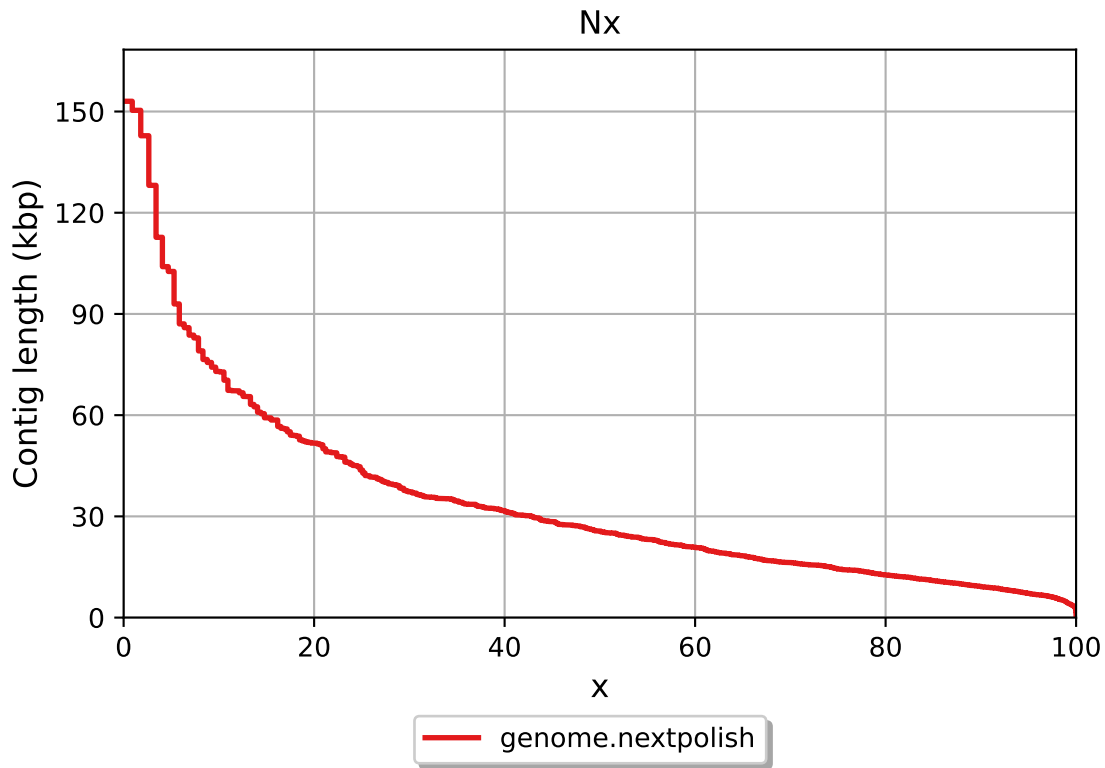


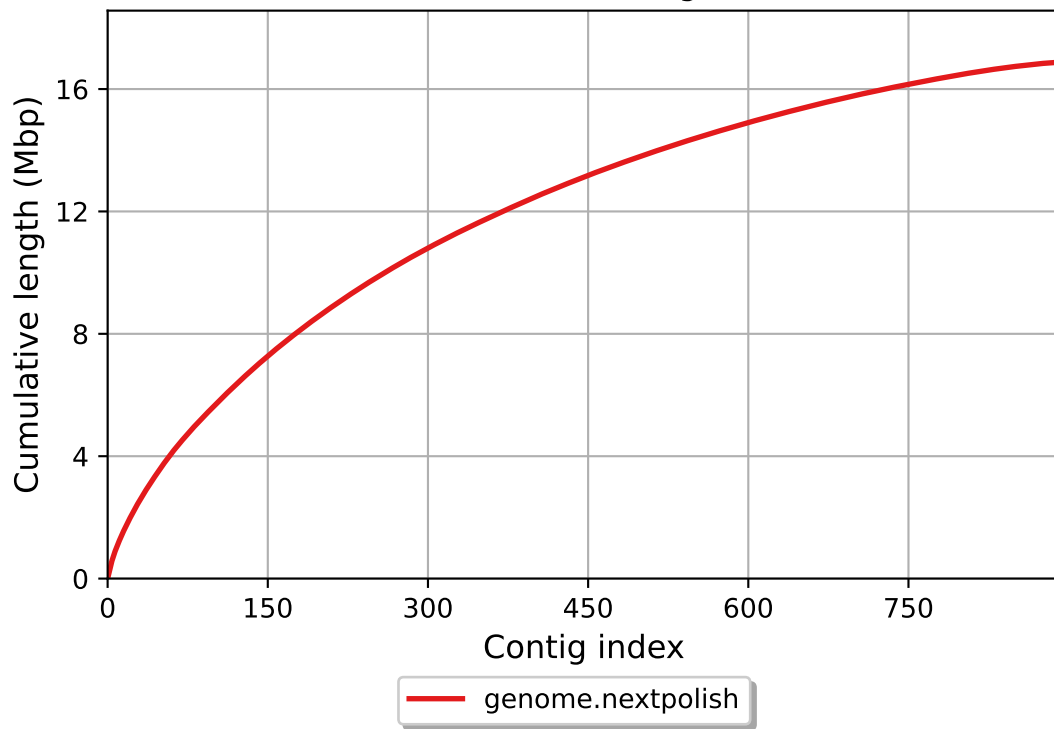
Report

	genome.nextpolish
# contigs (>= 0 bp)	892
# contigs (>= 1000 bp)	888
# contigs (>= 5000 bp)	833
# contigs (>= 10000 bp)	592
# contigs (>= 25000 bp)	203
# contigs (>= 50000 bp)	49
Total length (>= 0 bp)	16875454
Total length (>= 1000 bp)	16872548
Total length (>= 5000 bp)	16661328
