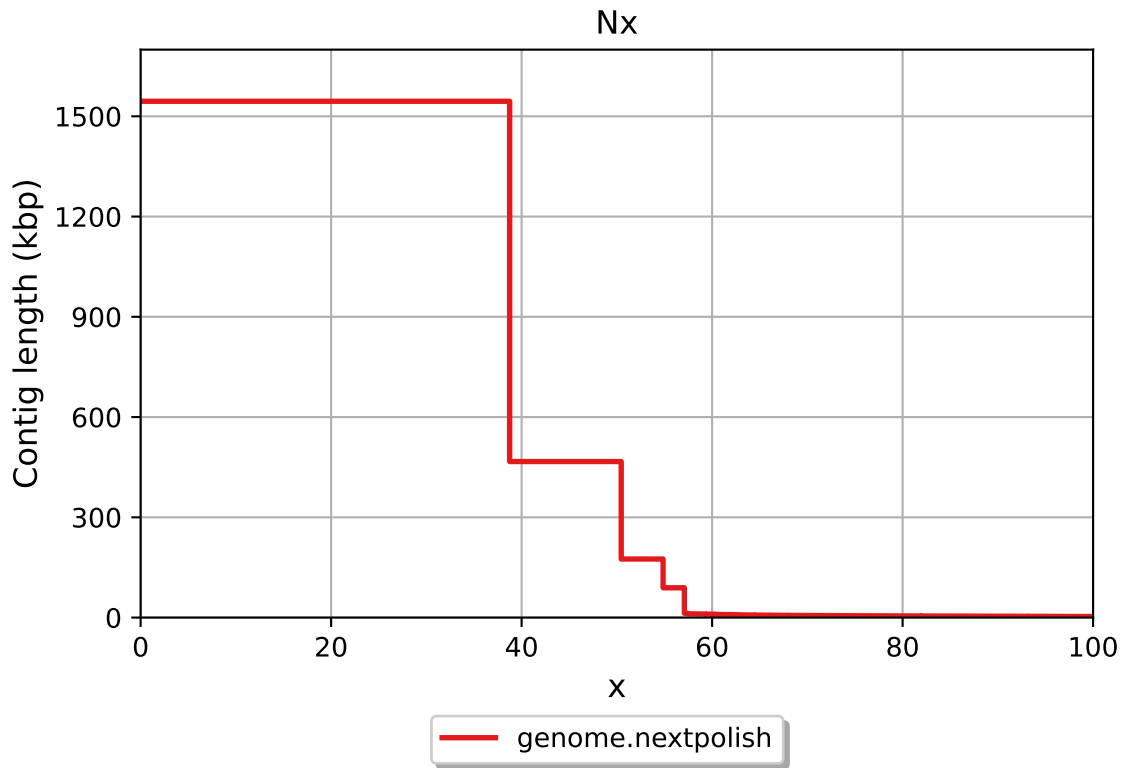
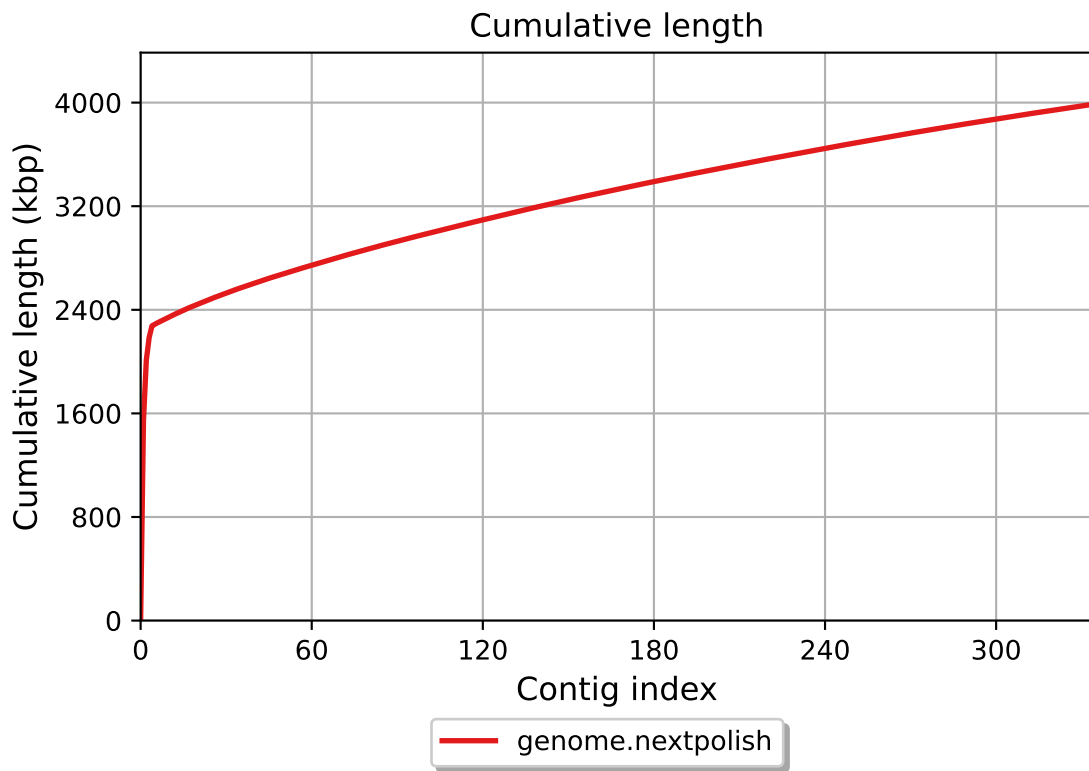


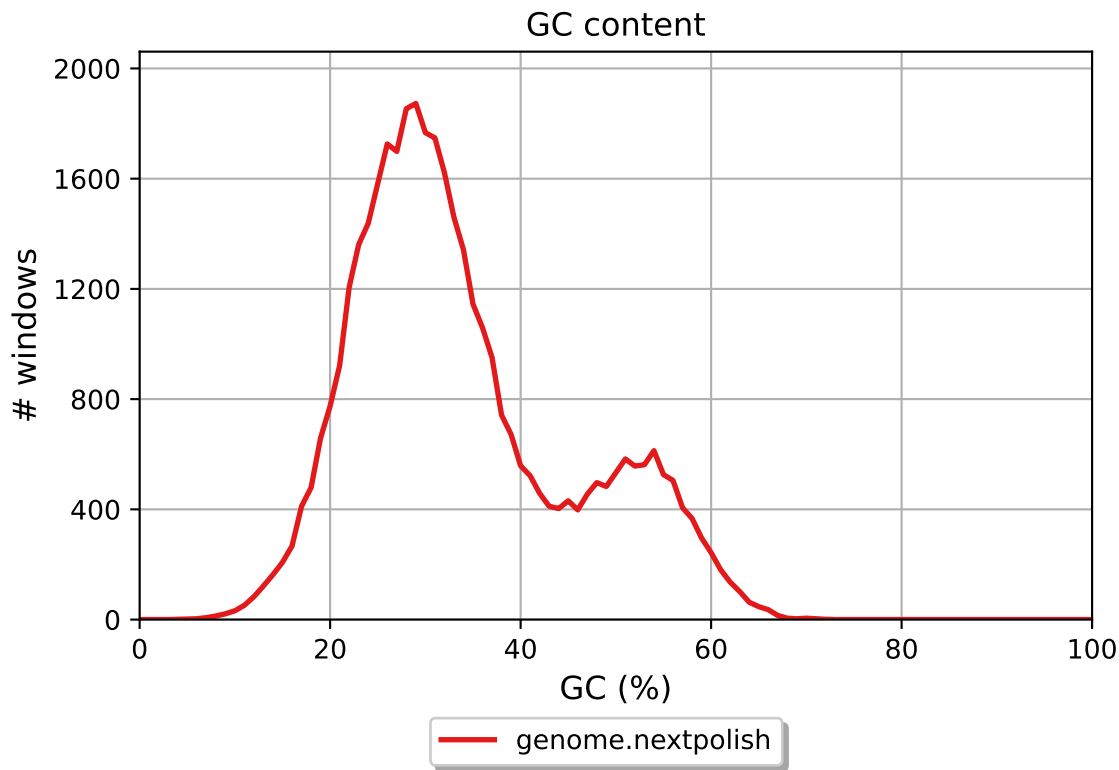
Report

	genome.nextpolish
# contigs (>= 0 bp)	334
# contigs (>= 1000 bp)	334
# contigs (>= 5000 bp)	145
# contigs (>= 10000 bp)	16
# contigs (>= 25000 bp)	4
# contigs (>= 50000 bp)	4
Total length (>= 0 bp)	3987189
Total length (>= 1000 bp)	3987189
Total length (>= 5000 bp)	3224146
Total length (>= 10000 bp)	2406502
Total length (>= 25000 bp)	2276294
Total length (>= 50000 bp)	2276294
# contigs	334
Largest contig	1544963
Total length	3987189
GC (%)	33.87
N50	466922
N90	4144
auN	665502.8
L50	2
L90	226
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	18 + 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

