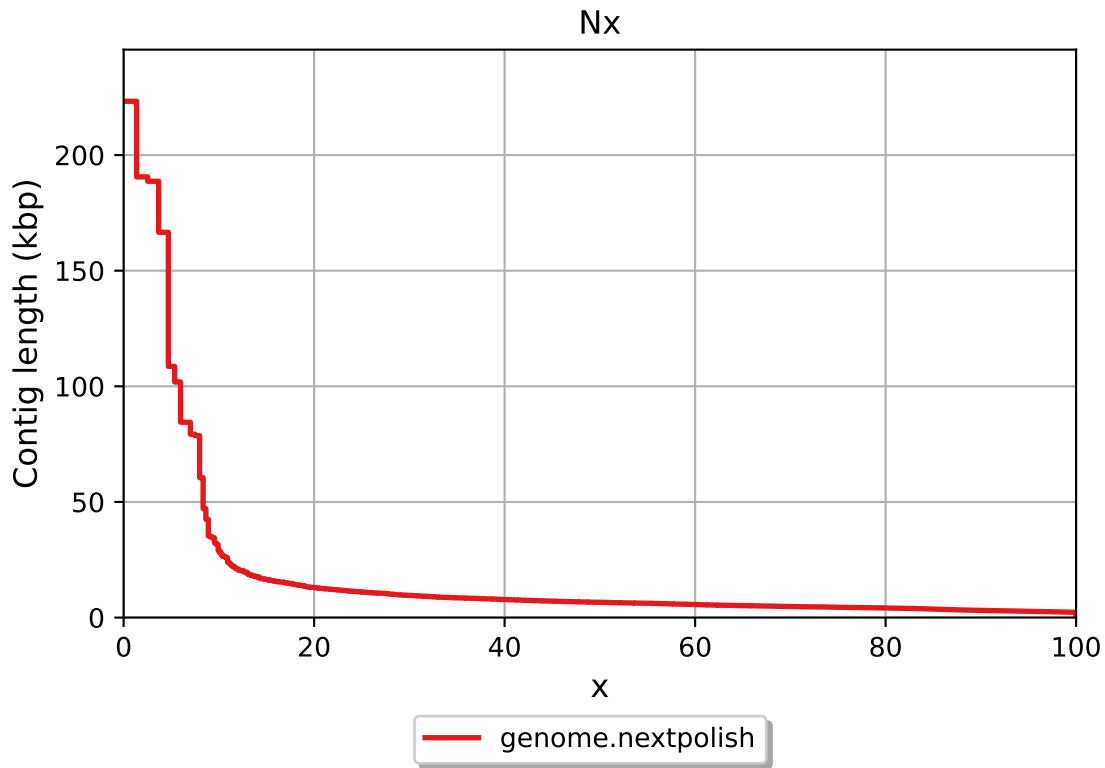


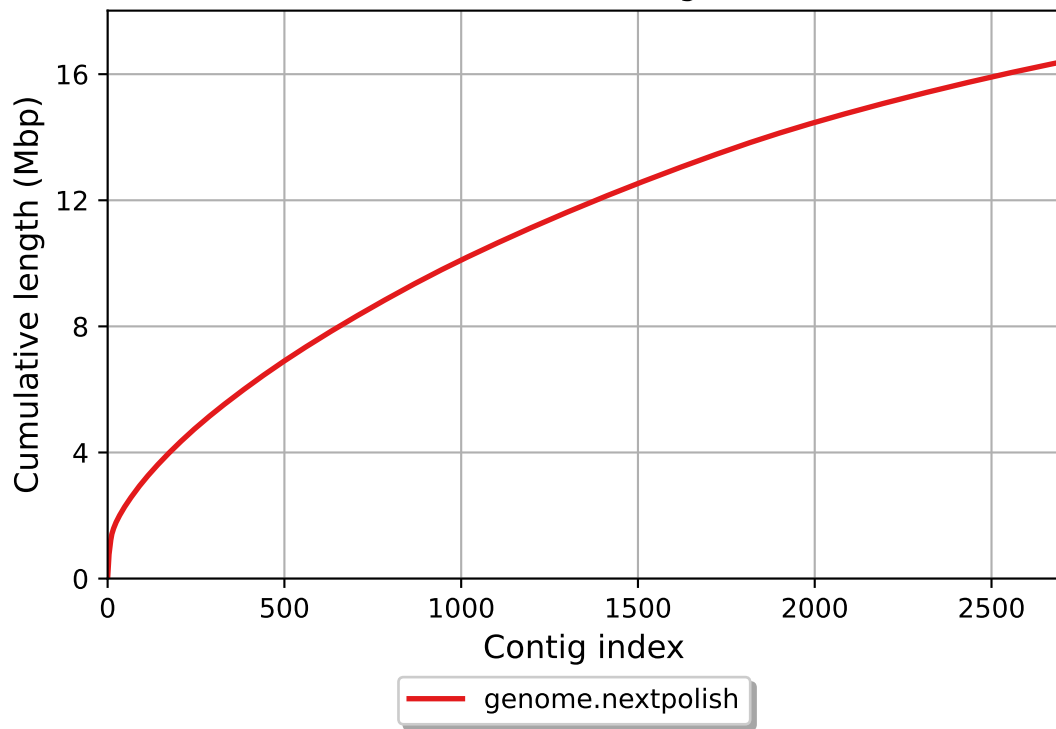
Report

	genome.nextpolish
# contigs (>= 0 bp)	2694
# contigs (>= 1000 bp)	2694
# contigs (>= 5000 bp)	1175
# contigs (>= 10000 bp)	237
# contigs (>= 25000 bp)	24
# contigs (>= 50000 bp)	11
Total length (>= 0 bp)	16377765
Total length (>= 1000 bp)	16377765
Total length (>= 5000 bp)	11015427
