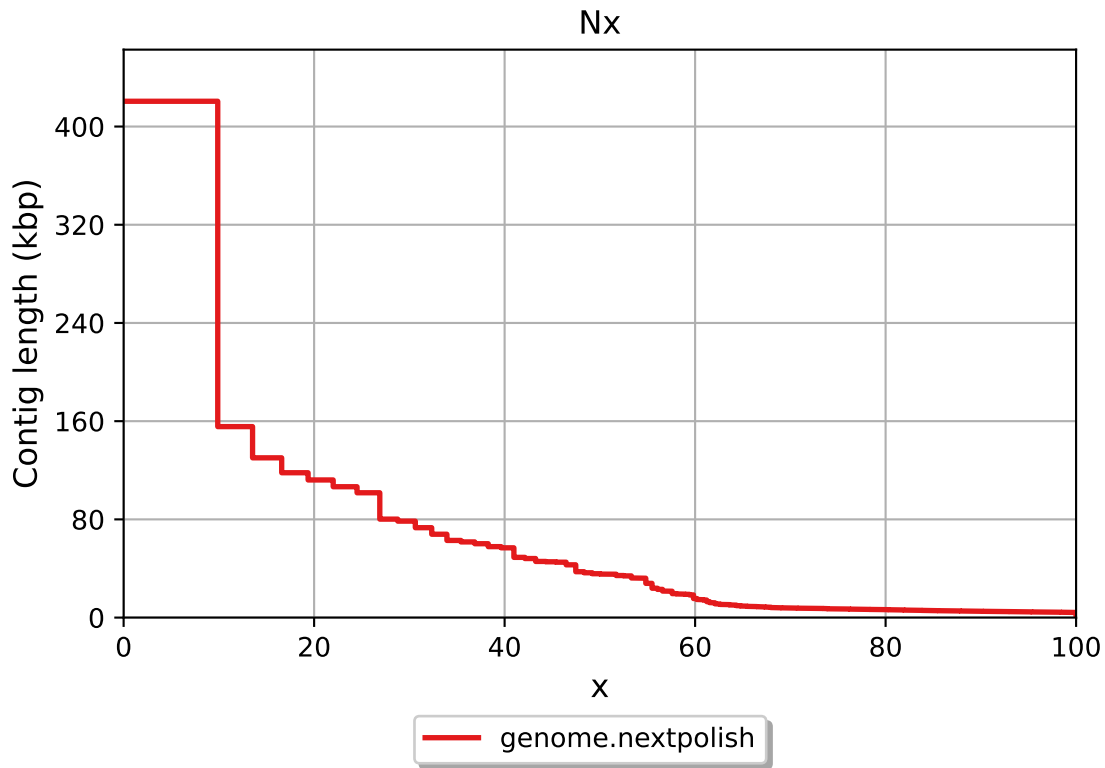
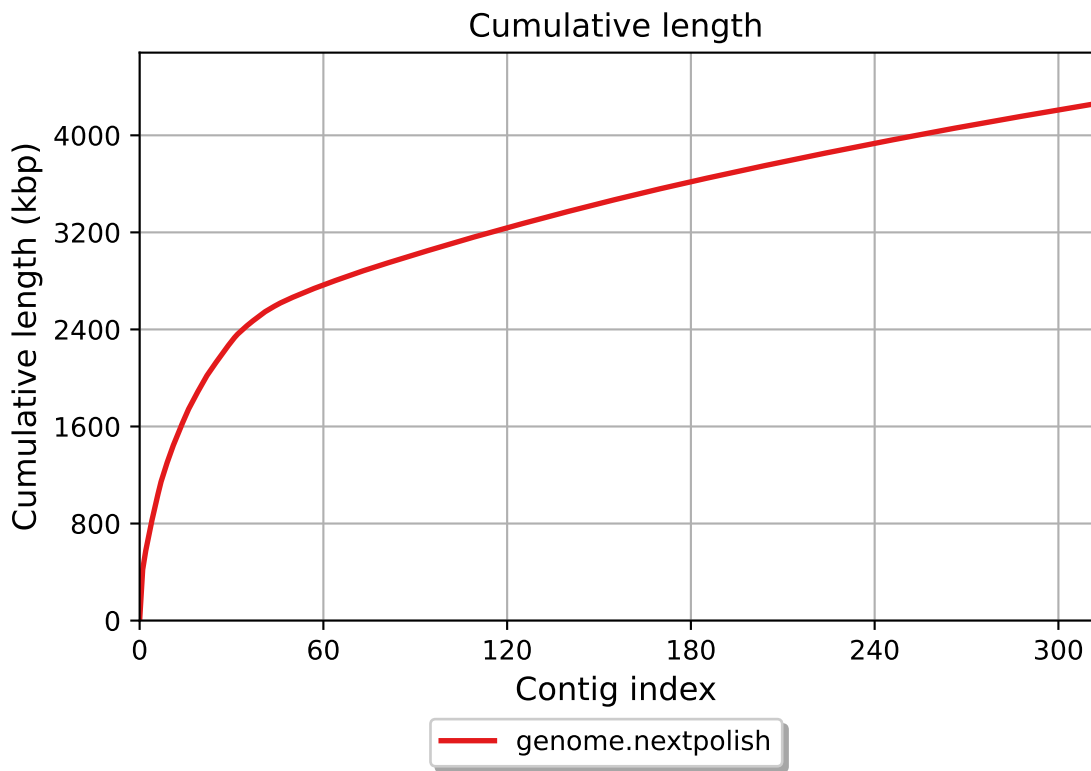


# Report

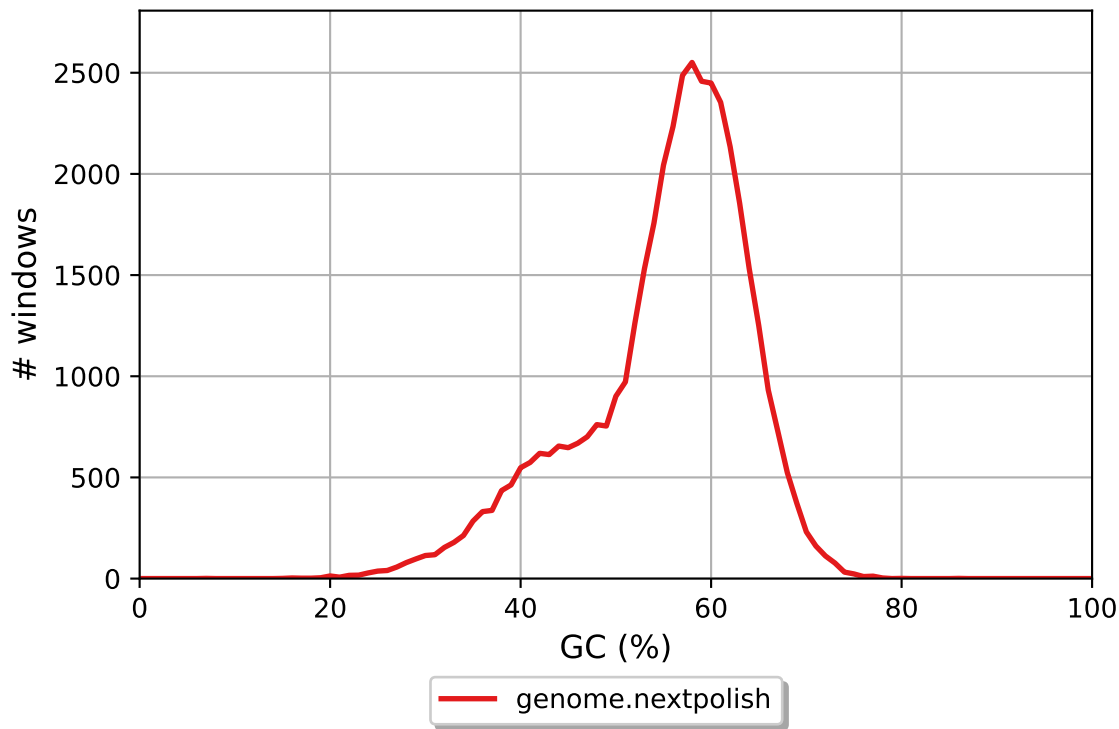
	genome.nextpolish
# contigs (>= 0 bp)	311
# contigs (>= 1000 bp)	311
# contigs (>= 5000 bp)	234
# contigs (>= 10000 bp)	57
# contigs (>= 25000 bp)	32
# contigs (>= 50000 bp)	16
Total length (>= 0 bp)	4255978
Total length (>= 1000 bp)	4255978
Total length (>= 5000 bp)	3904354
Total length (>= 10000 bp)	2737996
Total length (>= 25000 bp)	2361107
Total length (>= 50000 bp)	1743867
# contigs	311
Largest contig	420585
Total length	4255978
GC (%)	55.08
N50	35734
N90	5138
auN	81514.7
L50	25
L90	220
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 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

