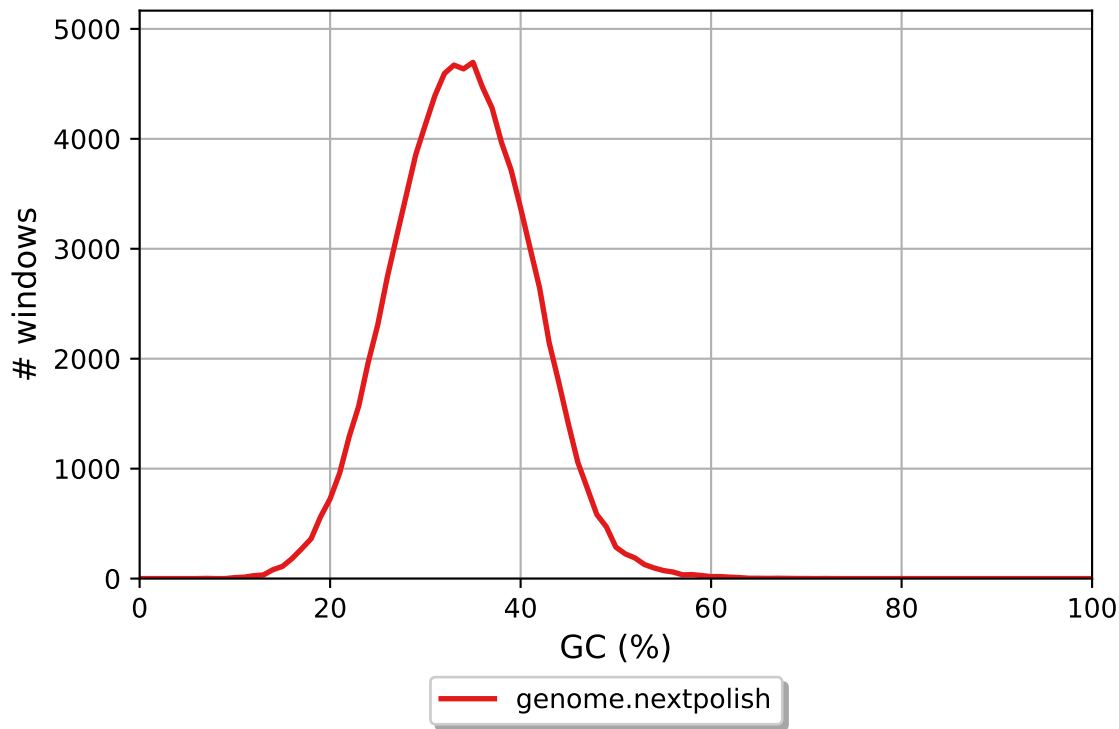
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

