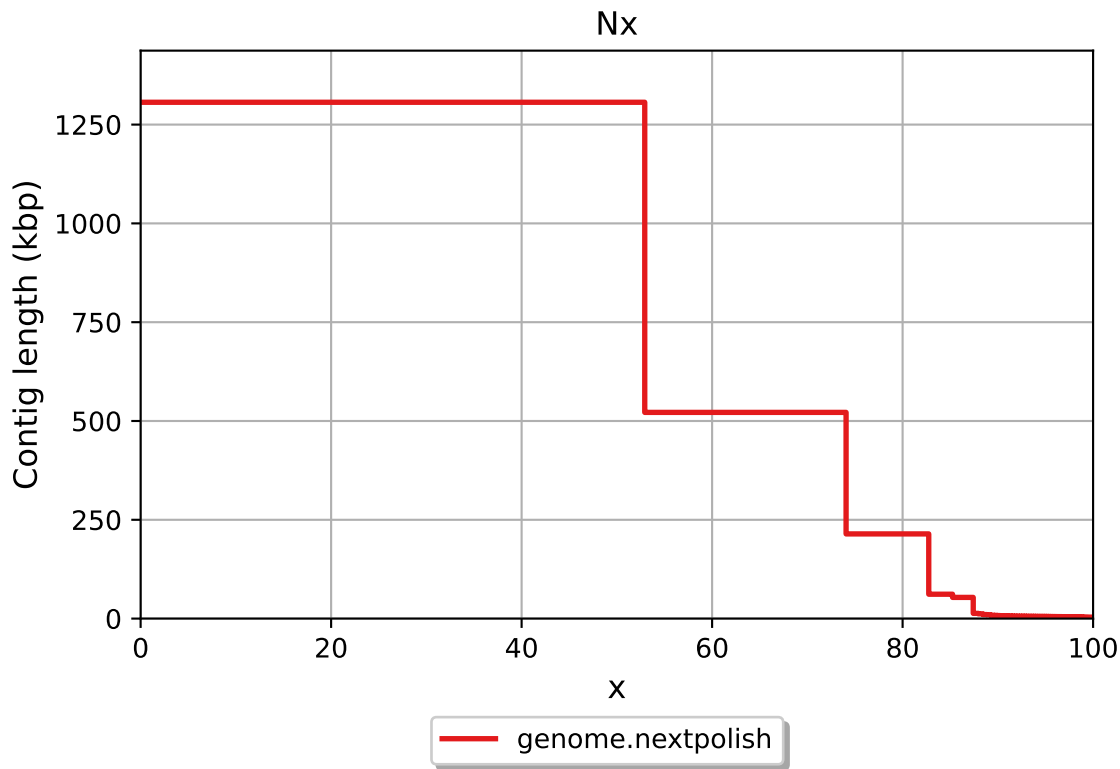
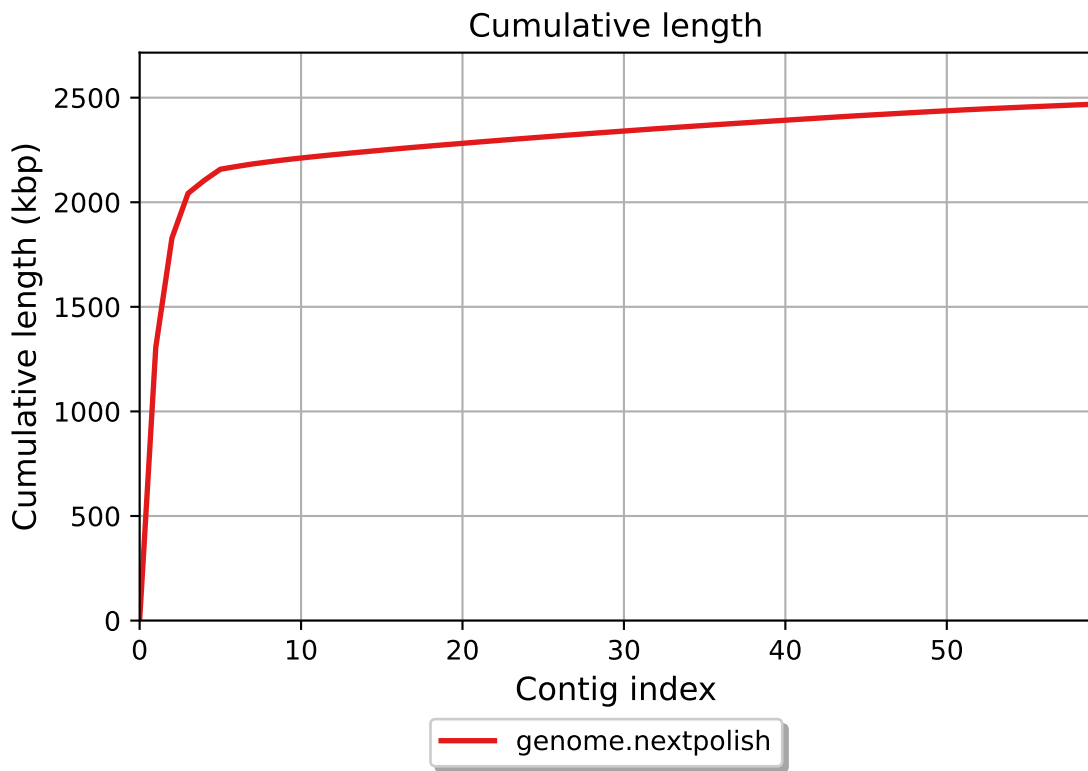


Report

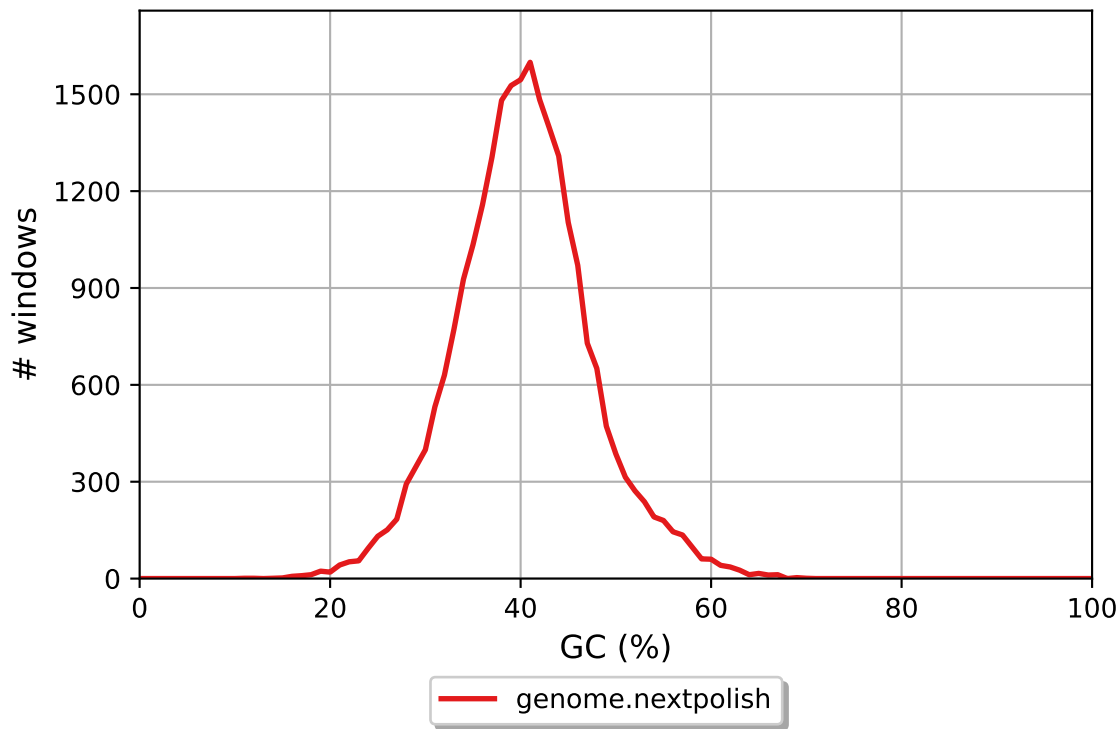
	genome.nextpolish
# contigs (>= 0 bp)	59
# contigs (>= 1000 bp)	59
# contigs (>= 5000 bp)	37
# contigs (>= 10000 bp)	8
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	5
Total length (>= 0 bp)	2468436
Total length (>= 1000 bp)	2468436
Total length (>= 5000 bp)	2377353
Total length (>= 10000 bp)	2193257
Total length (>= 25000 bp)	2157808
Total length (>= 50000 bp)	2157808
# contigs	59
Largest contig	1306495
Total length	2468436
GC (%)	40.35
N50	1306495
N90	7642
auN	823911.1
L50	1
L90	12
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 1 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

