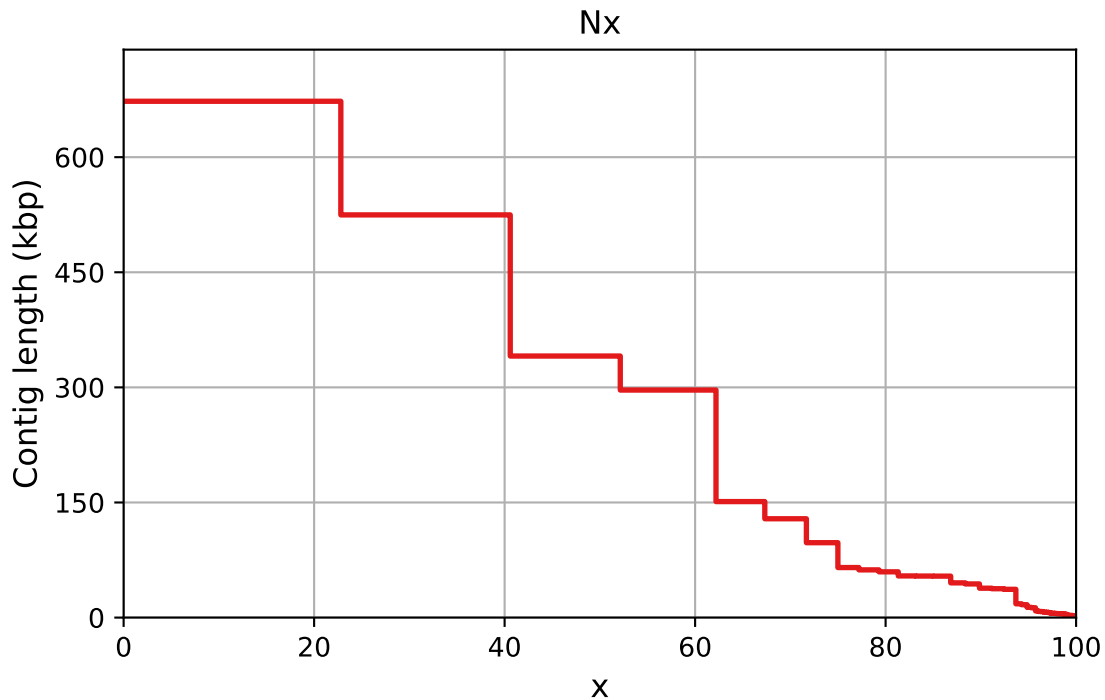


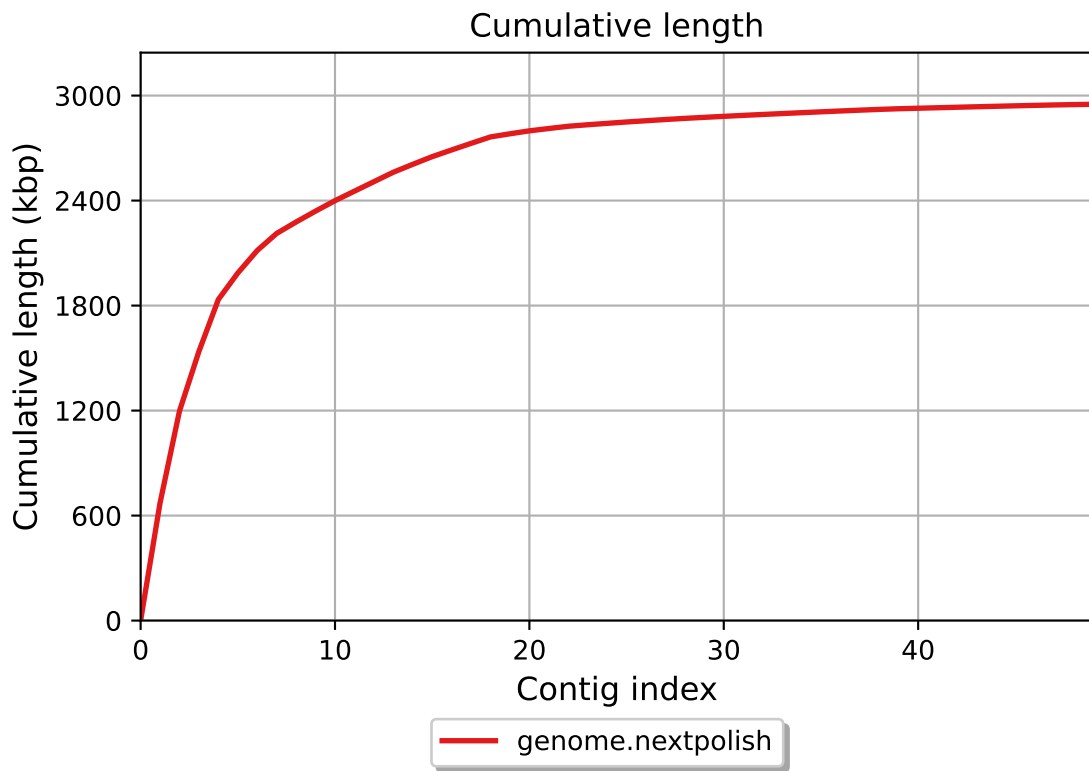
Report

	genome.nextpolish
# contigs (>= 0 bp)	49
# contigs (>= 1000 bp)	49
# contigs (>= 5000 bp)	33
# contigs (>= 10000 bp)	22
# contigs (>= 25000 bp)	18
# contigs (>= 50000 bp)	13
Total length (>= 0 bp)	2950378
Total length (>= 1000 bp)	2950378
Total length (>= 5000 bp)	2897252
Total length (>= 10000 bp)	2824680
