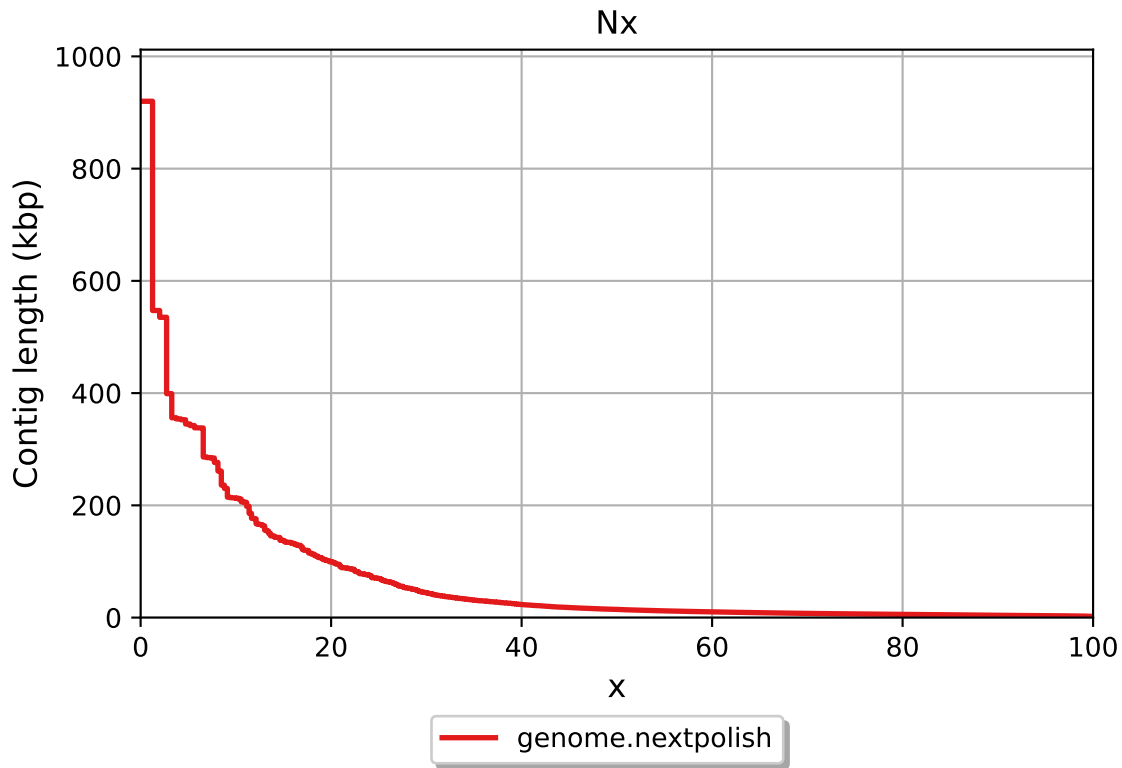


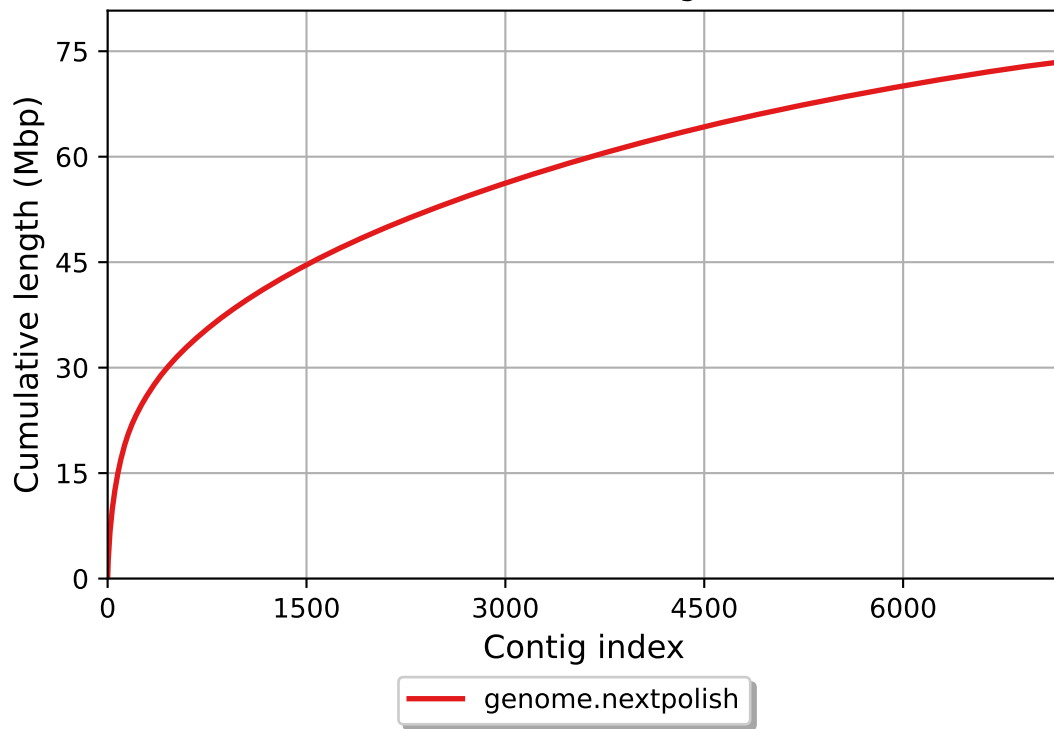
Report

	genome.nextpolish
# contigs (>= 0 bp)	7184
# contigs (>= 1000 bp)	7170
# contigs (>= 5000 bp)	4043
# contigs (>= 10000 bp)	1490
# contigs (>= 25000 bp)	392
# contigs (>= 50000 bp)	167
Total length (>= 0 bp)	73429689
Total length (>= 1000 bp)	73420818
Total length (>= 5000 bp)	62072681
