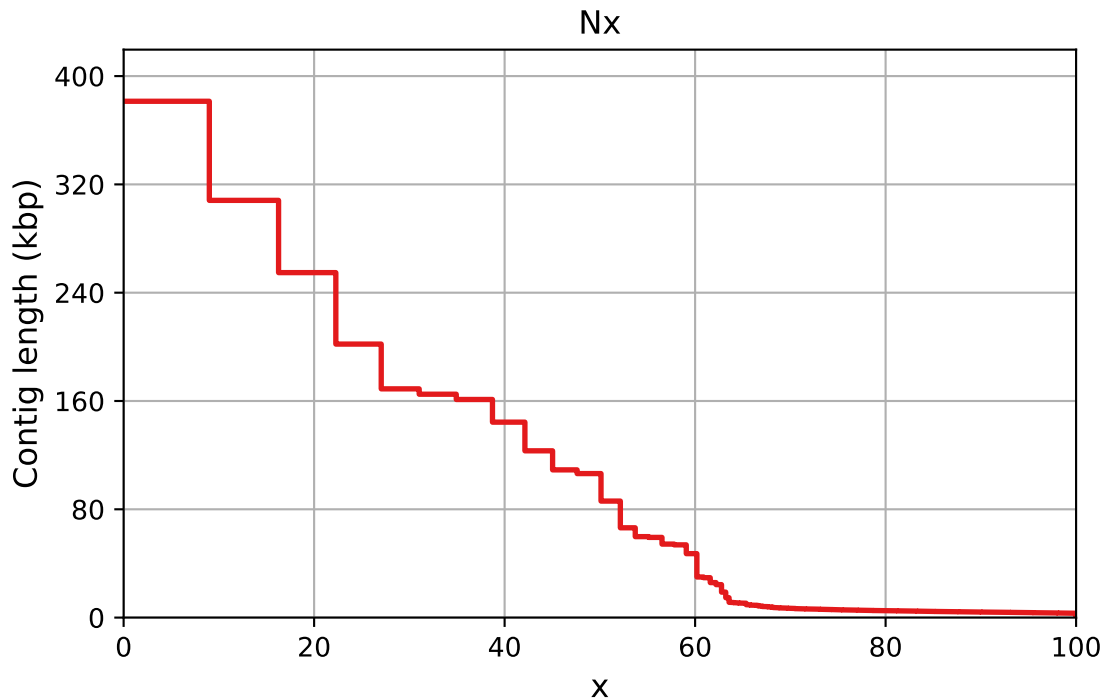


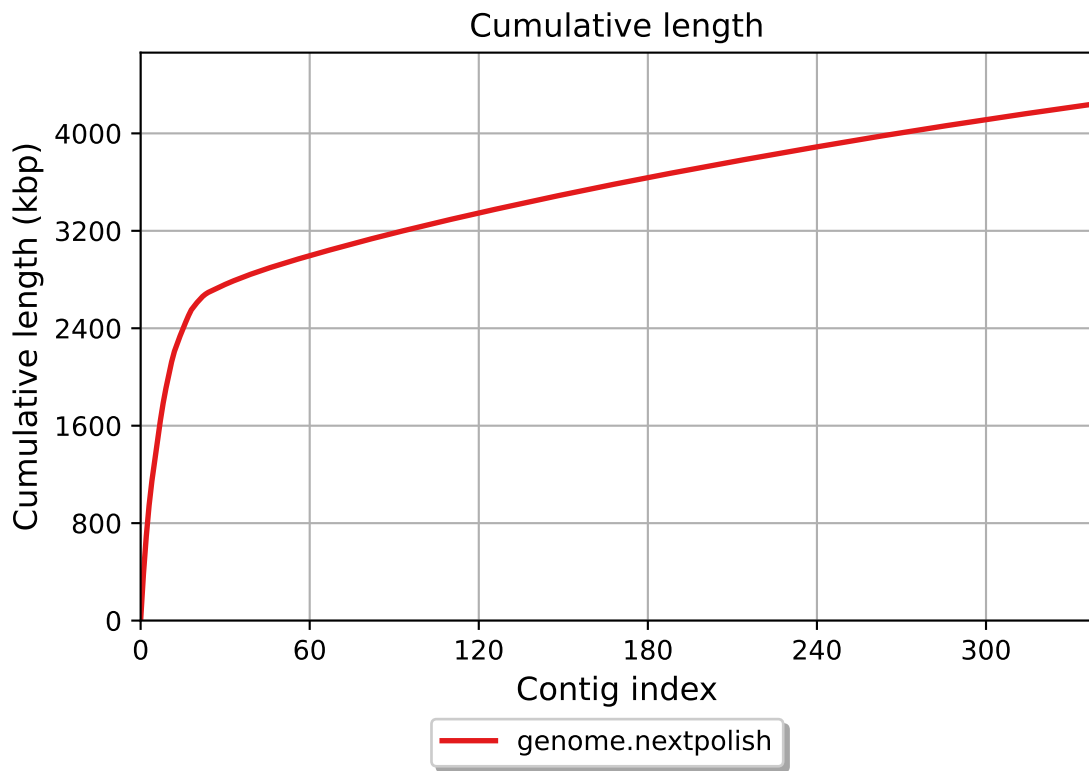
Report

	genome.nextpolish
# contigs (>= 0 bp)	338
# contigs (>= 1000 bp)	338
# contigs (>= 5000 bp)	139
# contigs (>= 10000 bp)	31
# contigs (>= 25000 bp)	21
# contigs (>= 50000 bp)	17
Total length (>= 0 bp)	4238966
Total length (>= 1000 bp)	4238966
Total length (>= 5000 bp)	3443300
Total length (>= 10000 bp)	2770585
