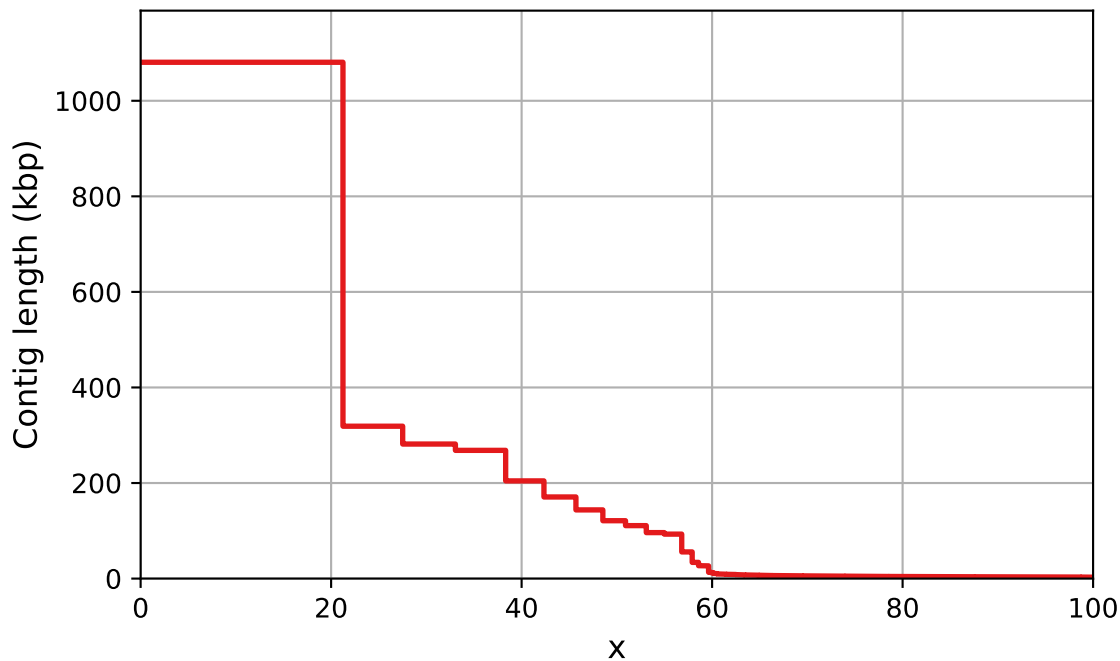


Report

	genome.nextpolish
# contigs (>= 0 bp)	474
# contigs (>= 1000 bp)	473
# contigs (>= 5000 bp)	131
# contigs (>= 10000 bp)	19
# contigs (>= 25000 bp)	15
# contigs (>= 50000 bp)	12
Total length (>= 0 bp)	5086545
Total length (>= 1000 bp)	5086012
Total length (>= 5000 bp)	3783010
Total length (>= 10000 bp)	3079721
Total length (>= 25000 bp)	3033045
Total length (>= 50000 bp)	2945509
# contigs	474
Largest contig	1080591
Total length	5086545
GC (%)	39.69
N50	121127
N90	3662
auN	309264.6
L50	8
L90	322
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	19 + 4 part