Total length (>= 10000 bp)	4659999
Total length (>= 25000 bp)	1787278
Total length (>= 50000 bp)	1366858
# contigs	2694
Largest contig	223253
Total length	16377765
GC (%)	38.54
N50	6605
N90	3122
auN	19637.6
L50	684
L90	2085
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	35 + 9 part

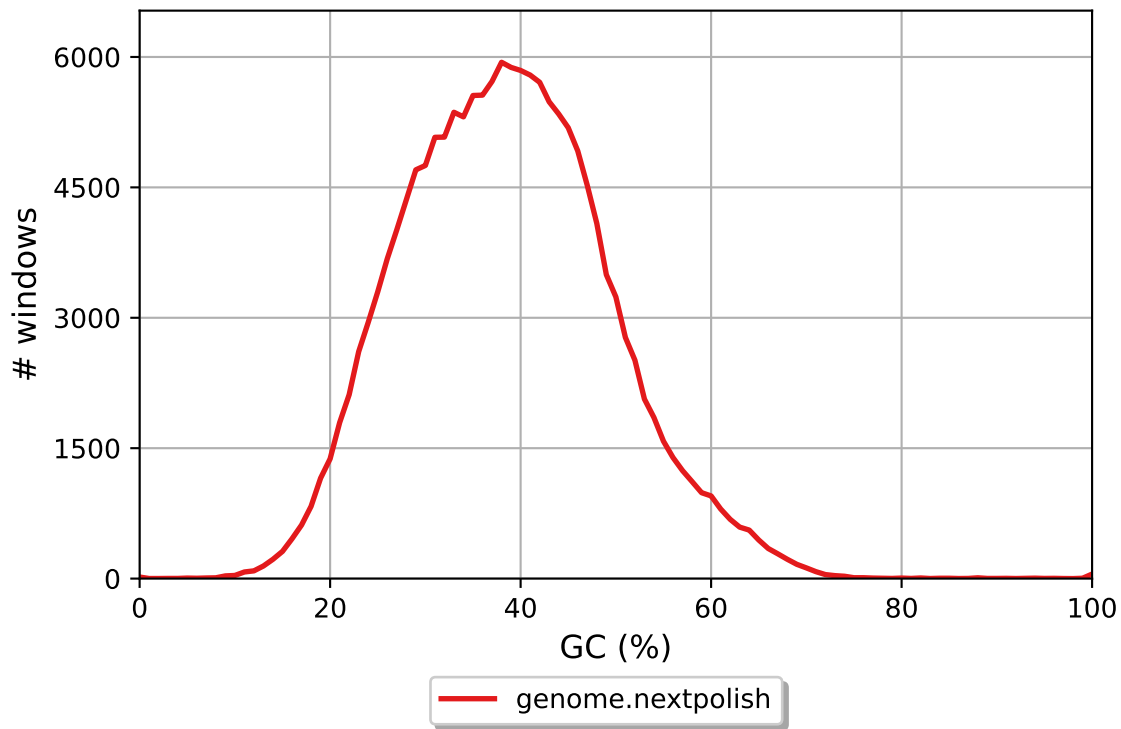
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

