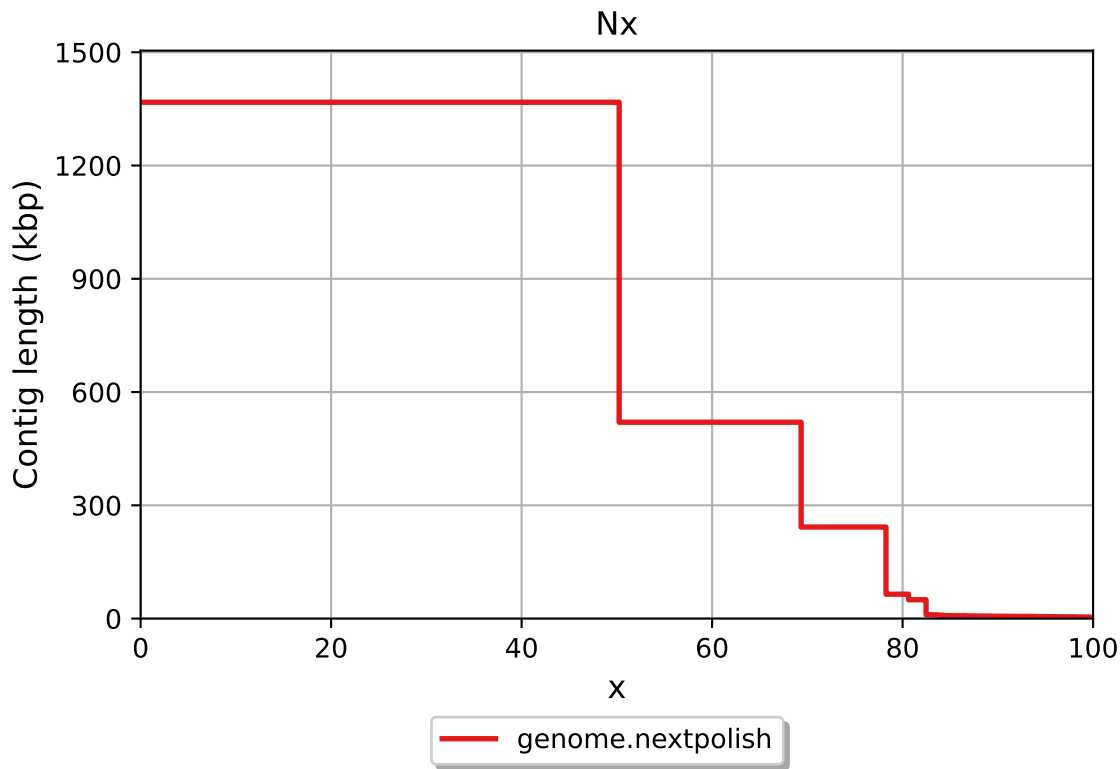
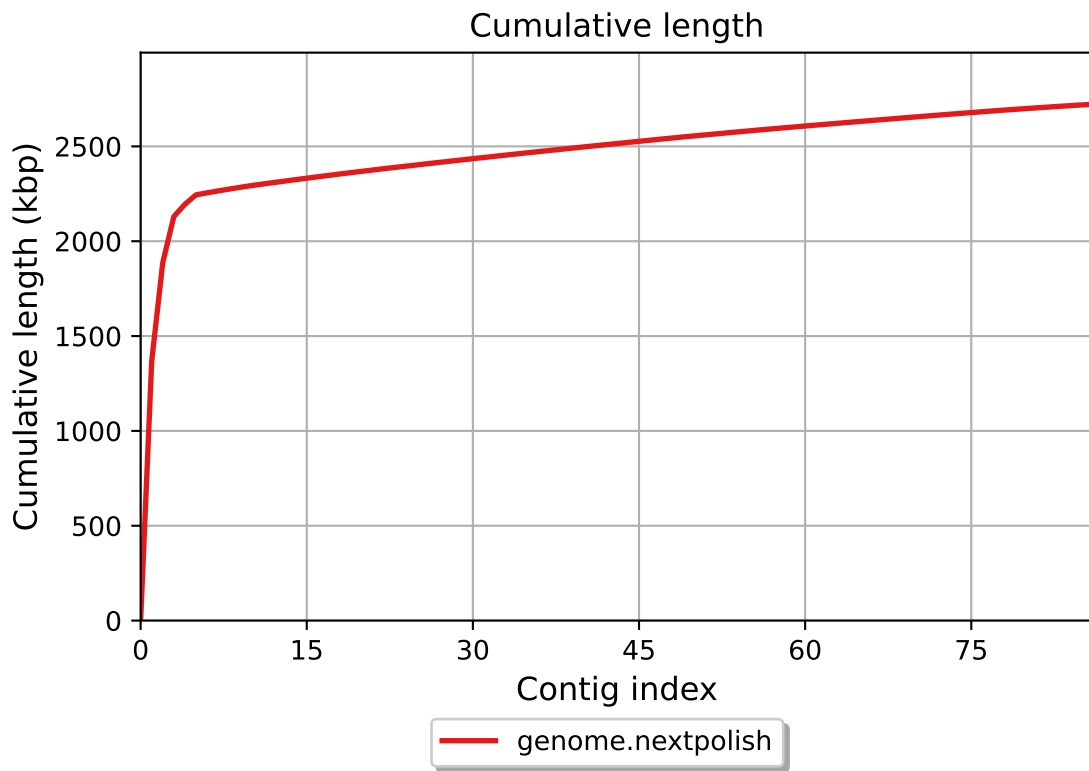


