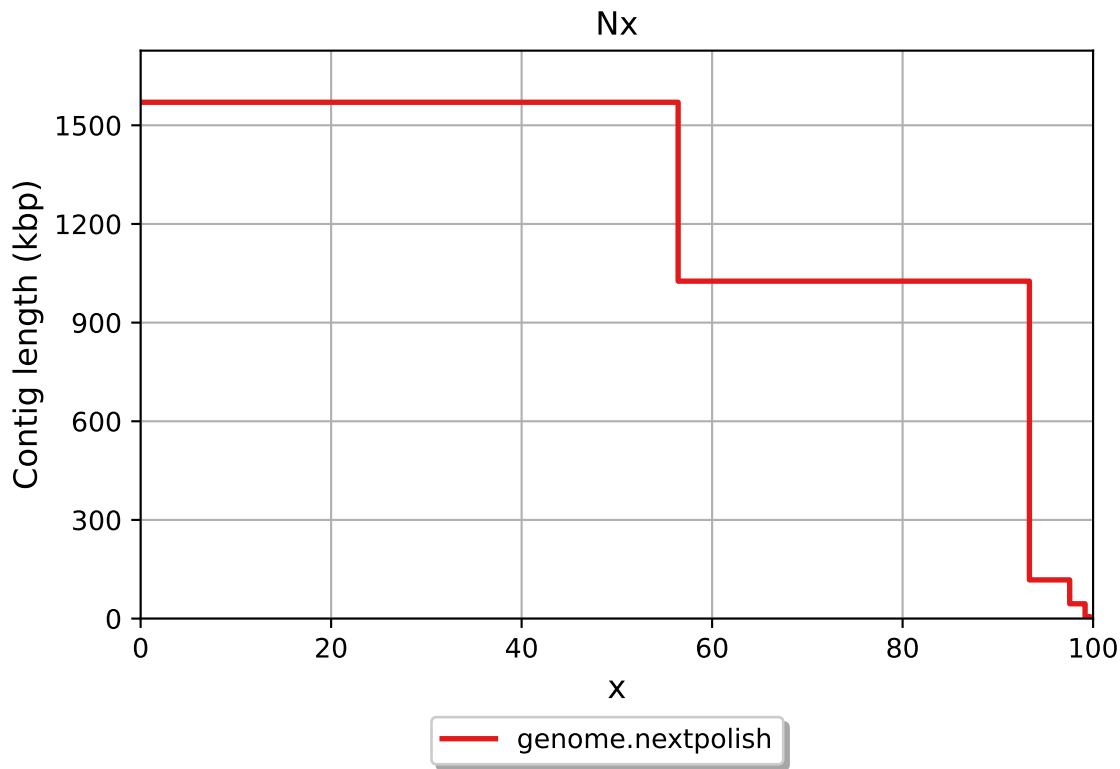
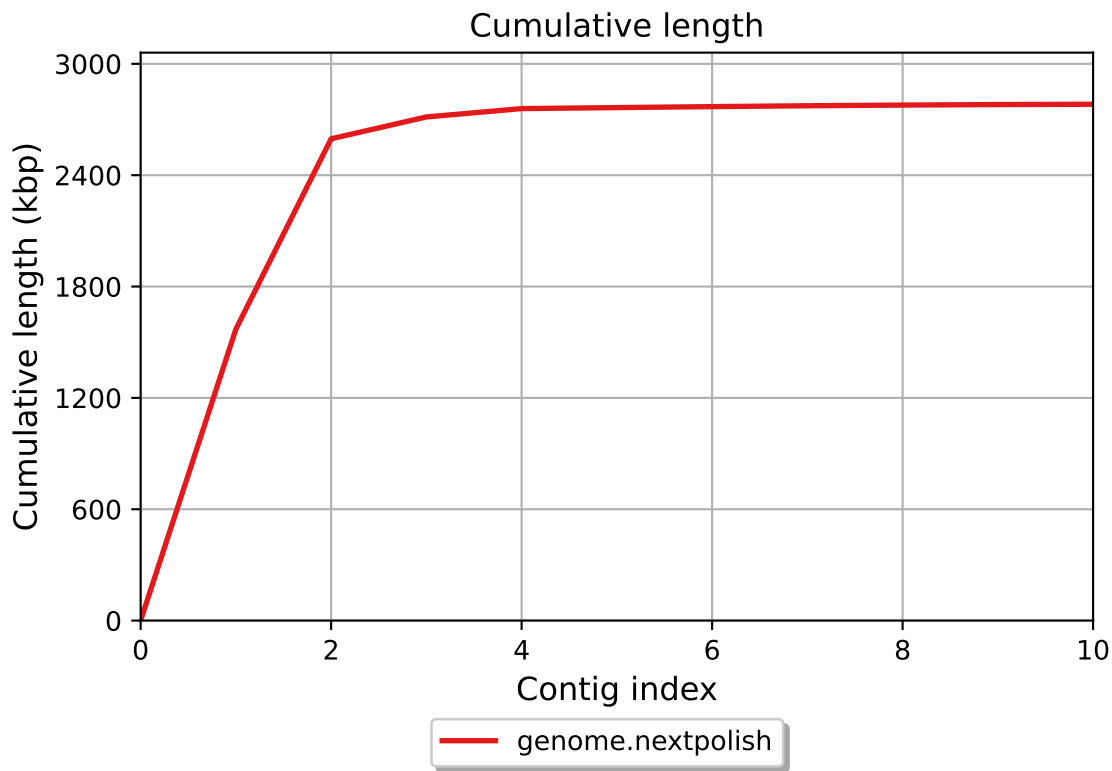


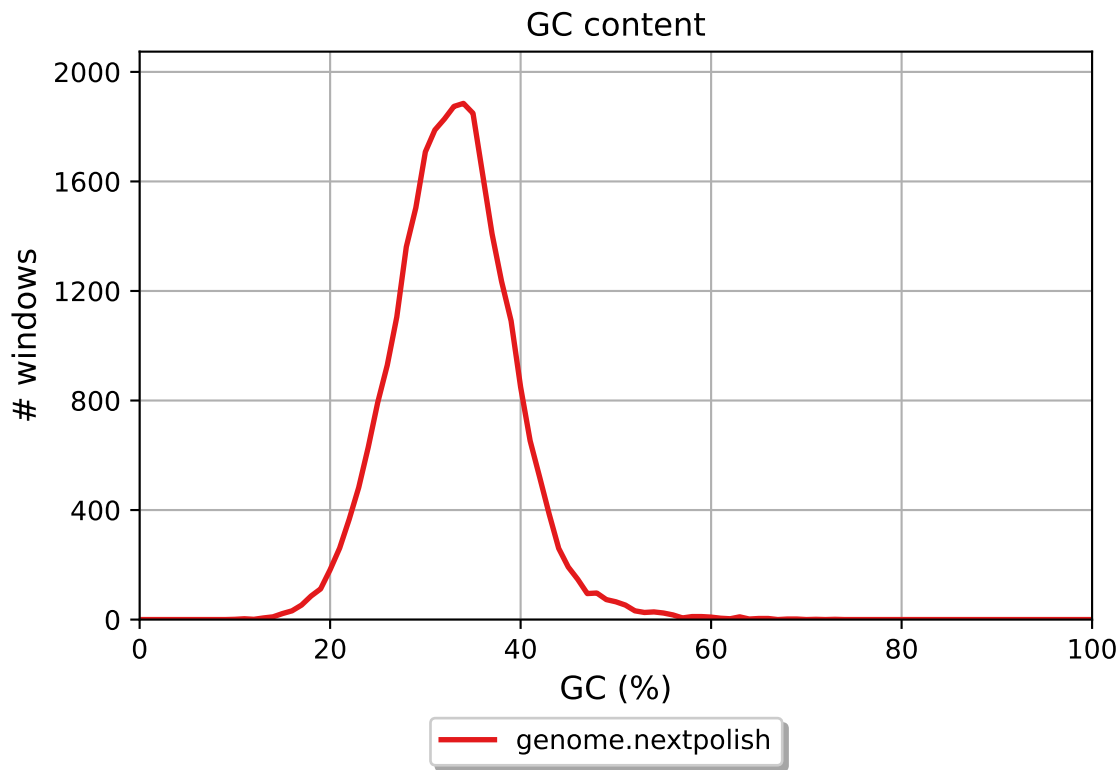
Report

	genome.nextpolish
# contigs (>= 0 bp)	10
# contigs (>= 1000 bp)	10
# contigs (>= 5000 bp)	6
# contigs (>= 10000 bp)	4
# contigs (>= 25000 bp)	4
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	2782121
Total length (>= 1000 bp)	2782121
Total length (>= 5000 bp)	2769422
Total length (>= 10000 bp)	2758770
Total length (>= 25000 bp)	2758770
Total length (>= 50000 bp)	2713750
# contigs	10
Largest contig	1570021
Total length	2782121
GC (%)	32.94
N50	1570021
N90	1026069
auN	1270166.3
L50	1
L90	2
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 9 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