Total length (>= 25000 bp)	2636583
Total length (>= 50000 bp)	2504047
# contigs	338
Largest contig	381402
Total length	4238966
GC (%)	57.02
N50	106381
N90	4095
auN	124116.4
L50	11
L90	222
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	17 + 4 part

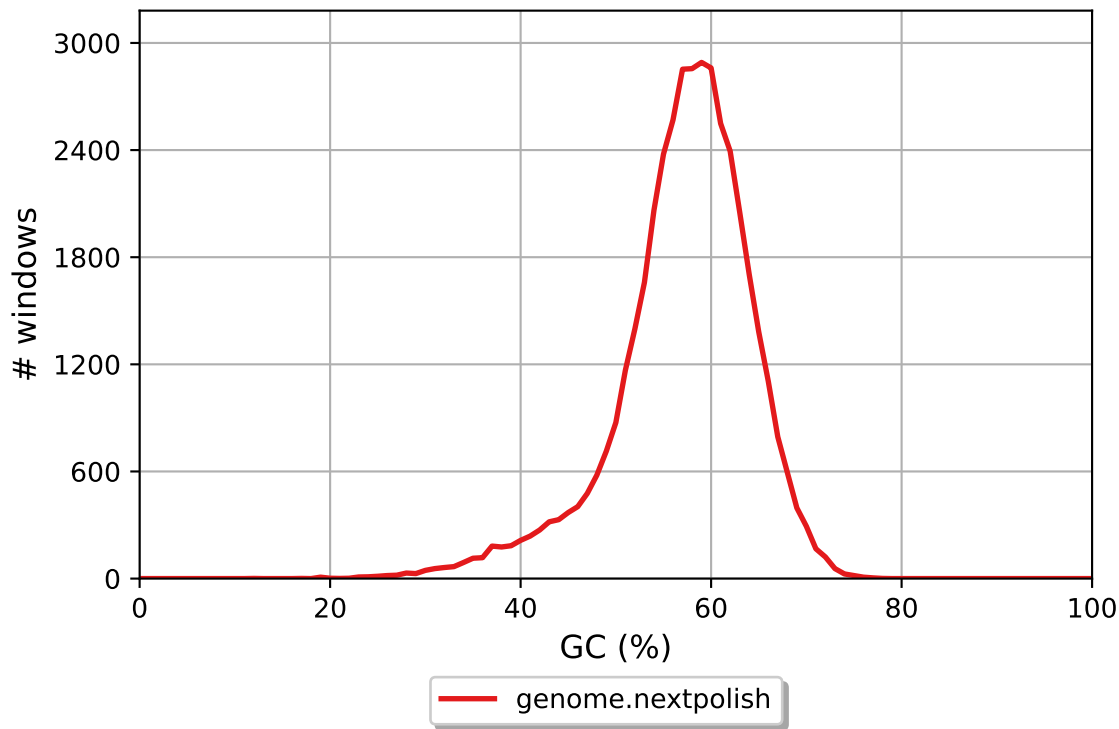
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



GC content



genome.nextpolish GC content