Total length (>= 25000 bp)	2763807
Total length (>= 50000 bp)	2561957
# contigs	49
Largest contig	672836
Total length	2950378
GC (%)	32.90
N50	340837
N90	38289
auN	342821.9
L50	3
L90	16
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	10 + 11 part

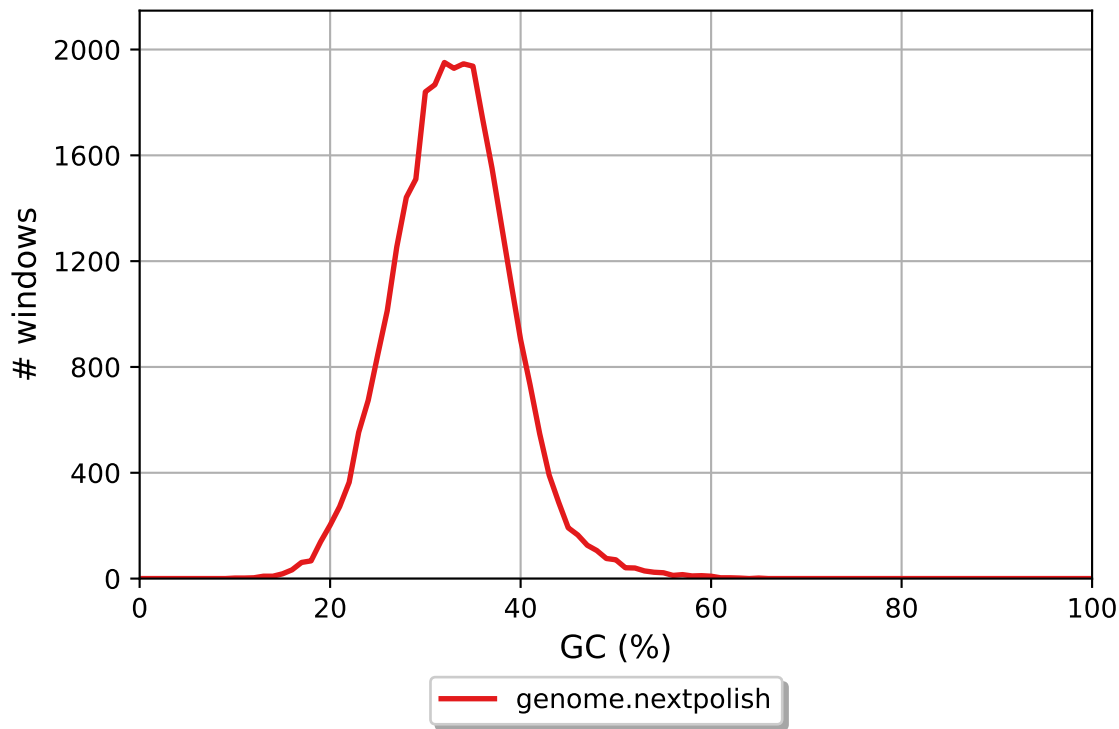
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



GC content



genome.nextpolish GC content

