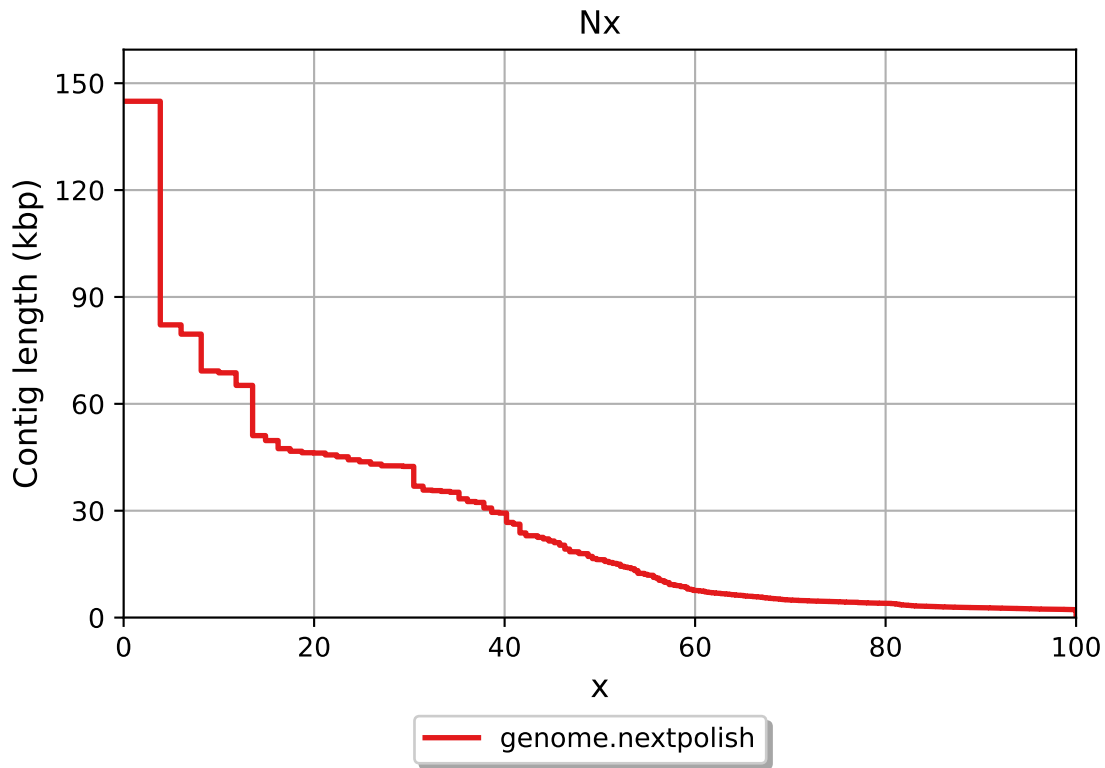
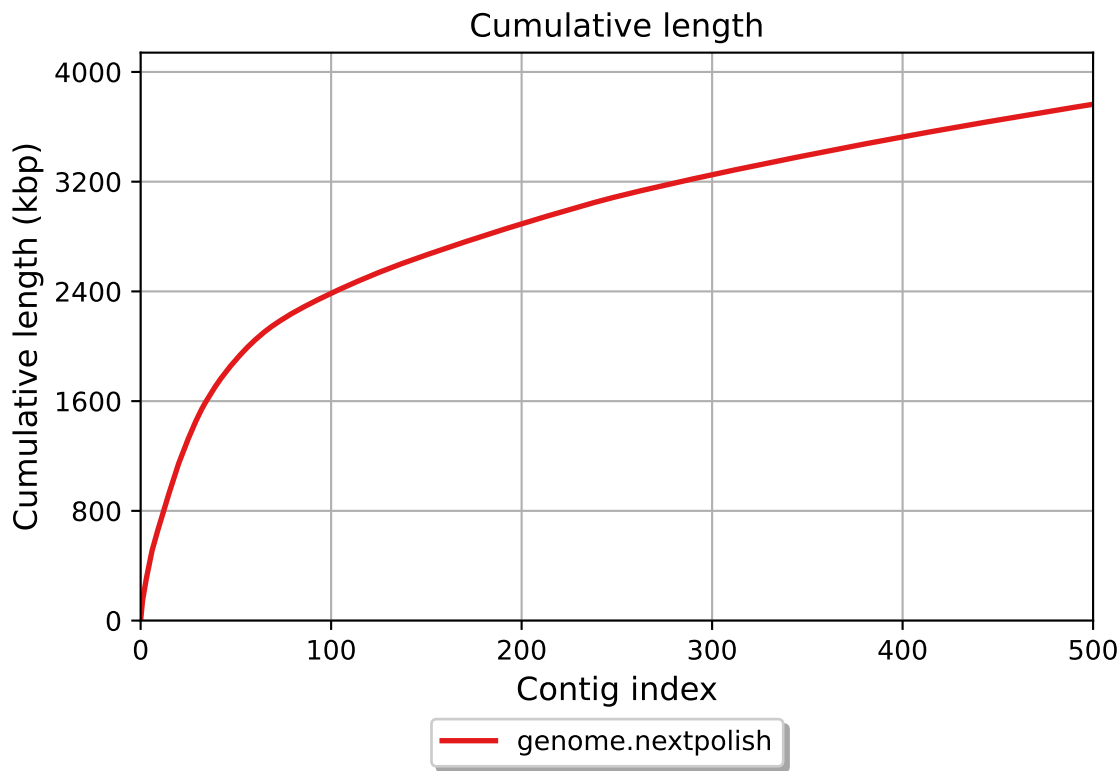


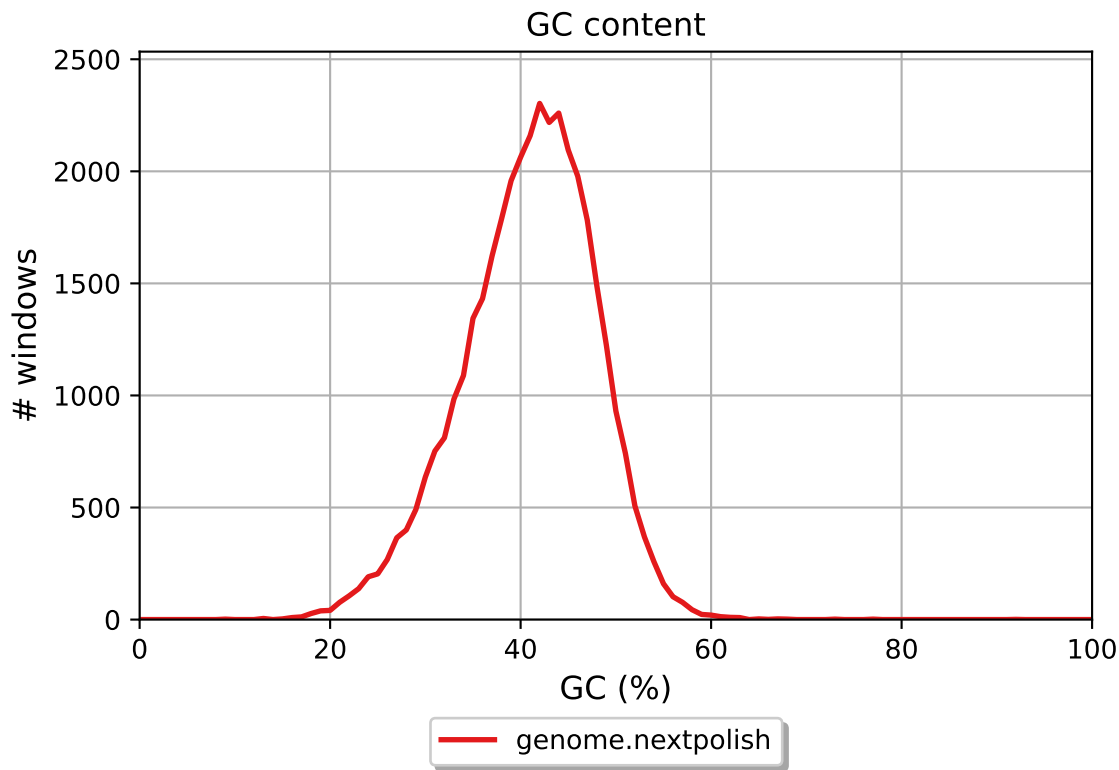
Report

	genome.nextpolish
# contigs (>= 0 bp)	500
# contigs (>= 1000 bp)	500
# contigs (>= 5000 bp)	141
# contigs (>= 10000 bp)	68
# contigs (>= 25000 bp)	33
# contigs (>= 50000 bp)	7
Total length (>= 0 bp)	3764388
Total length (>= 1000 bp)	3764388
Total length (>= 5000 bp)	2621975
Total length (>= 10000 bp)	2136583
Total length (>= 25000 bp)	1566289
Total length (>= 50000 bp)	560848
# contigs	500
Largest contig	144930
Total length	3764388
GC (%)	40.82
N50	16287
N90	2768
auN	28601.1
L50	49
L90	348
# 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

