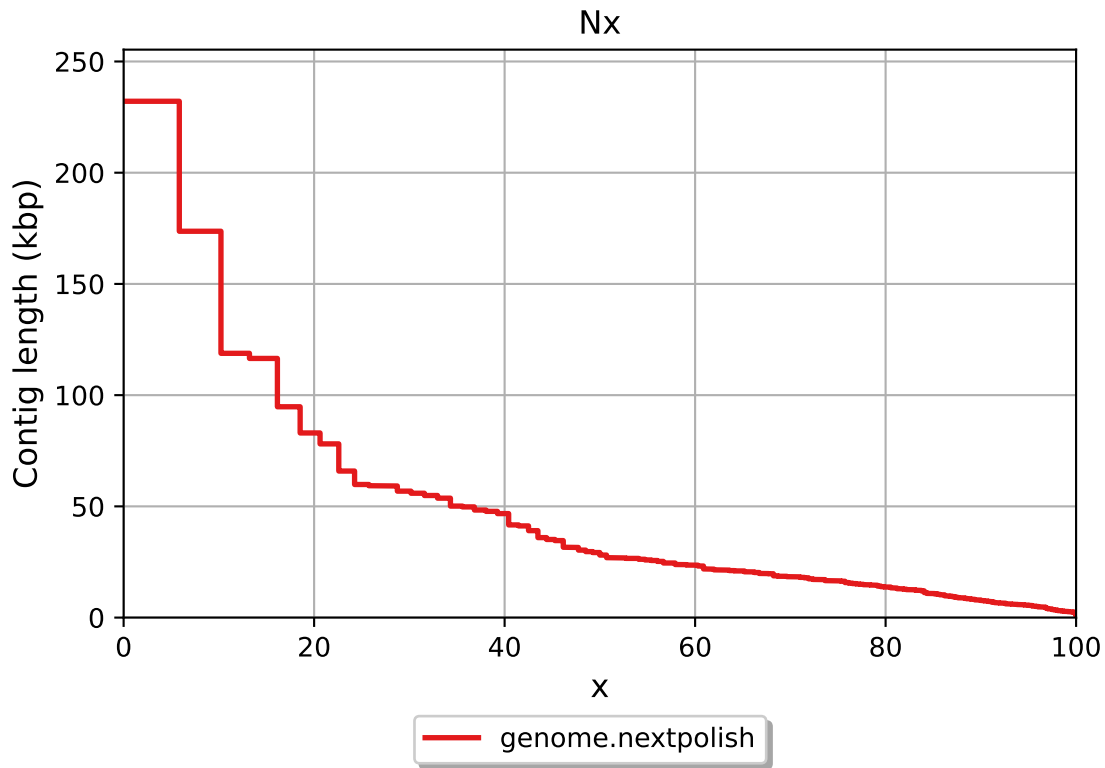
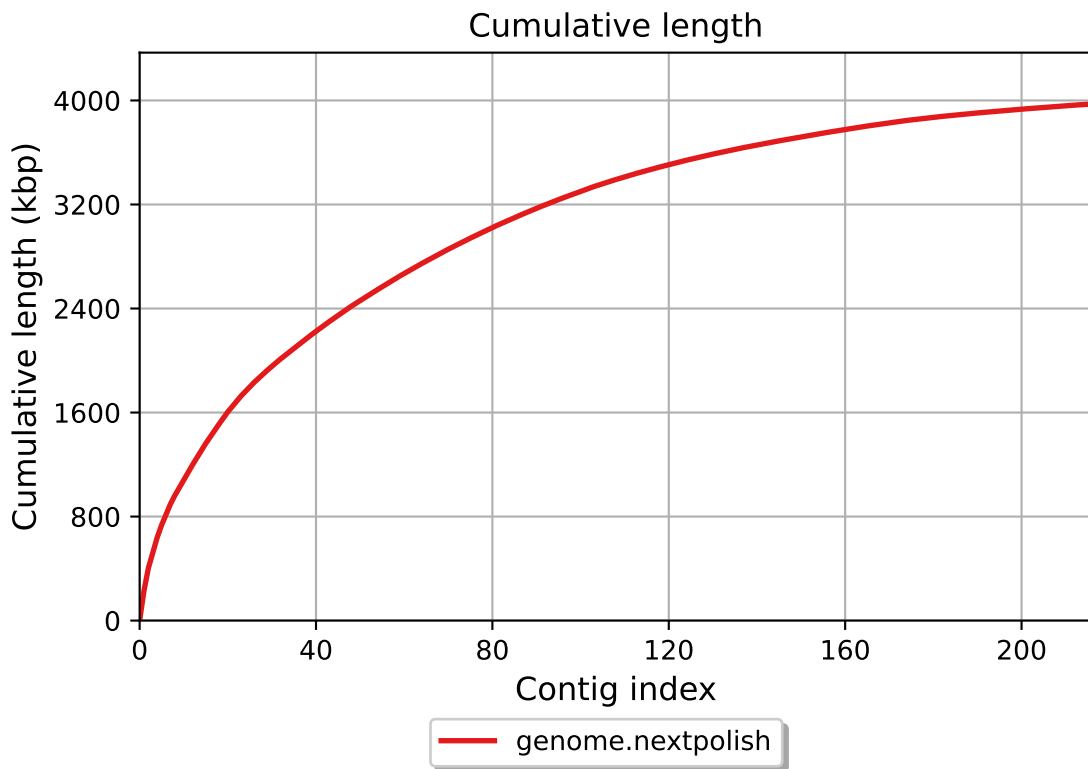


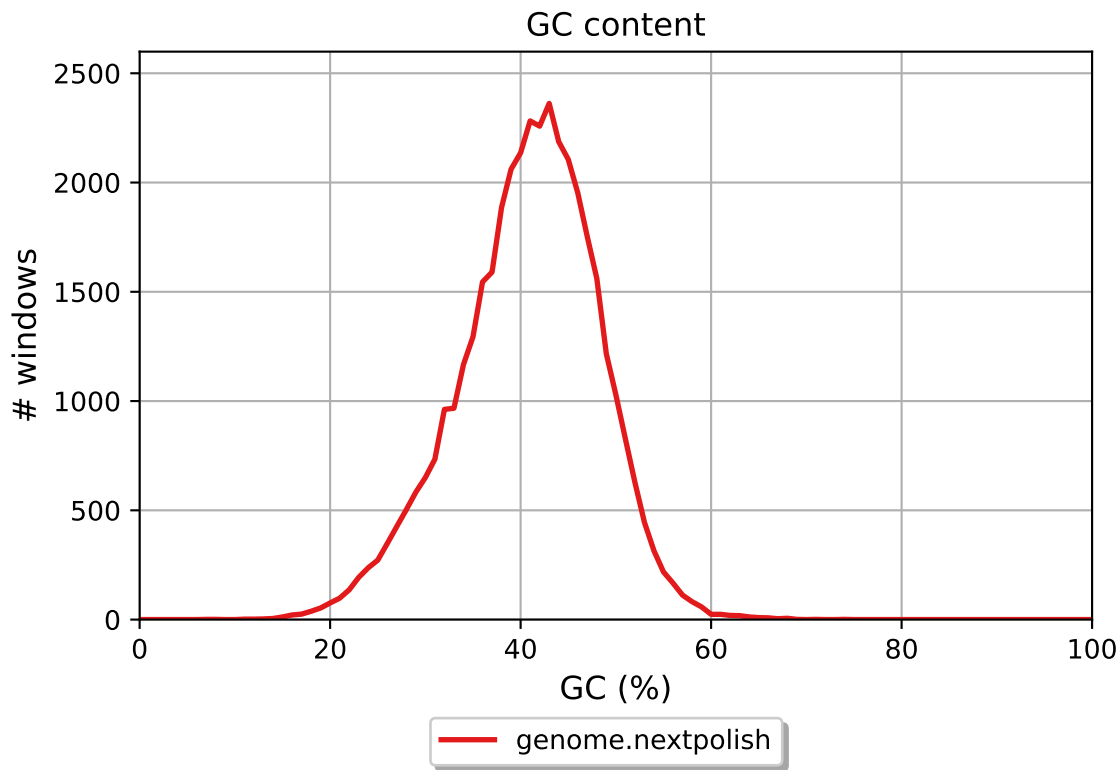
Report

	genome.nextpolish
# contigs (>= 0 bp)	216
# contigs (>= 1000 bp)	215
# contigs (>= 5000 bp)	166
# contigs (>= 10000 bp)	111
# contigs (>= 25000 bp)	41
# contigs (>= 50000 bp)	16
Total length (>= 0 bp)	3970727
Total length (>= 1000 bp)	3970216
Total length (>= 5000 bp)	3808317
Total length (>= 10000 bp)	3422799
Total length (>= 25000 bp)	2250225
Total length (>= 50000 bp)	1412569
# contigs	216
Largest contig	232115
Total length	3970727
GC (%)	40.74
N50	28102
N90	7726
auN	54705.0
L50	32
L90	129
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 2 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

