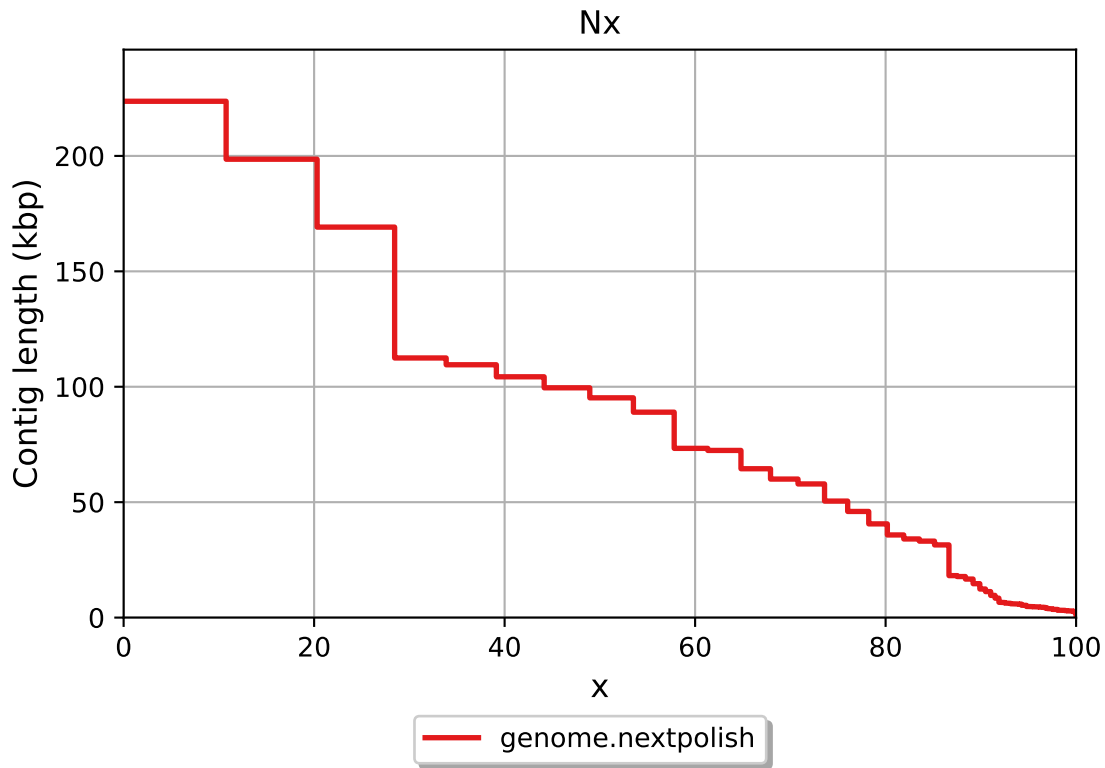
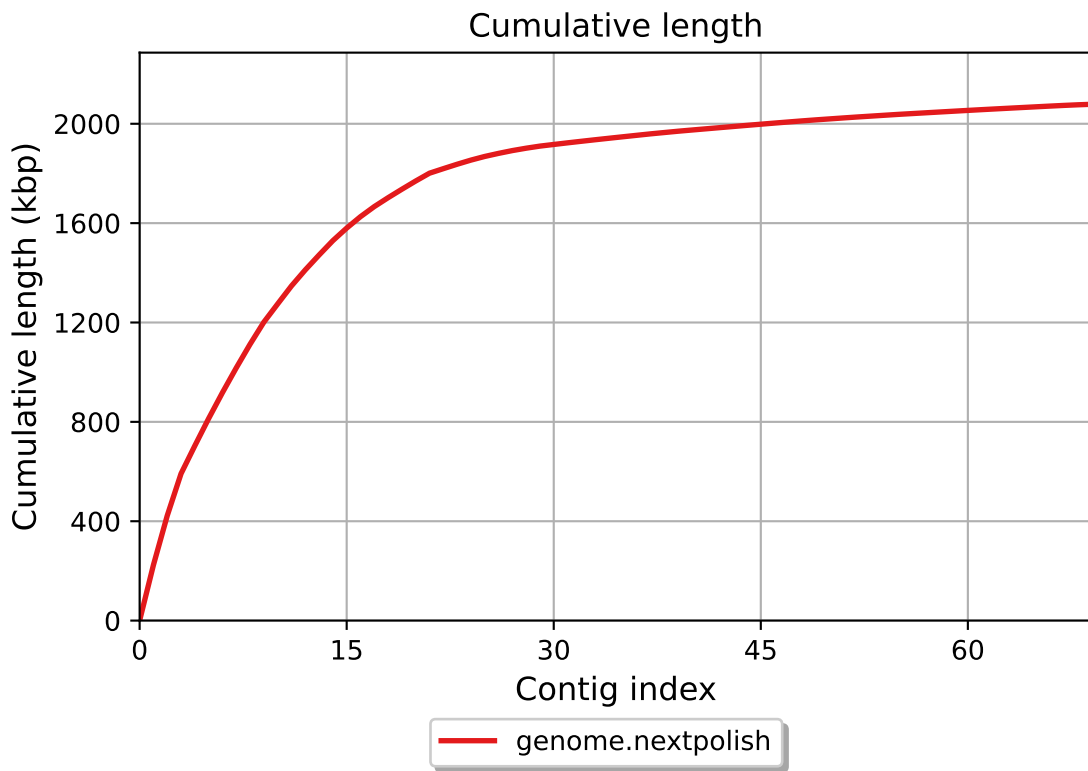


