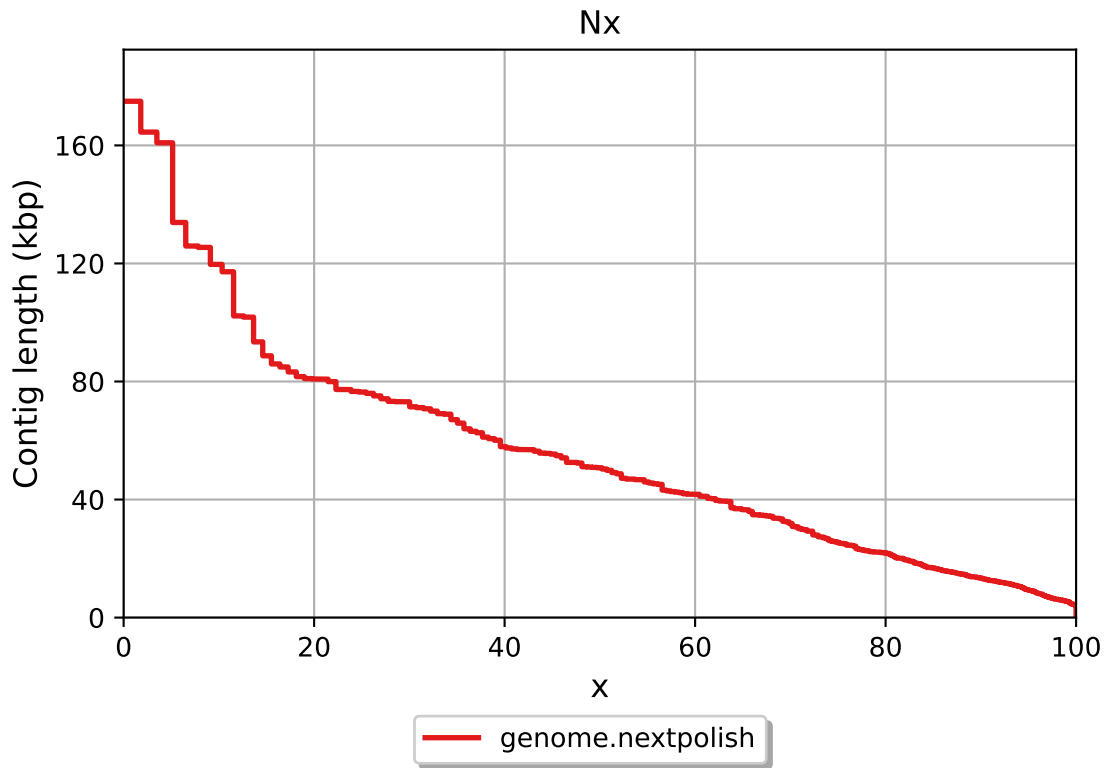


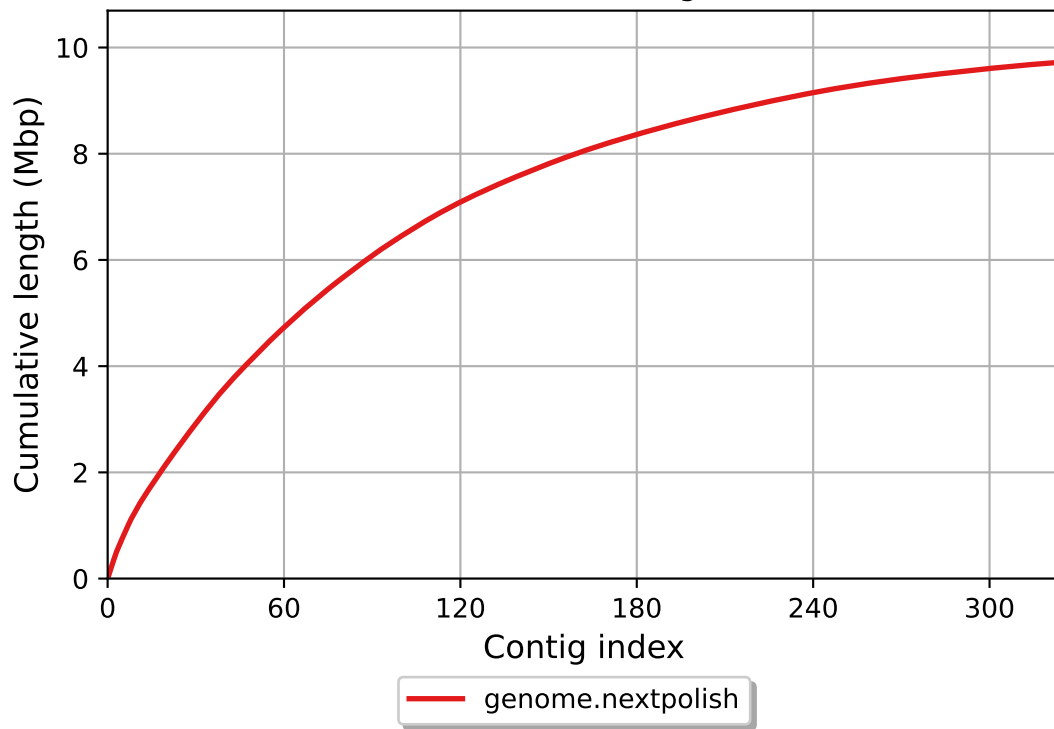
Report

	genome.nextpolish
# contigs (>= 0 bp)	324
# contigs (>= 1000 bp)	324
# contigs (>= 5000 bp)	309
# contigs (>= 10000 bp)	244
# contigs (>= 25000 bp)	131
# contigs (>= 50000 bp)	64
Total length (>= 0 bp)	9722237
Total length (>= 1000 bp)	9722237
Total length (>= 5000 bp)	9654528