Total length (>= 10000 bp)	14825976
Total length (>= 25000 bp)	8707903
Total length (>= 50000 bp)	3580332
# contigs	892
Largest contig	153037
Total length	16875454
GC (%)	54.01
N50	25635
N90	9237
auN	34907.6
L50	193
L90	630
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	38 + 7 part

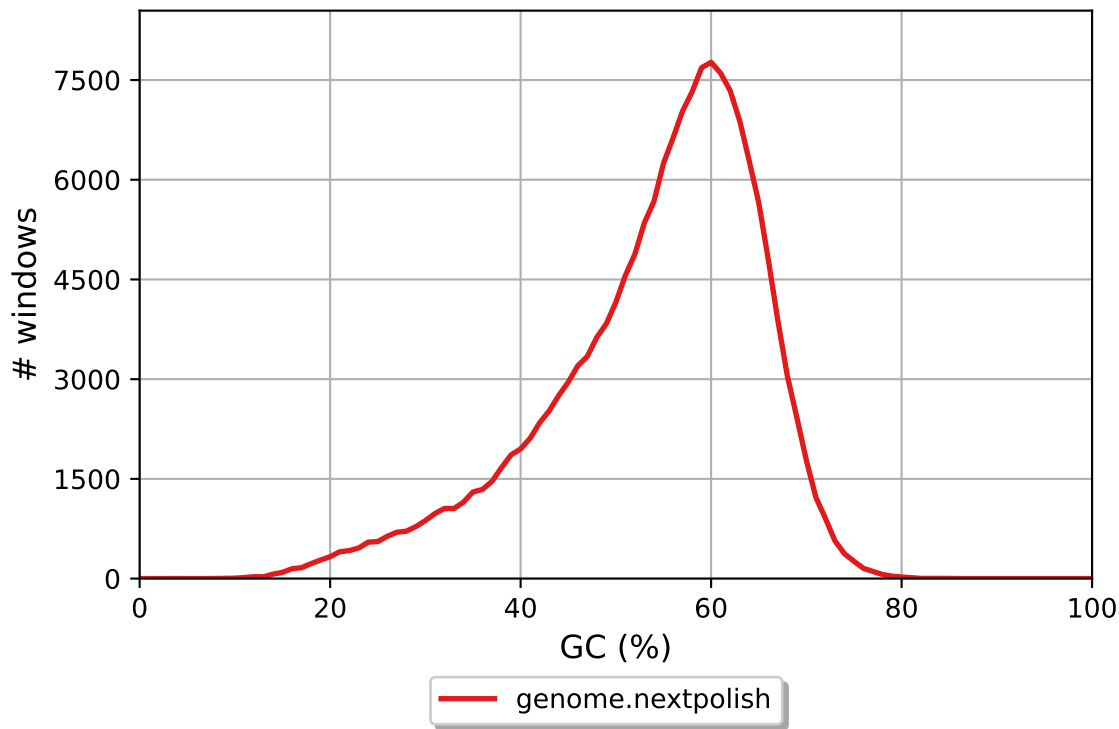
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

