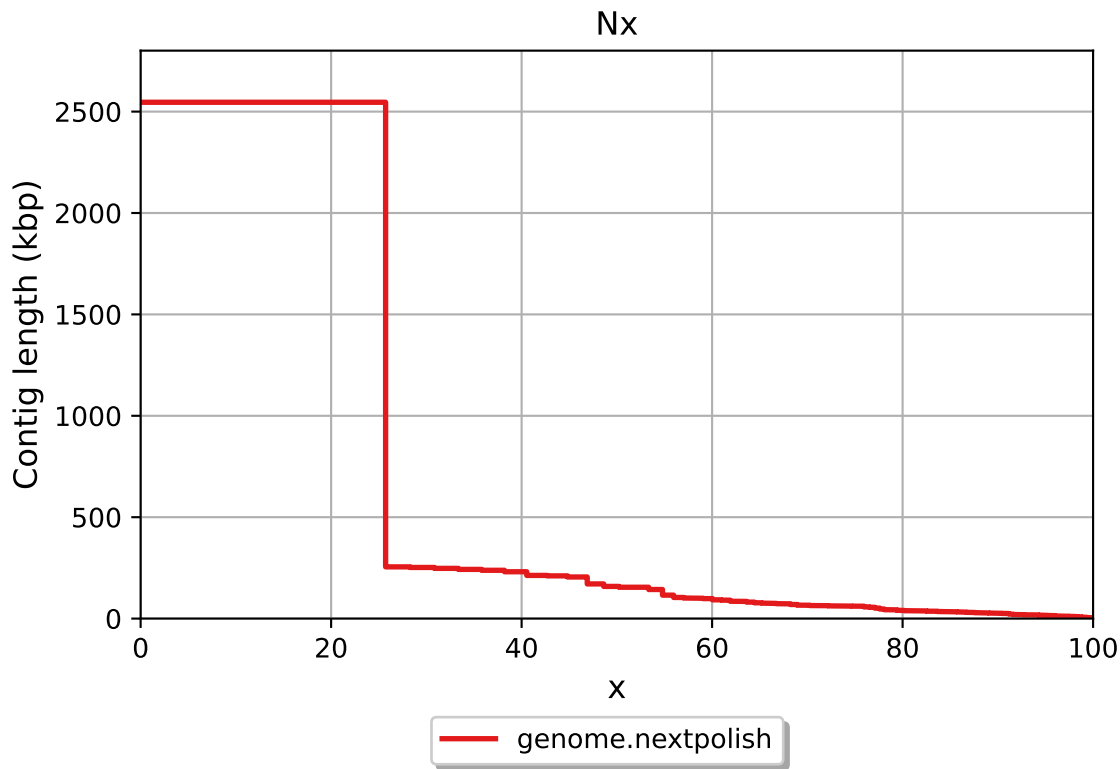


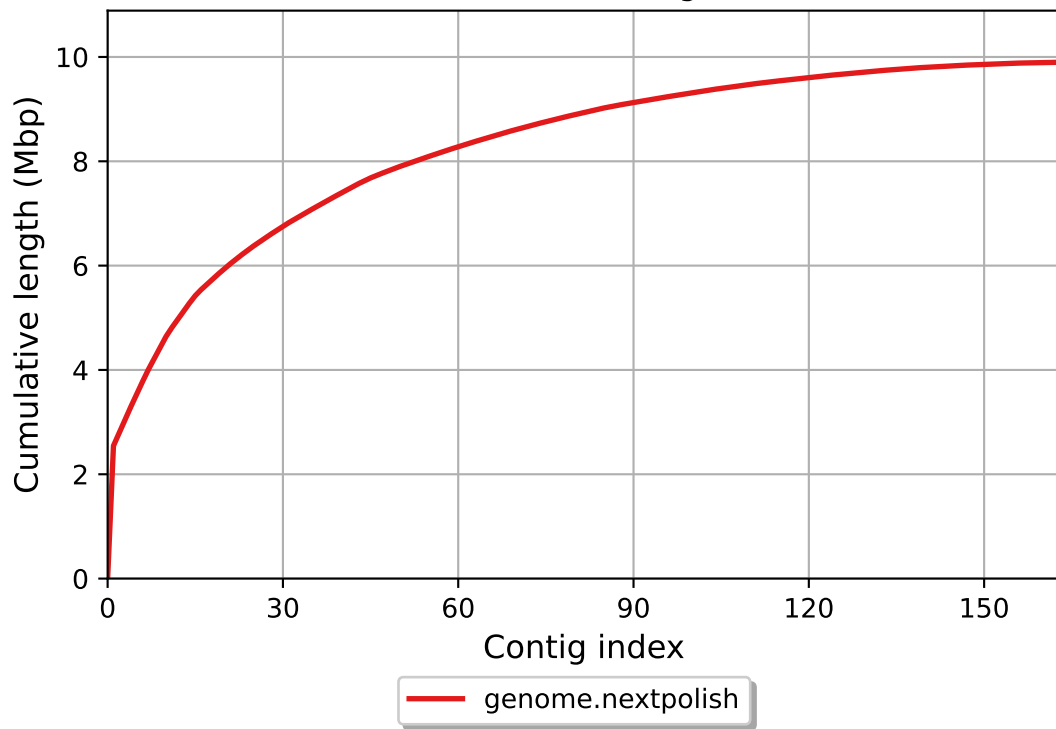
# Report

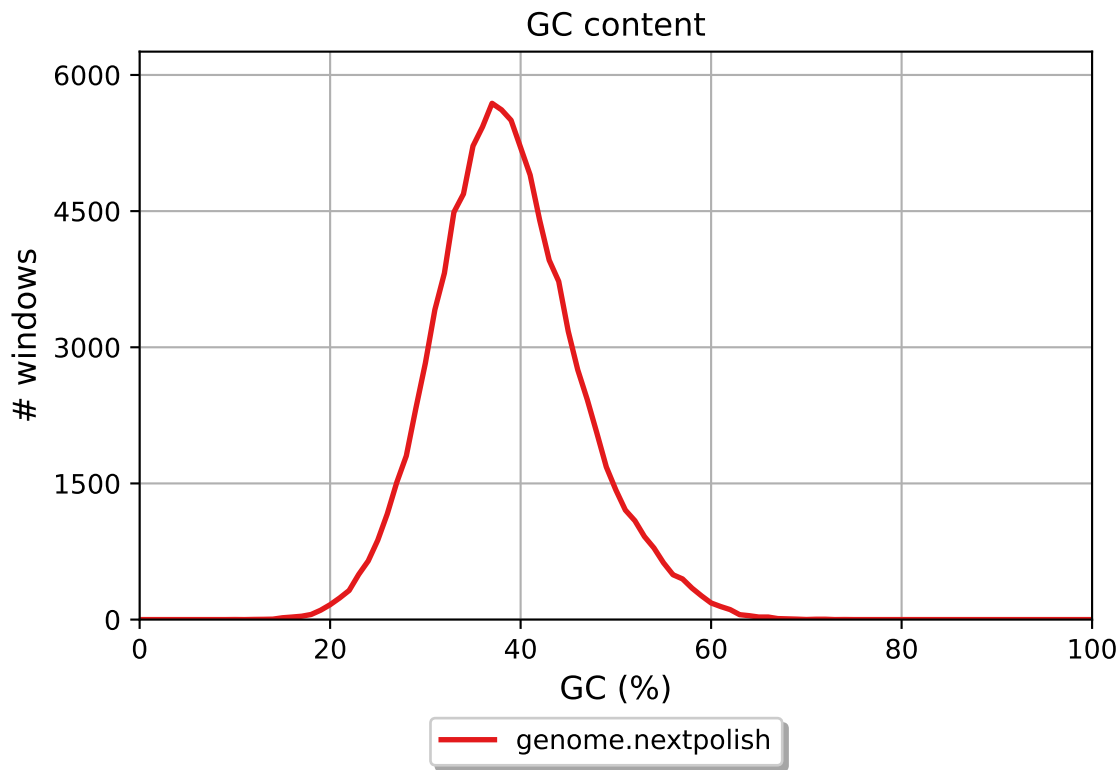
	genome.nextpolish
# contigs (>= 0 bp)	163
# contigs (>= 1000 bp)	163
# contigs (>= 5000 bp)	147
# contigs (>= 10000 bp)	131
# contigs (>= 25000 bp)	84
# contigs (>= 50000 bp)	45
Total length (>= 0 bp)	9898370
Total length (>= 1000 bp)	9898370
Total length (>= 5000 bp)	9843897
Total length (>= 10000 bp)	9725026
Total length (>= 25000 bp)	8998624
Total length (>= 50000 bp)	7681461
# contigs	163
Largest contig	2546040
Total length	9898370
GC (%)	38.69
N50	158663
N90	26098
auN	740778.2
L50	12
L90	81
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	38 + 0 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





genome.nextpolish GC content