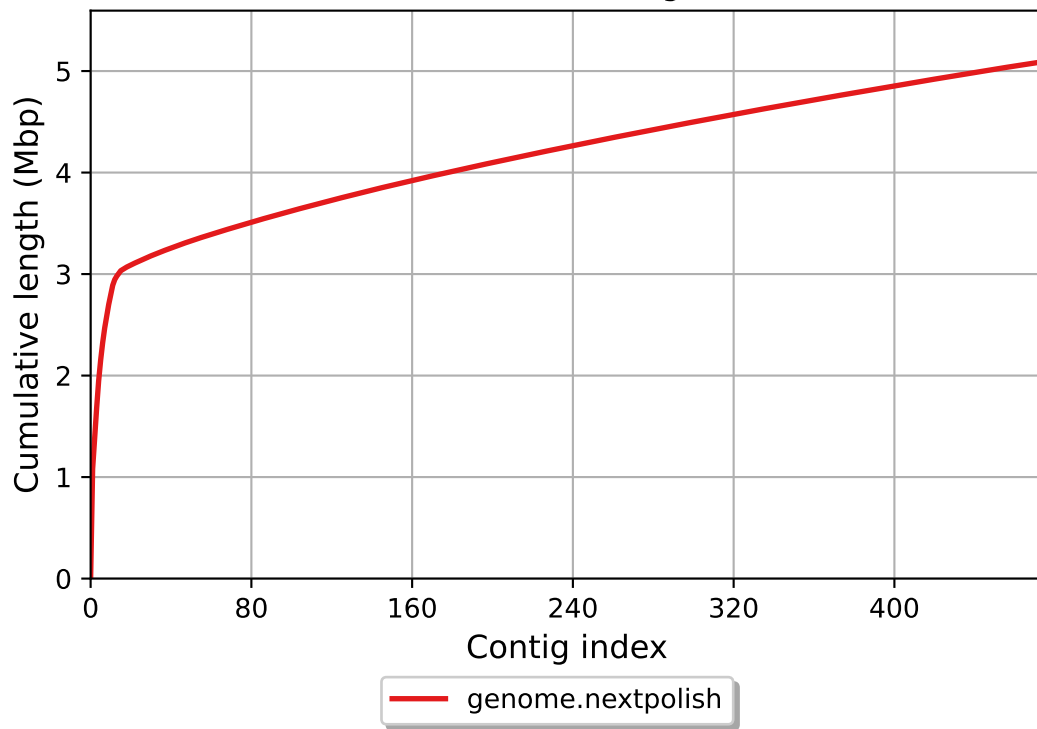
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

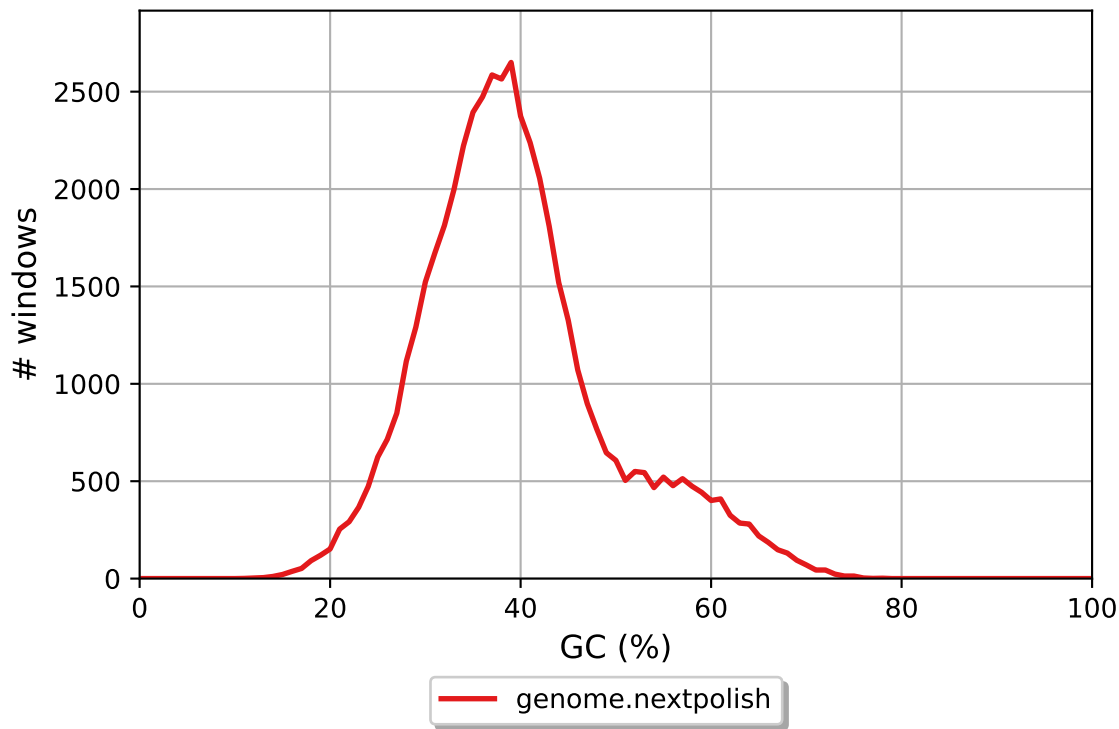


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