Total length (>= 10000 bp)	44523030
Total length (>= 25000 bp)	28661587
Total length (>= 50000 bp)	21168049
# contigs	7184
Largest contig	920031
Total length	73429689
GC (%)	41.18
N50	14600
N90	4111
auN	71872.8
L50	832
L90	4923
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	151 + 23 part

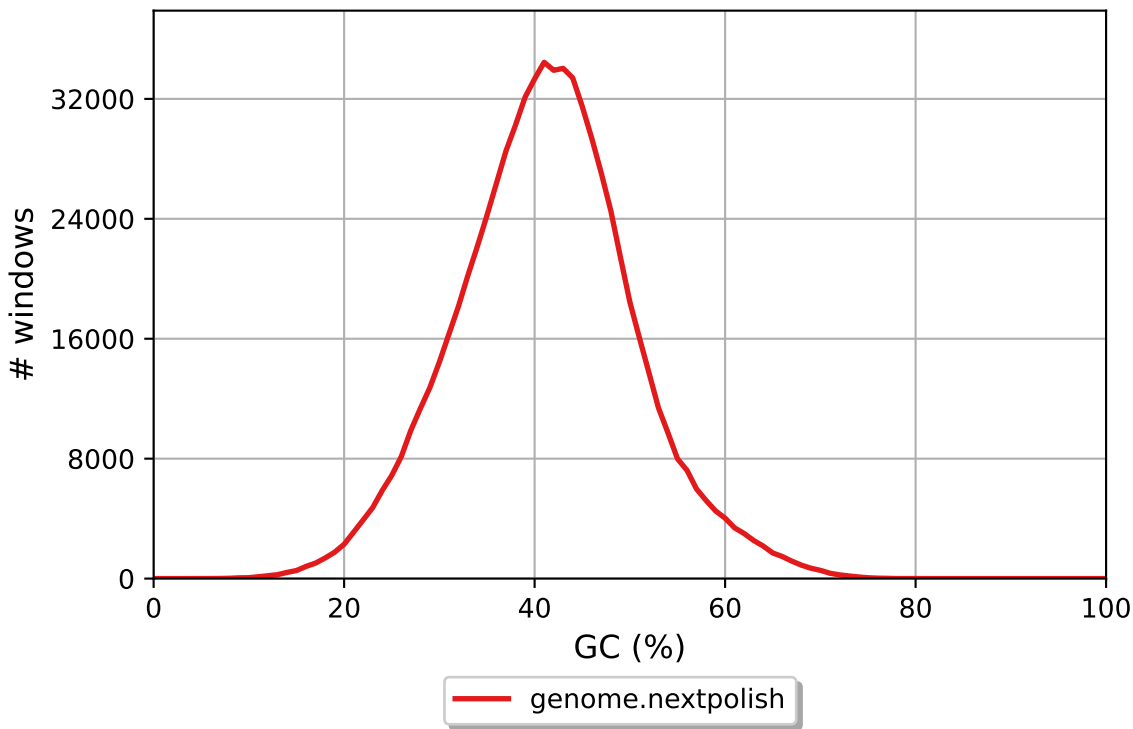
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

