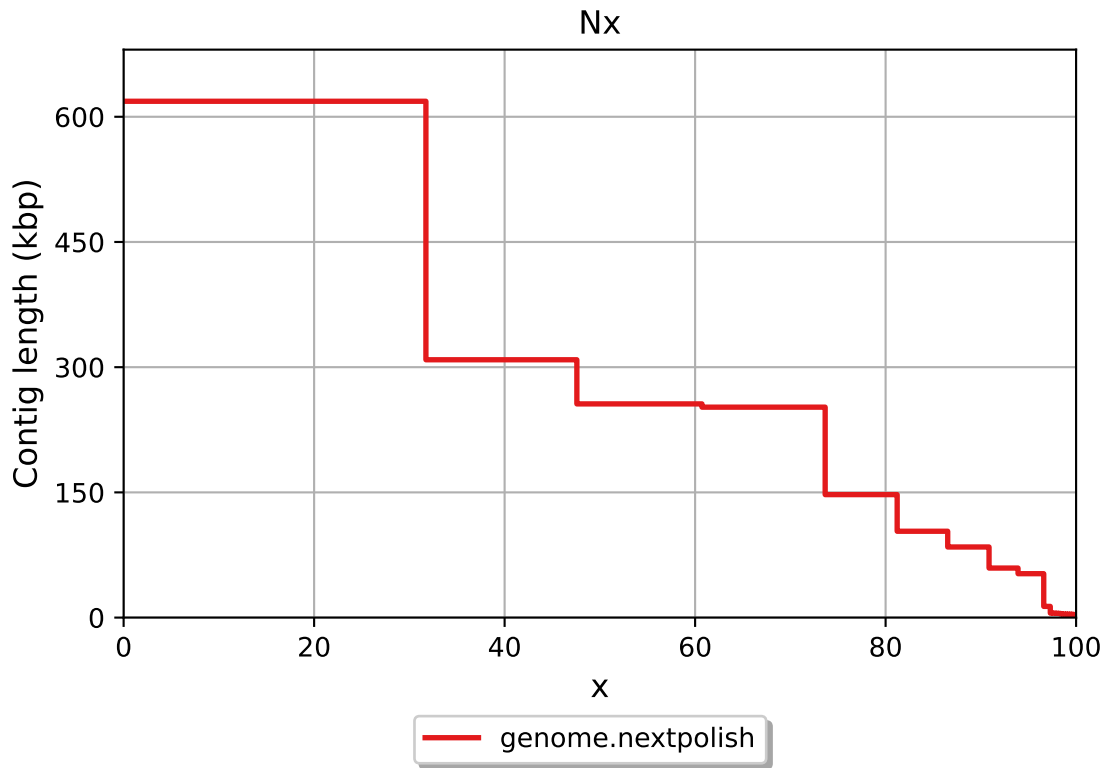
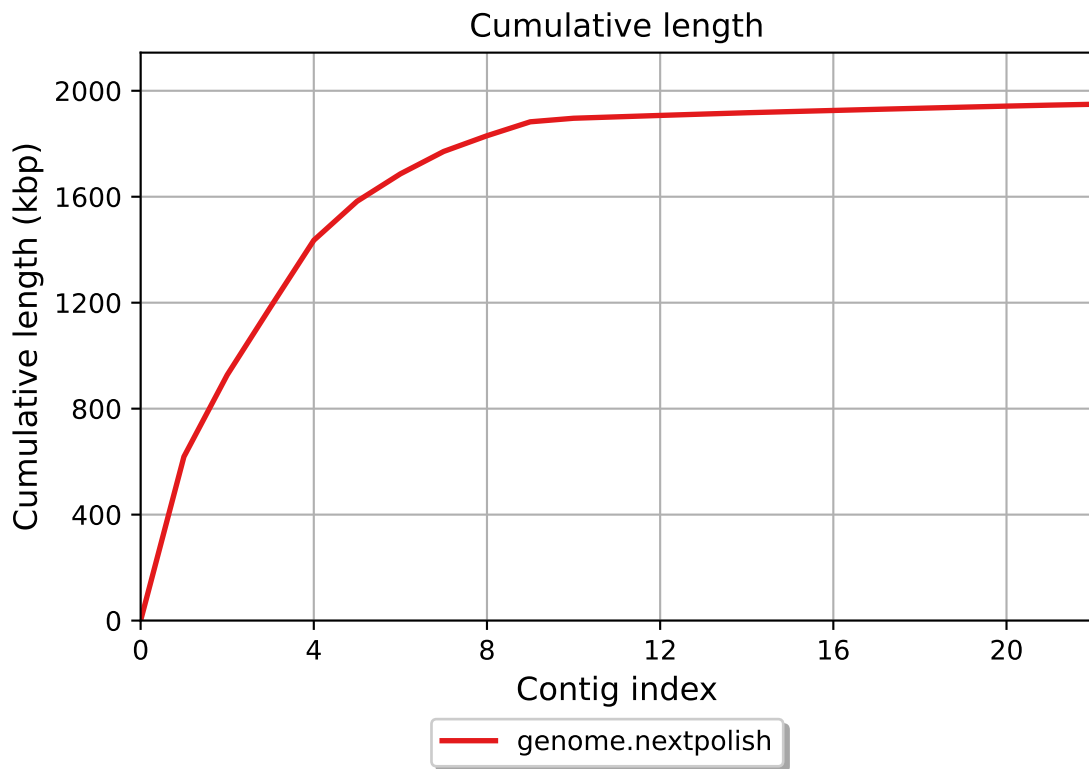


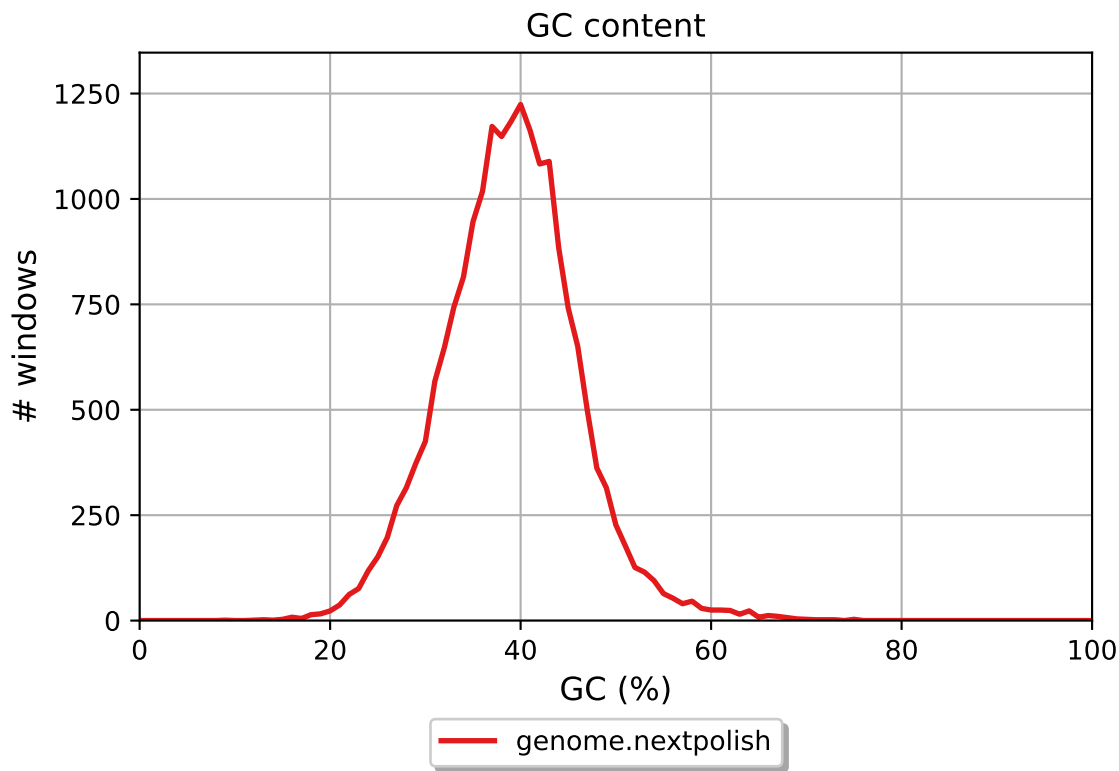
Report

	genome.nextpolish
# contigs (>= 0 bp)	22
# contigs (>= 1000 bp)	22
# contigs (>= 5000 bp)	13
# contigs (>= 10000 bp)	10
# contigs (>= 25000 bp)	9
# contigs (>= 50000 bp)	9
Total length (>= 0 bp)	1949044
Total length (>= 1000 bp)	1949044
Total length (>= 5000 bp)	1912056
Total length (>= 10000 bp)	1896242
Total length (>= 25000 bp)	1882833
Total length (>= 50000 bp)	1882833
# contigs	22
Largest contig	618571
Total length	1949044
GC (%)	38.96
N50	256008
N90	84584
auN	335239.5
L50	3
L90	7
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	18 + 6 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