Total length (>= 10000 bp)	9191833
Total length (>= 25000 bp)	7376867
Total length (>= 50000 bp)	4932482
# contigs	324
Largest contig	174956
Total length	9722237
GC (%)	56.13
N50	50776
N90	13403
auN	57351.4
L50	63
L90	207
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	18 + 2 part

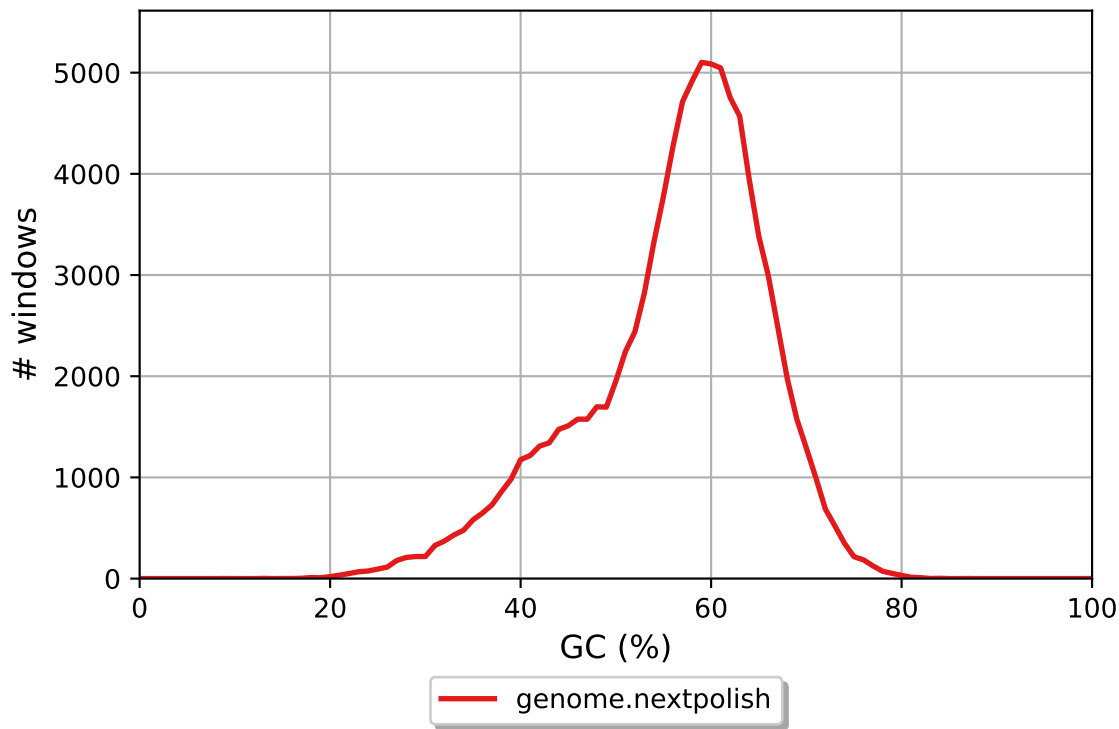
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

