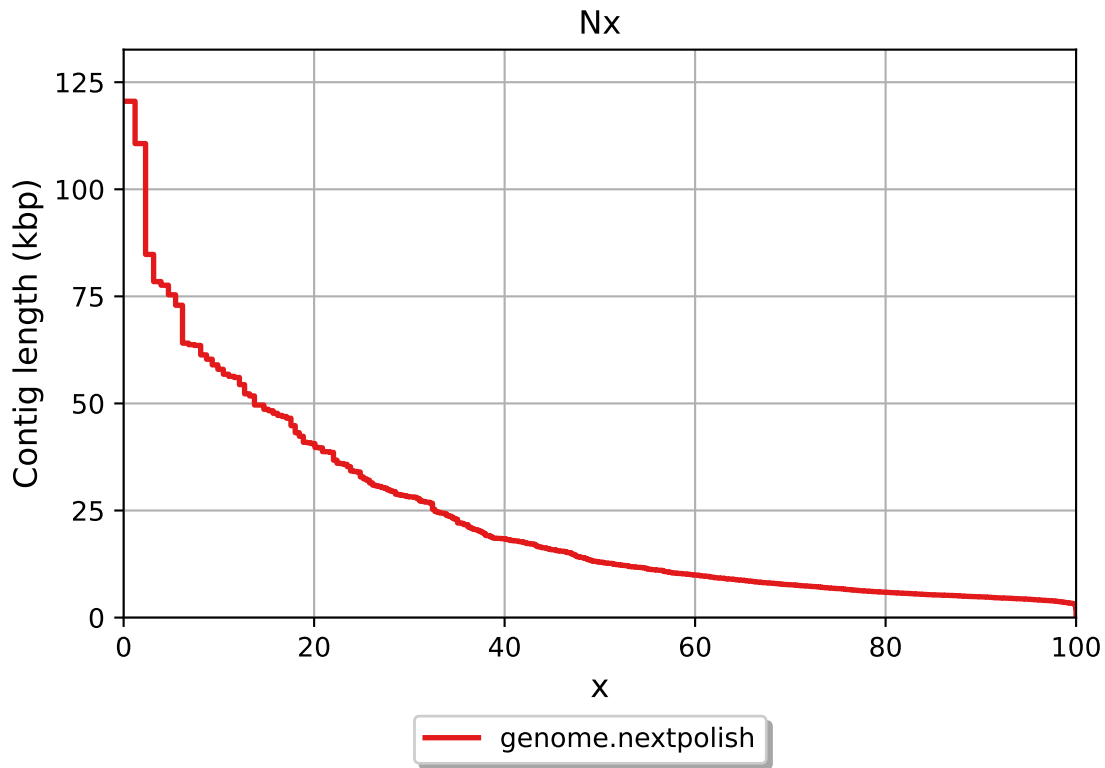


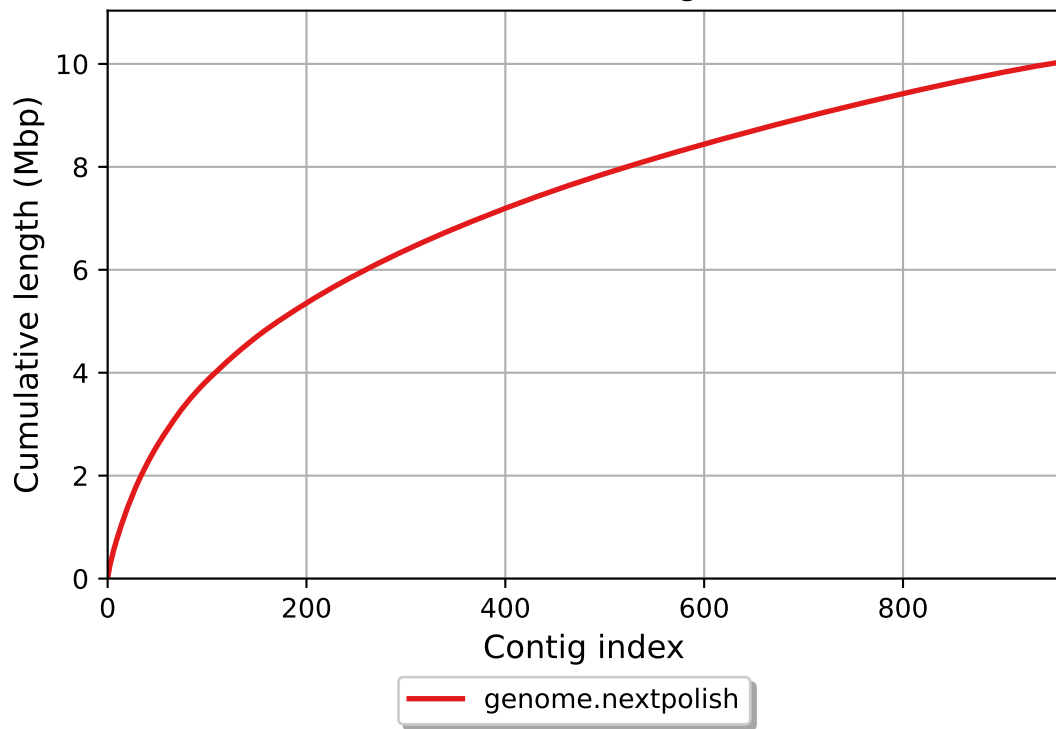
# Report

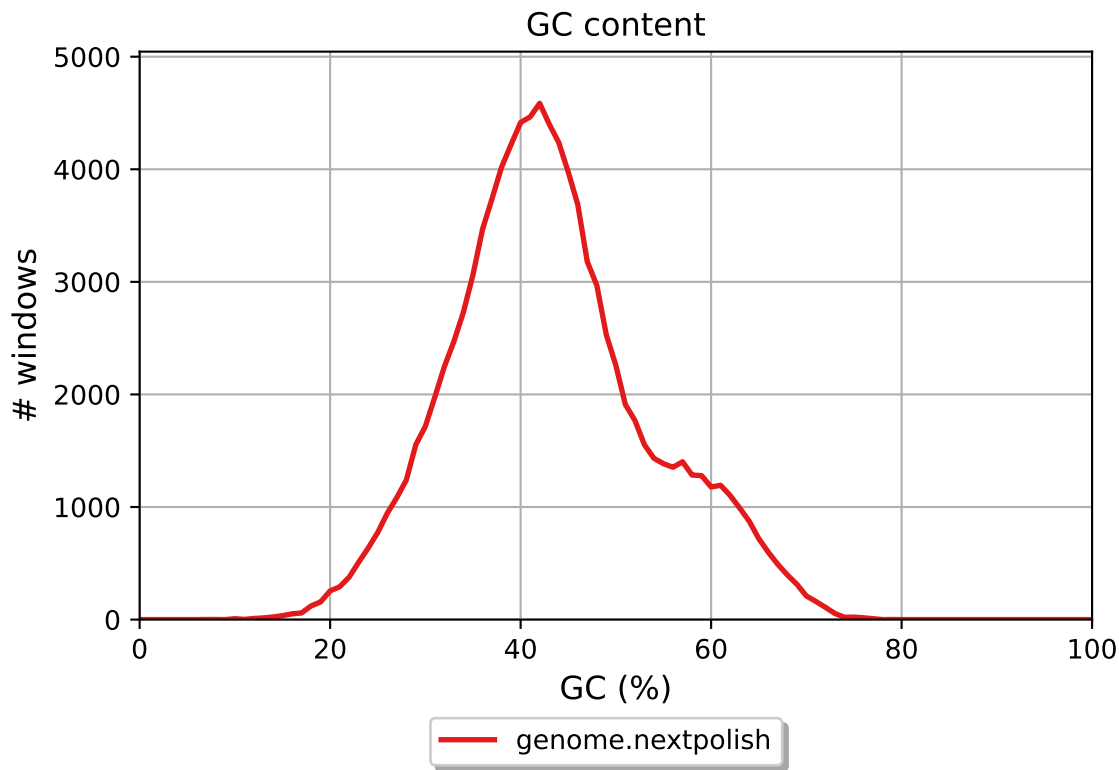
	genome.nextpolish
# contigs (>= 0 bp)	958
# contigs (>= 1000 bp)	958
# contigs (>= 5000 bp)	678
# contigs (>= 10000 bp)	258
# contigs (>= 25000 bp)	74
# contigs (>= 50000 bp)	20
Total length (>= 0 bp)	10032176
Total length (>= 1000 bp)	10032176
Total length (>= 5000 bp)	8848910
Total length (>= 10000 bp)	5985924
Total length (>= 25000 bp)	3278241
Total length (>= 50000 bp)	1377886
# contigs	958
Largest contig	120550
Total length	10032176
GC (%)	43.17
N50	12943
N90	4842
auN	23851.7
L50	173
L90	715
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	13 + 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).



Cumulative length





genome.nextpolish GC content