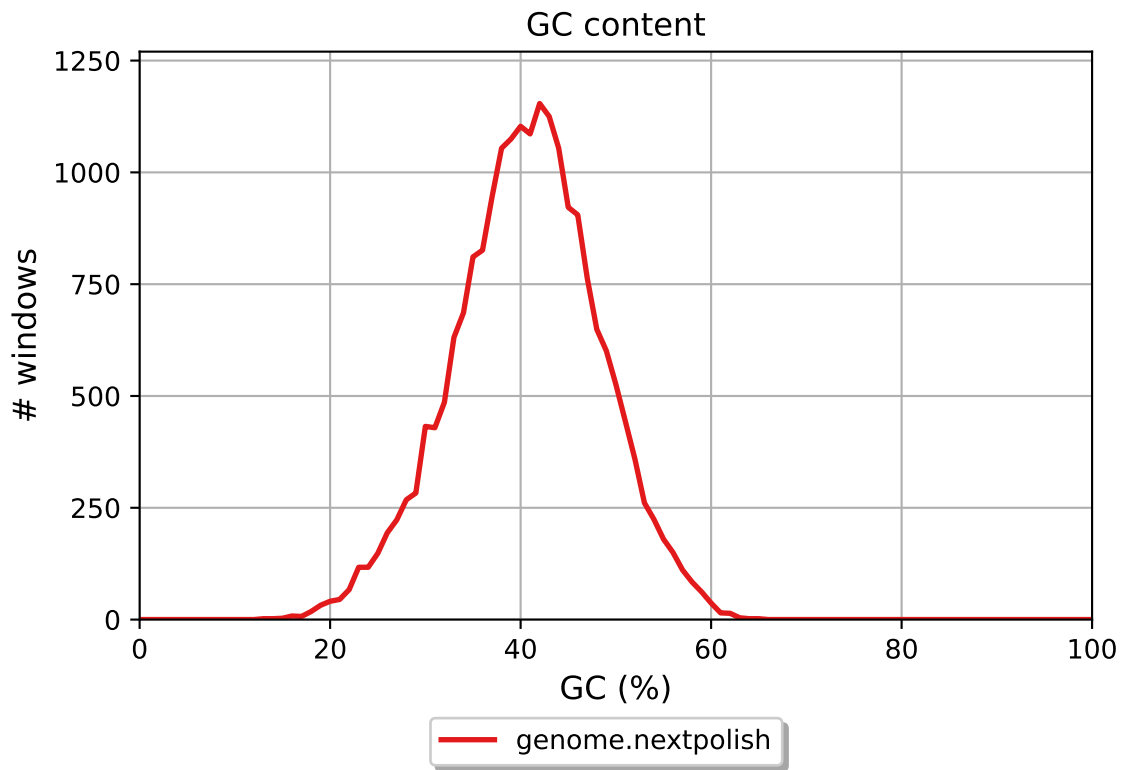
# Report

	genome.nextpolish
# contigs (>= 0 bp)	69
# contigs (>= 1000 bp)	69
# contigs (>= 5000 bp)	39
# contigs (>= 10000 bp)	27
# contigs (>= 25000 bp)	21
# contigs (>= 50000 bp)	15
Total length (>= 0 bp)	2078513
Total length (>= 1000 bp)	2078513
Total length (>= 5000 bp)	1969989
Total length (>= 10000 bp)	1892061
Total length (>= 25000 bp)	1801079
Total length (>= 50000 bp)	1580035
# contigs	69
Largest contig	223676
Total length	2078513
GC (%)	40.61
N50	95206
N90	12379
auN	103641.5
L50	8
L90	26
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	11 + 3 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).







genome.nextpolish GC content

