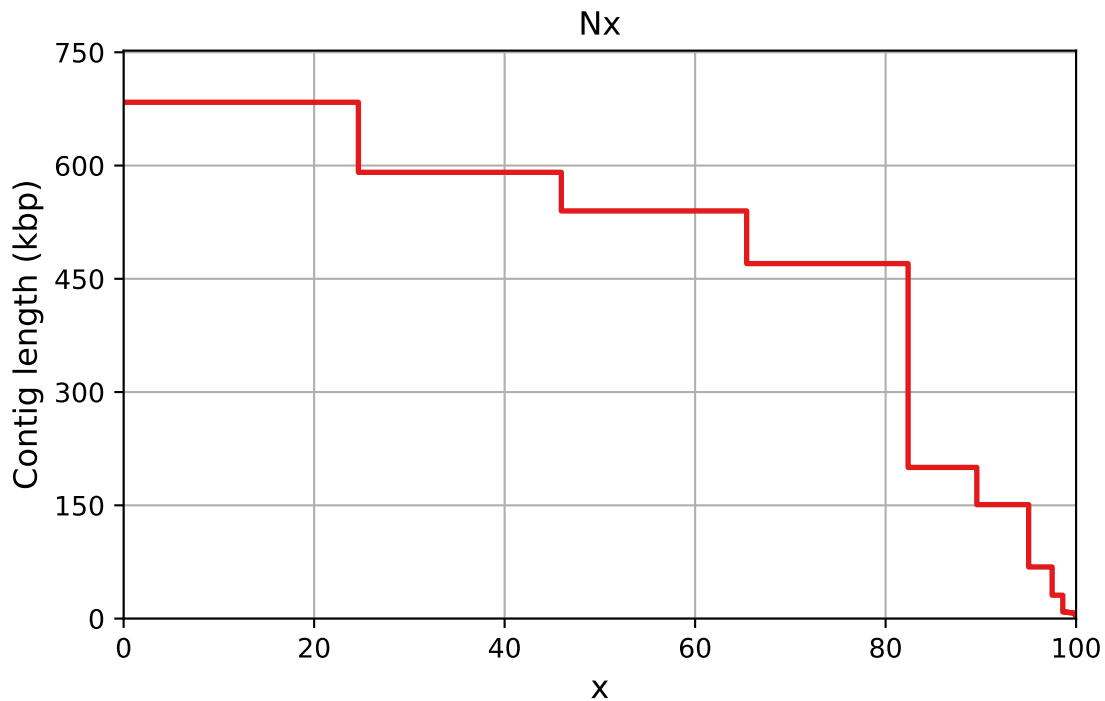


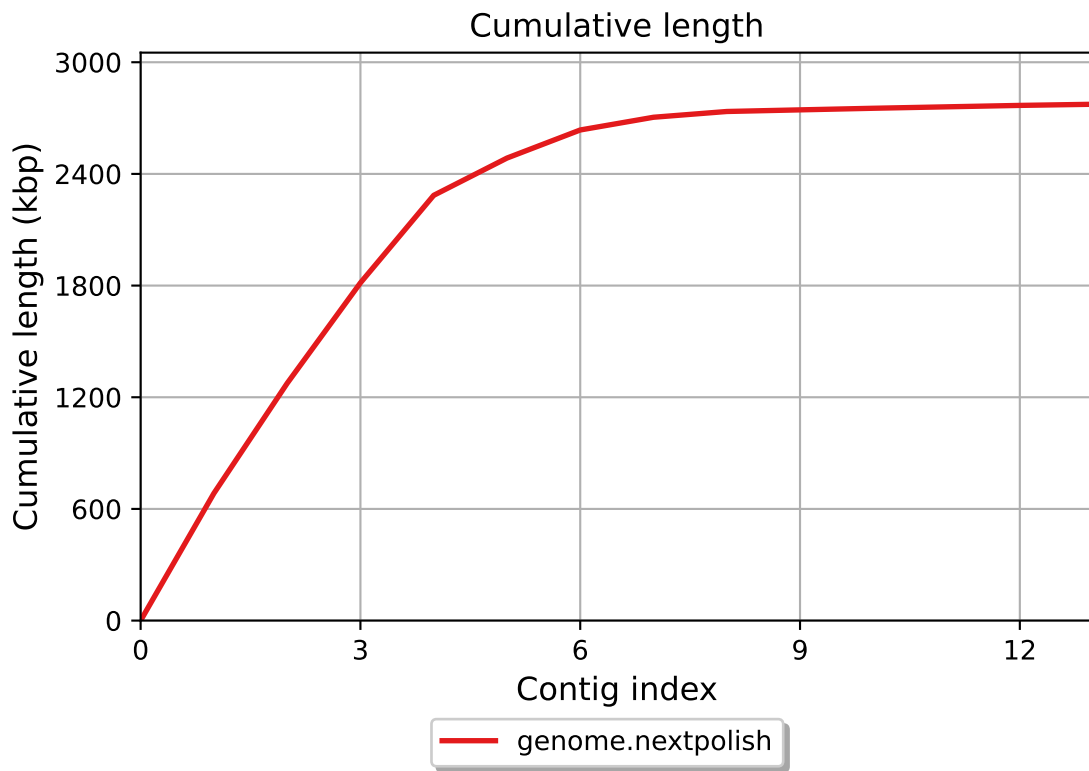
# Report

	genome.nextpolish
# contigs (>= 0 bp)	13
# contigs (>= 1000 bp)	13
# contigs (>= 5000 bp)	13
# contigs (>= 10000 bp)	8
# contigs (>= 25000 bp)	8
# contigs (>= 50000 bp)	7
Total length (>= 0 bp)	2774168
Total length (>= 1000 bp)	2774168
Total length (>= 5000 bp)	2774168
Total length (>= 10000 bp)	2735370
