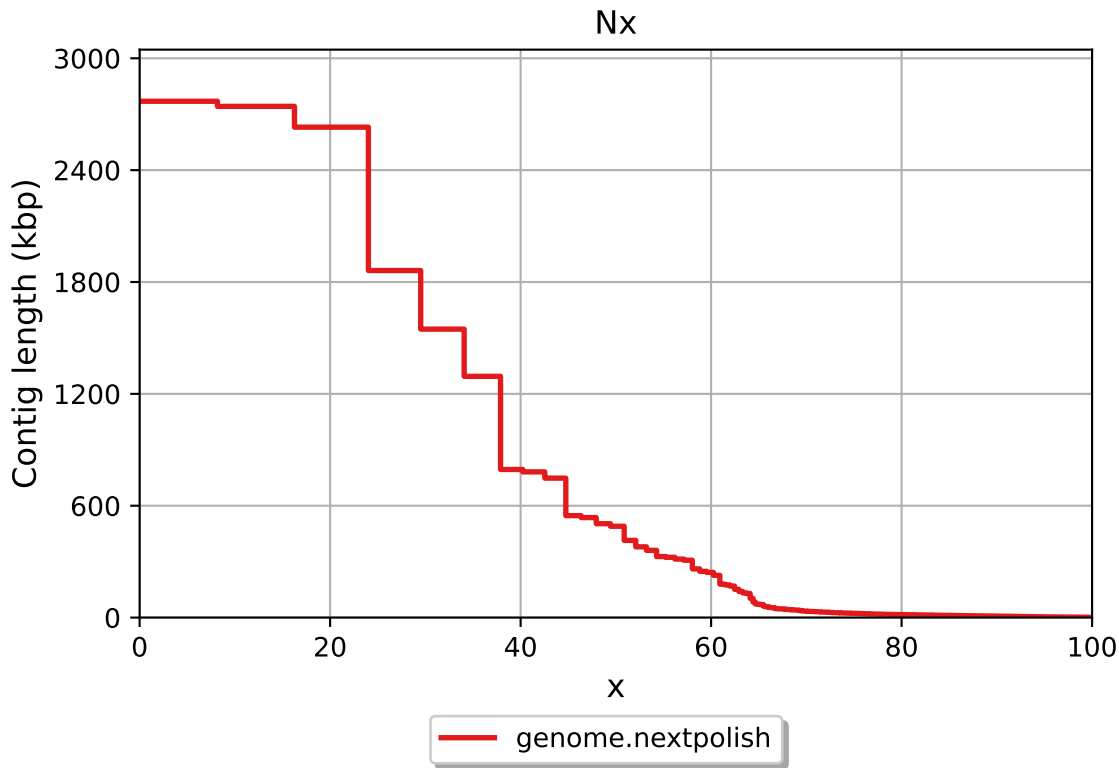


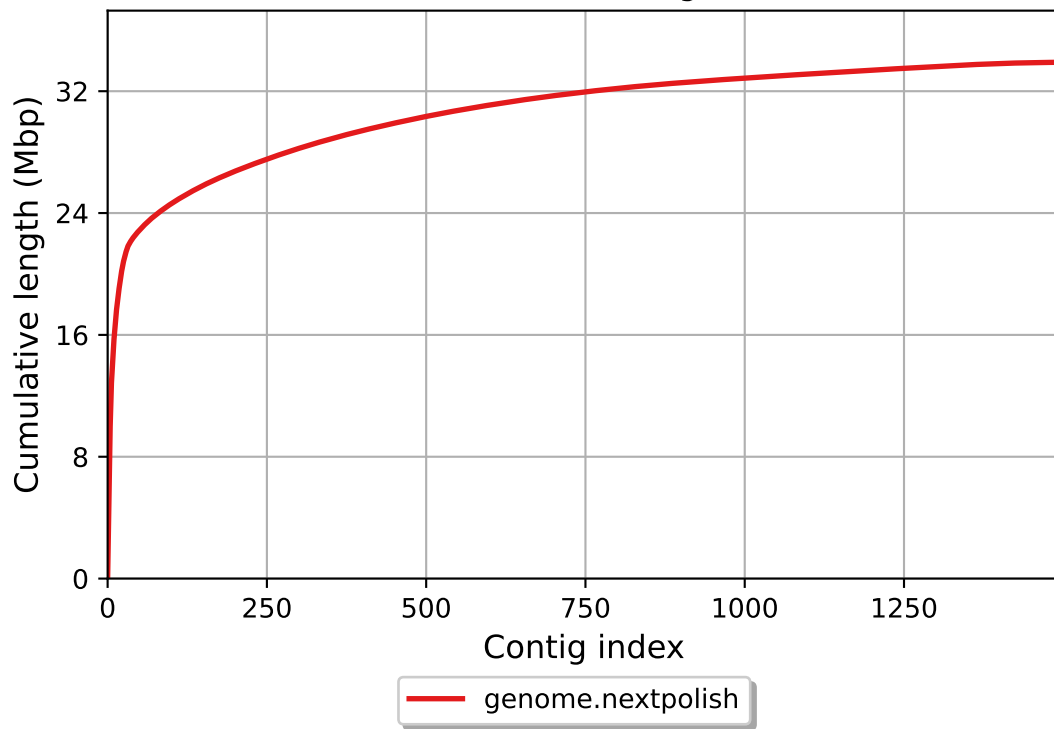
Report

	genome.nextpolish
# contigs (>= 0 bp)	1496
# contigs (>= 1000 bp)	1435
# contigs (>= 5000 bp)	732
# contigs (>= 10000 bp)	413
# contigs (>= 25000 bp)	114
# contigs (>= 50000 bp)	44
Total length (>= 0 bp)	33900284
Total length (>= 1000 bp)	33860313
Total length (>= 5000 bp)	31868267