Report

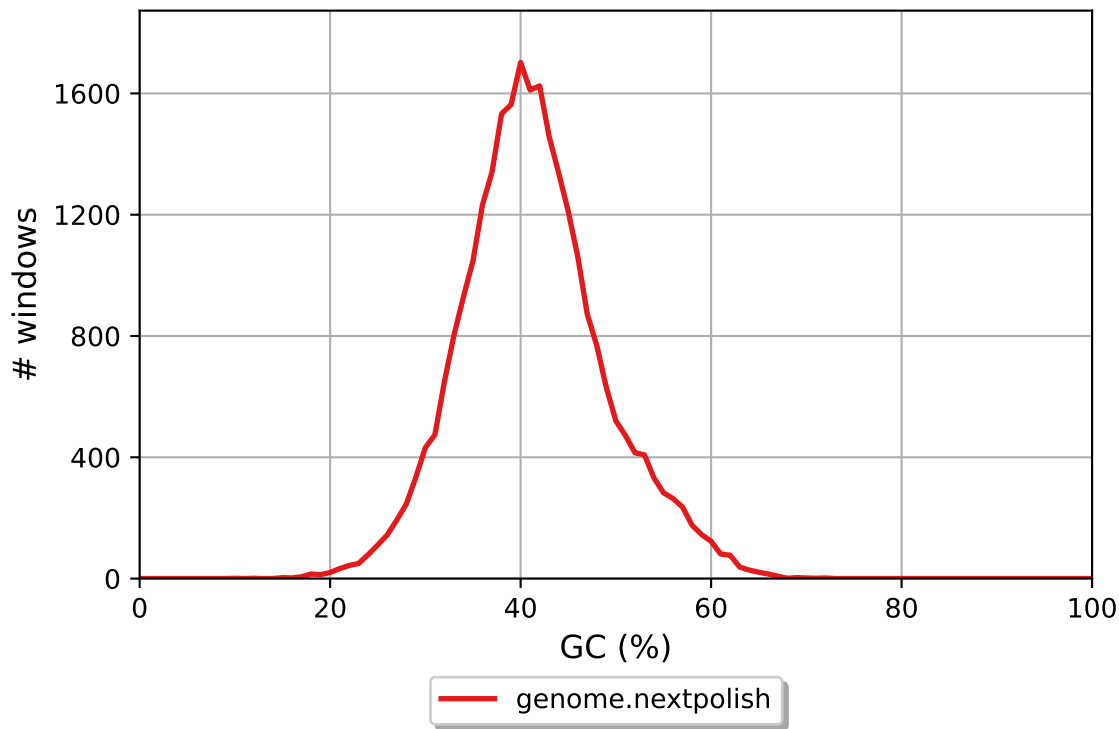
	genome.nextpolish
# contigs (>= 0 bp)	86
# contigs (>= 1000 bp)	86
# contigs (>= 5000 bp)	58
# contigs (>= 10000 bp)	7
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	4
Total length (>= 0 bp)	2722045
Total length (>= 1000 bp)	2722045
Total length (>= 5000 bp)	2597442
Total length (>= 10000 bp)	2265294
Total length (>= 25000 bp)	2244597
Total length (>= 50000 bp)	2194654
# contigs	86
Largest contig	1367551
Total length	2722045
GC (%)	41.32
N50	1367551
N90	6283
auN	811570.3
L50	1
L90	33
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 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).





GC content



genome.nextpolish GC content

