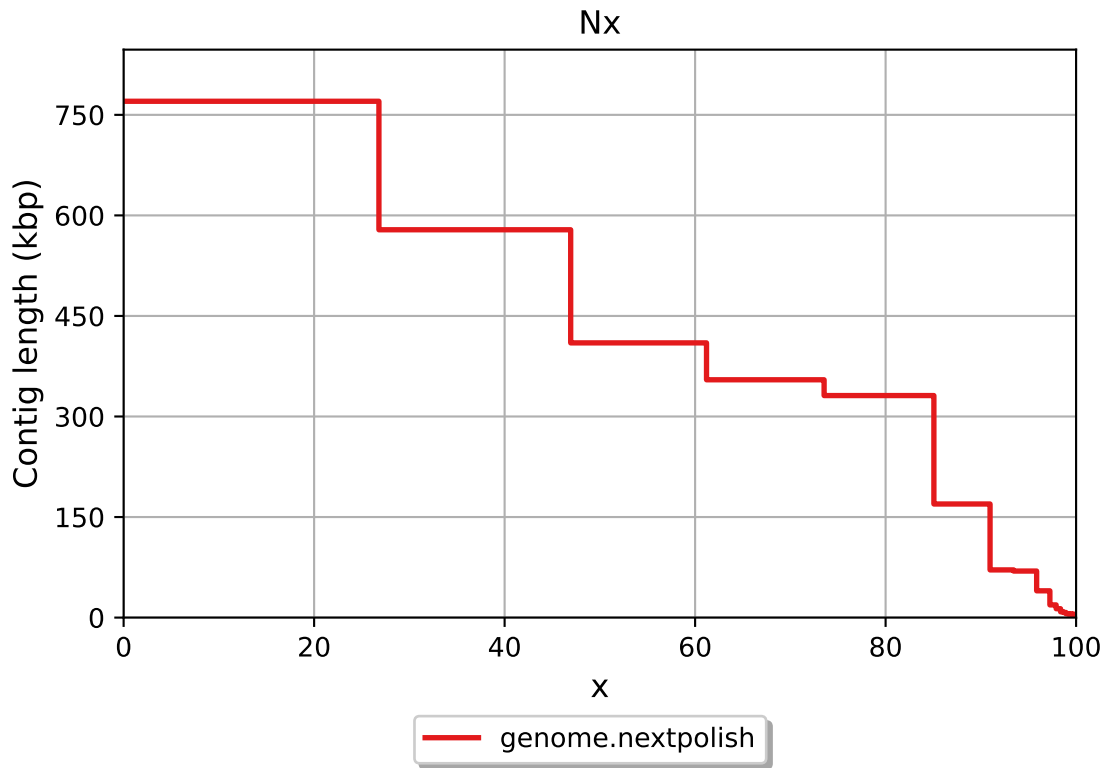
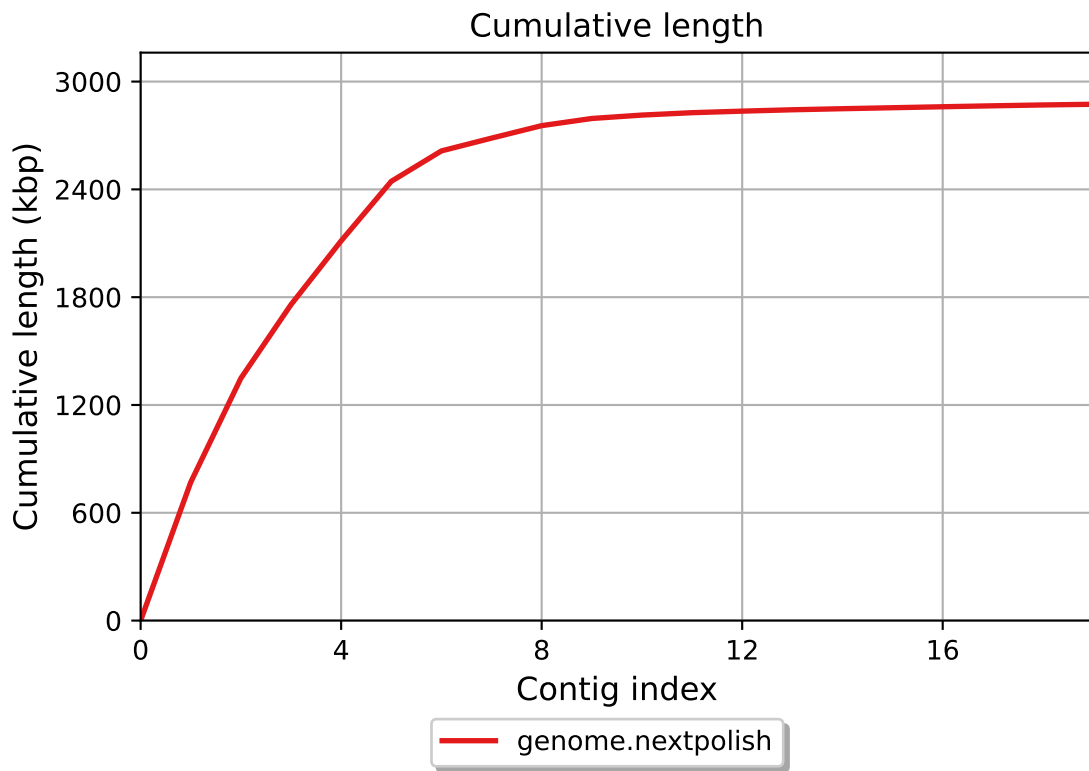


# Report

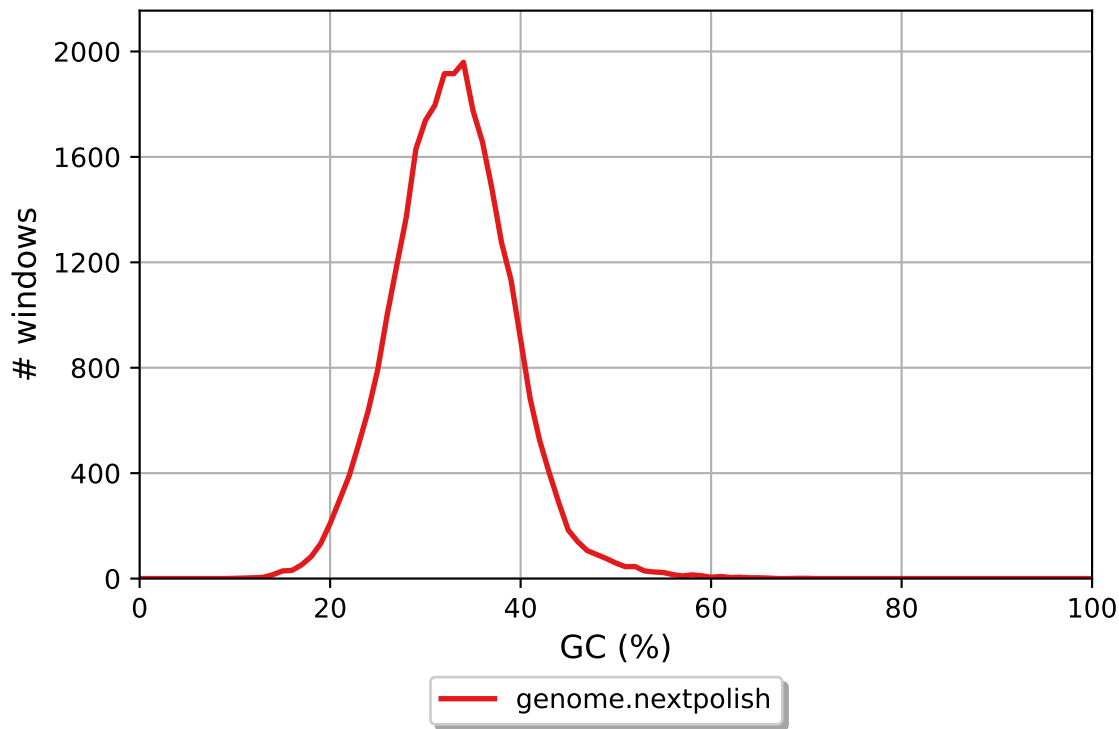
	genome.nextpolish
# contigs (>= 0 bp)	19
# contigs (>= 1000 bp)	19
# contigs (>= 5000 bp)	17
# contigs (>= 10000 bp)	11
# contigs (>= 25000 bp)	9
# contigs (>= 50000 bp)	8
Total length (>= 0 bp)	2873711
Total length (>= 1000 bp)	2873711
Total length (>= 5000 bp)	2865156
Total length (>= 10000 bp)	2826809
Total length (>= 25000 bp)	2794623
Total length (>= 50000 bp)	2754674
# contigs	19
Largest contig	770263
Total length	2873711
GC (%)	32.87
N50	409816
N90	169466
auN	477637.8
L50	3
L90	6
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 7 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