Total length (>= 25000 bp)	2735370
Total length (>= 50000 bp)	2704387
# contigs	13
Largest contig	683796
Total length	2774168
GC (%)	58.44
N50	539881
N90	150924
auN	504014.8
L50	3
L90	6
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	12 + 2 part

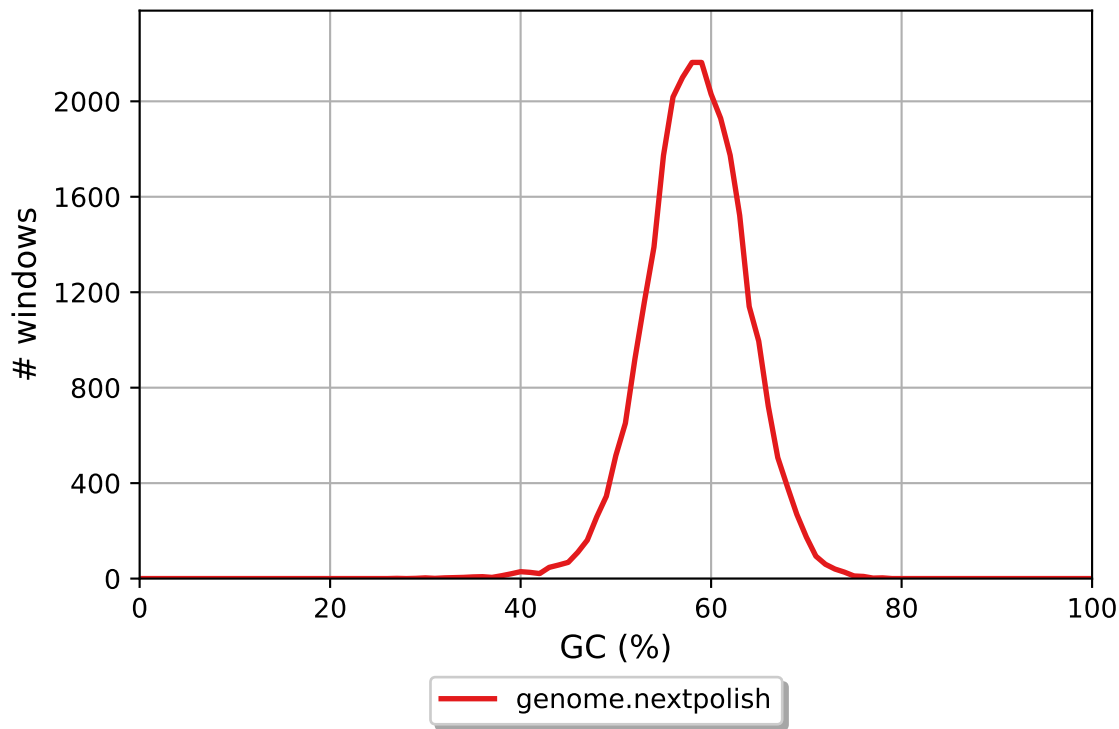
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



## GC content



genome.nextpolish GC content