Total length (>= 10000 bp)	29546074
Total length (>= 25000 bp)	24985375
Total length (>= 50000 bp)	22593209
# contigs	1495
Largest contig	2769437
Total length	33899788
GC (%)	40.92
N50	489803
N90	7974
auN	1003740.2
L50	13
L90	521
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	90 + 17 part

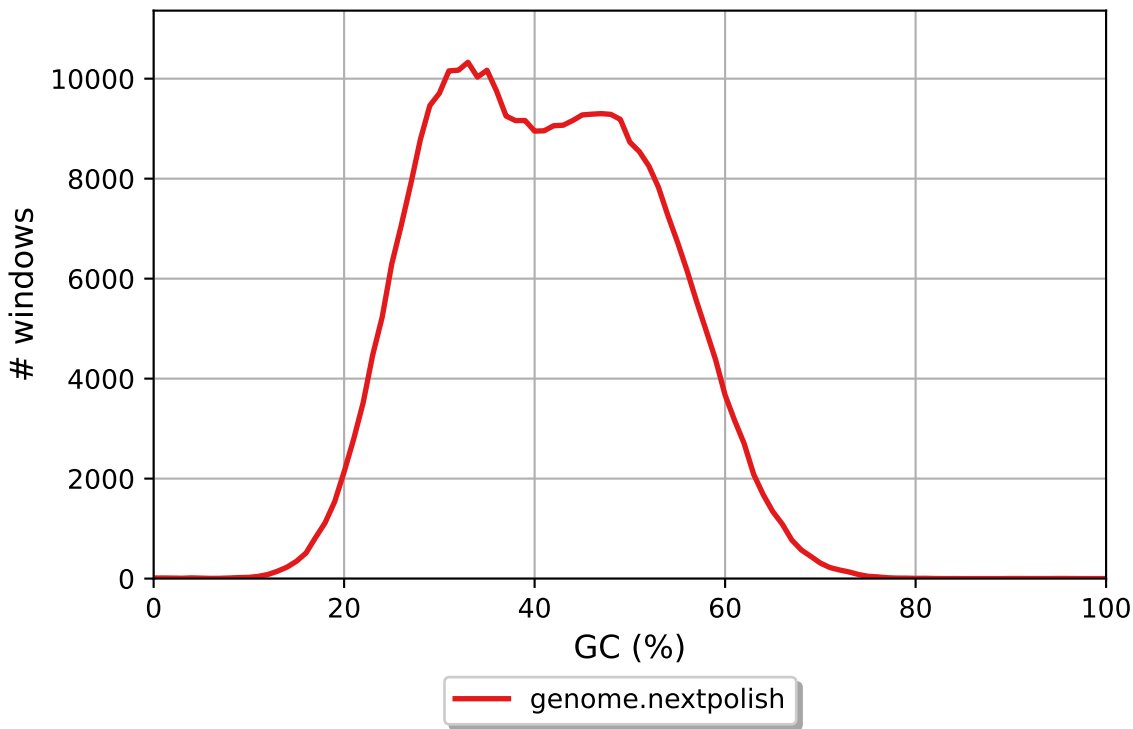
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

