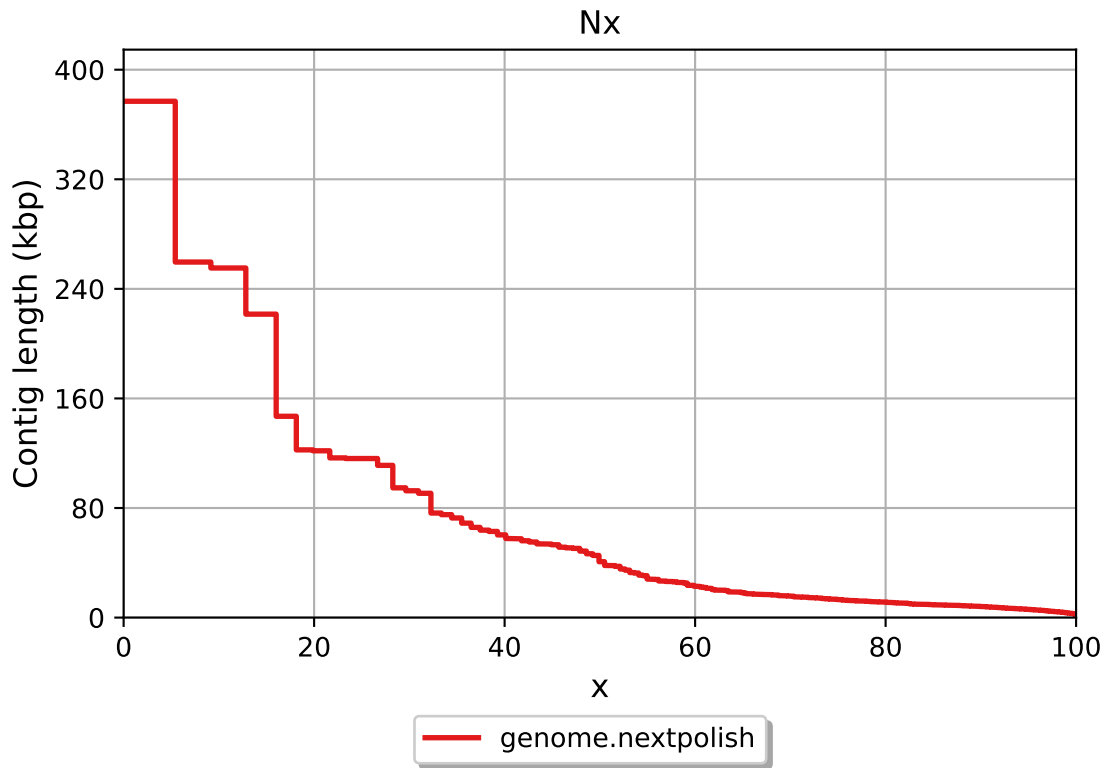
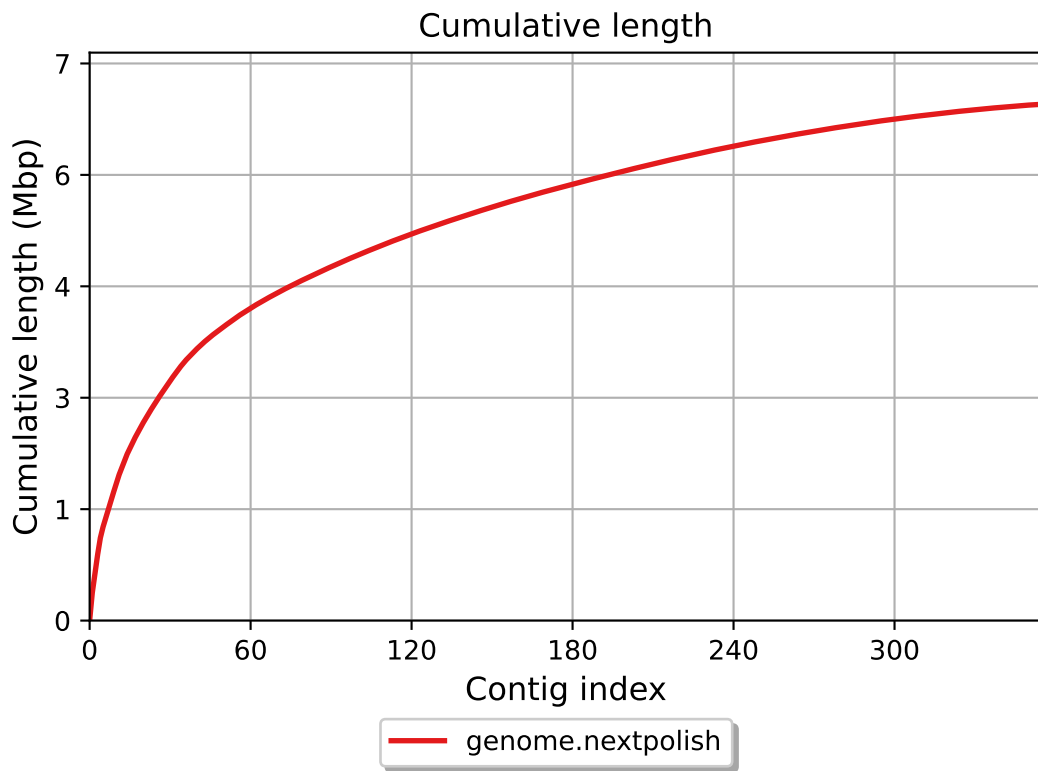


Report

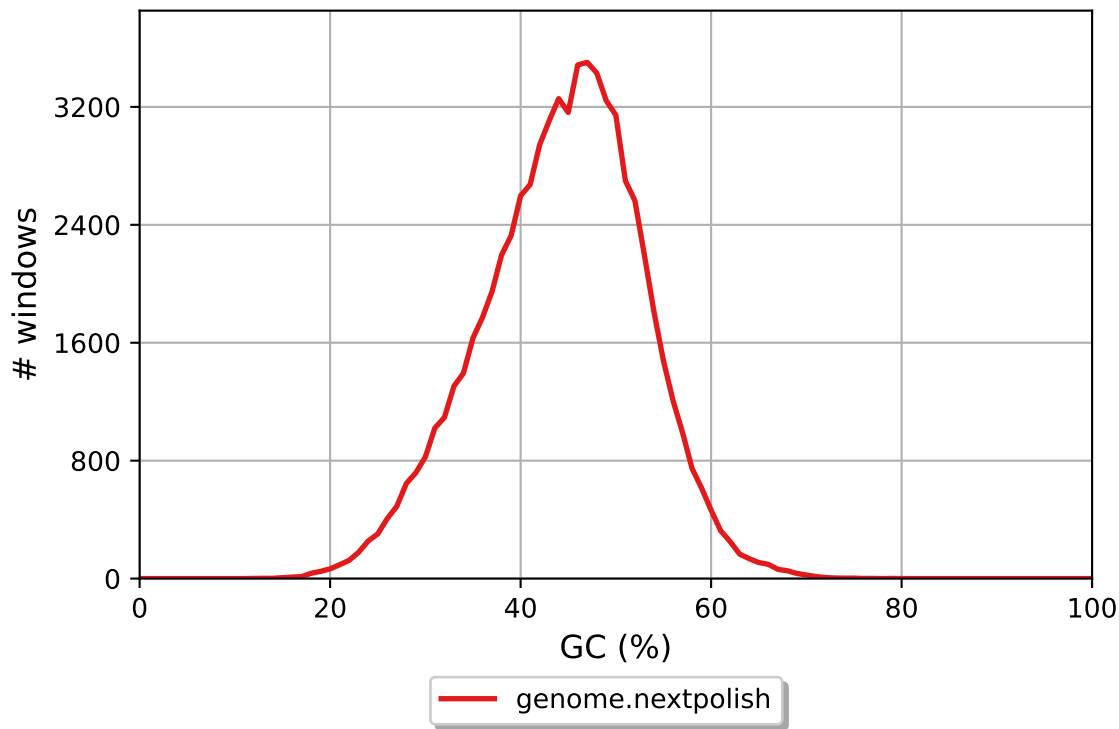
	genome.nextpolish
# contigs (>= 0 bp)	355
# contigs (>= 1000 bp)	355
# contigs (>= 5000 bp)	295
# contigs (>= 10000 bp)	167
# contigs (>= 25000 bp)	56
# contigs (>= 50000 bp)	32
Total length (>= 0 bp)	6950715
Total length (>= 1000 bp)	6950715
Total length (>= 5000 bp)	6727172
Total length (>= 10000 bp)	5748338
Total length (>= 25000 bp)	4112817
Total length (>= 50000 bp)	3328563
# contigs	355
Largest contig	376950
Total length	6950715
GC (%)	44.30
N50	40797
N90	8116
auN	83649.6
L50	36
L90	223
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	26 + 4 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

