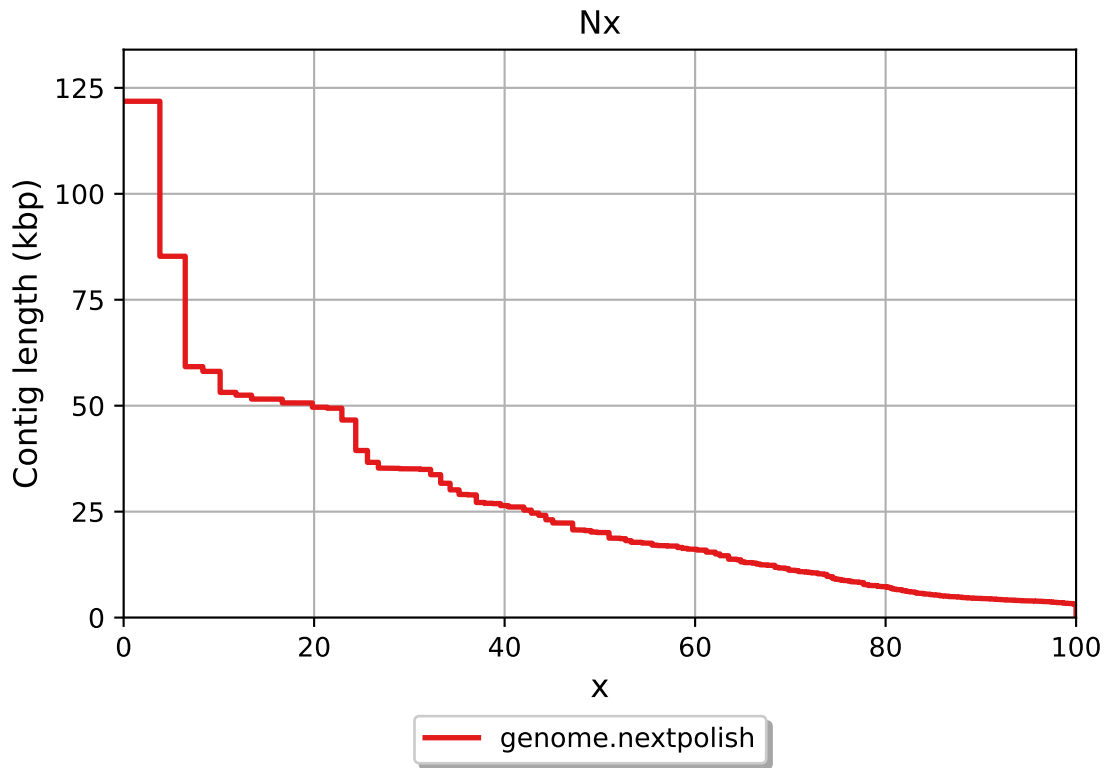


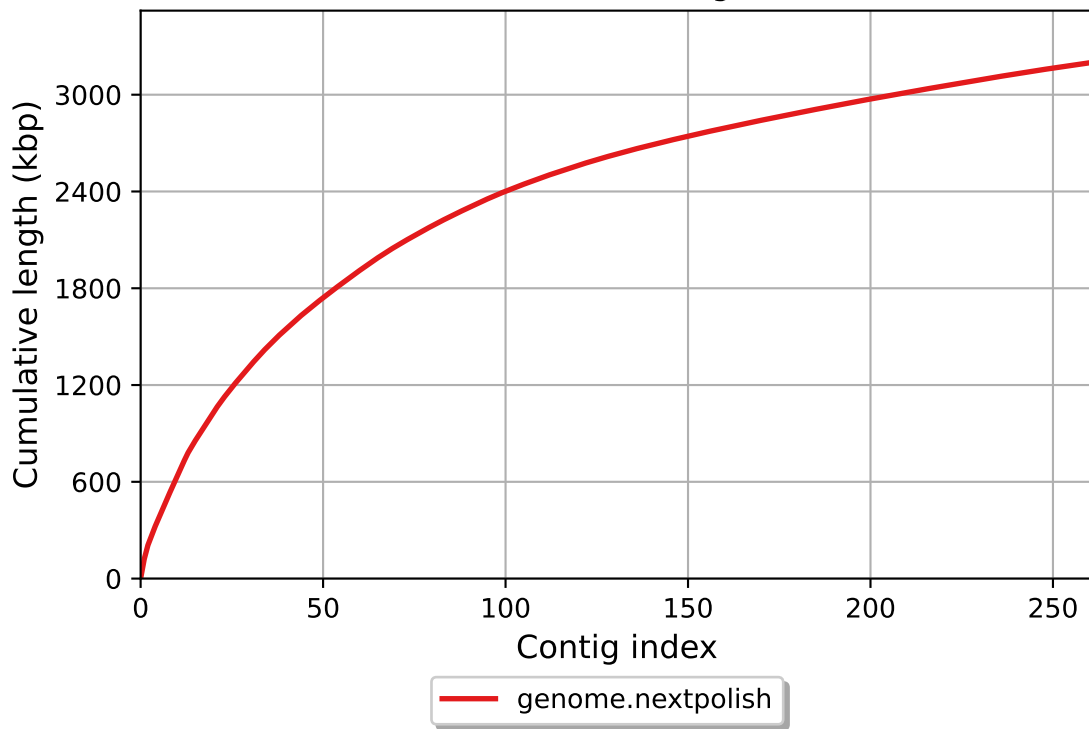
# Report

	genome.nextpolish
# contigs (>= 0 bp)	261
# contigs (>= 1000 bp)	261
# contigs (>= 5000 bp)	156
# contigs (>= 10000 bp)	96
# contigs (>= 25000 bp)	32
# contigs (>= 50000 bp)	10
Total length (>= 0 bp)	3201117
Total length (>= 1000 bp)	3201117
Total length (>= 5000 bp)	2773560
Total length (>= 10000 bp)	2363748
Total length (>= 25000 bp)	1370381
Total length (>= 50000 bp)	634438
# contigs	261
Largest contig	121835
Total length	3201117
GC (%)	56.30
N50	20071
N90	4522
auN	28699.0
L50	43
L90	179
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	12 + 1 part

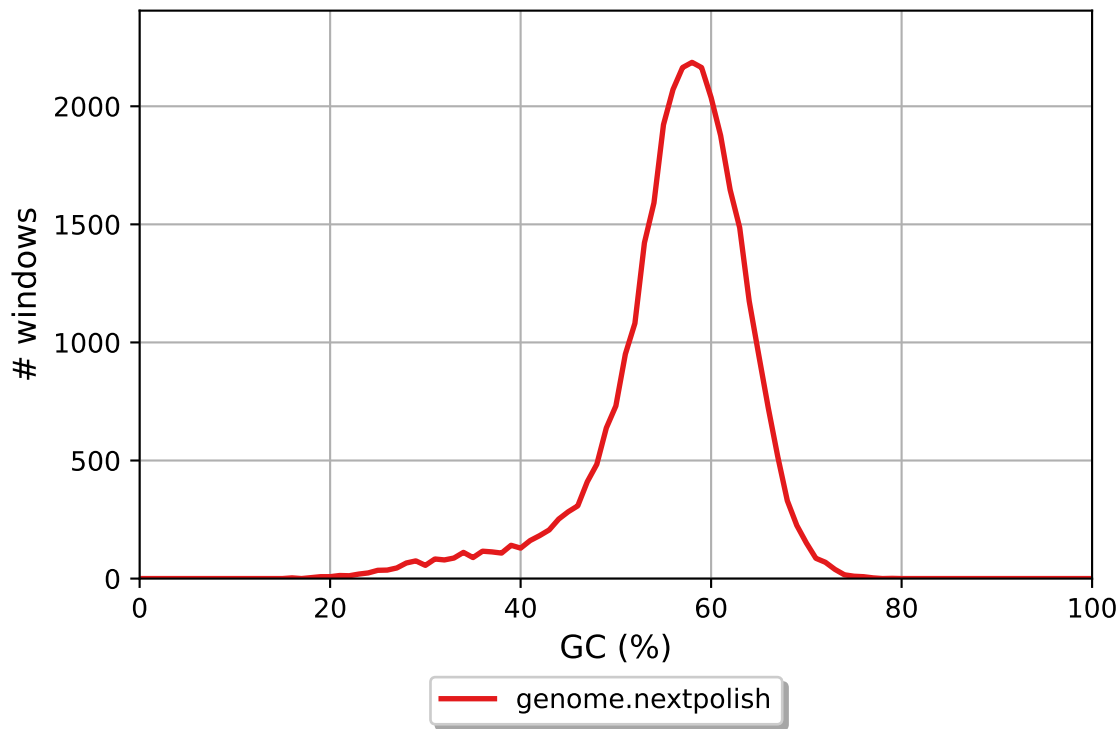
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

