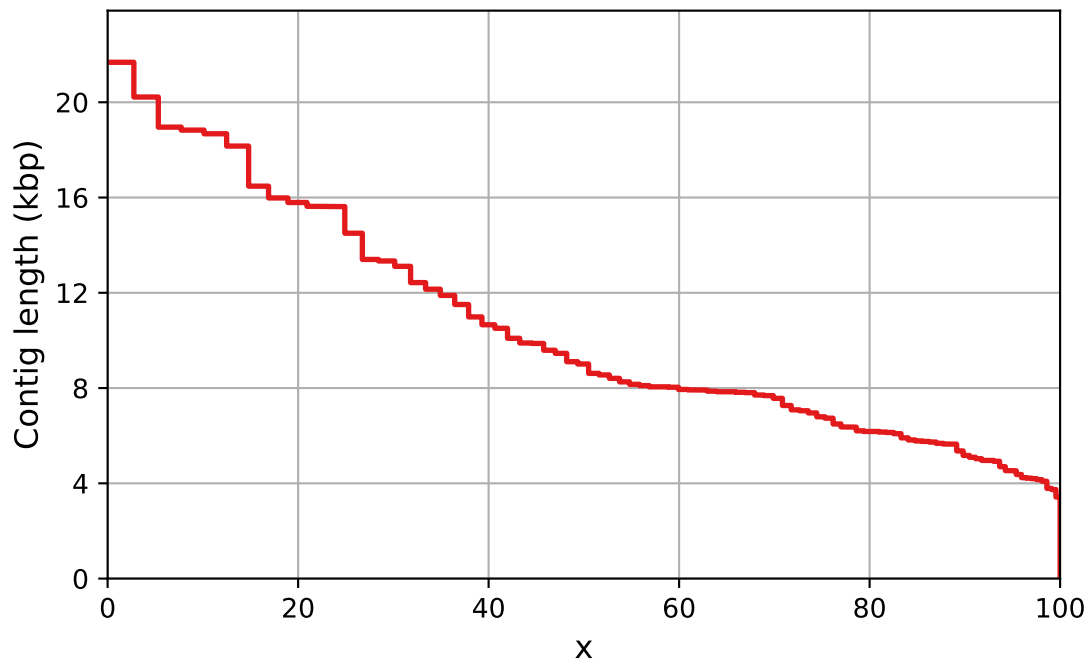


Report

	genome.nextpolish
# contigs (>= 0 bp)	91
# contigs (>= 1000 bp)	91
# contigs (>= 5000 bp)	76
# contigs (>= 10000 bp)	23
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	787162
Total length (>= 1000 bp)	787162
Total length (>= 5000 bp)	722342
Total length (>= 10000 bp)	340591
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	91
Largest contig	21679
Total length	787162
GC (%)	50.31
N50	9010
N90	5170
auN	10696.5
L50	29
L90	74
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 2 part

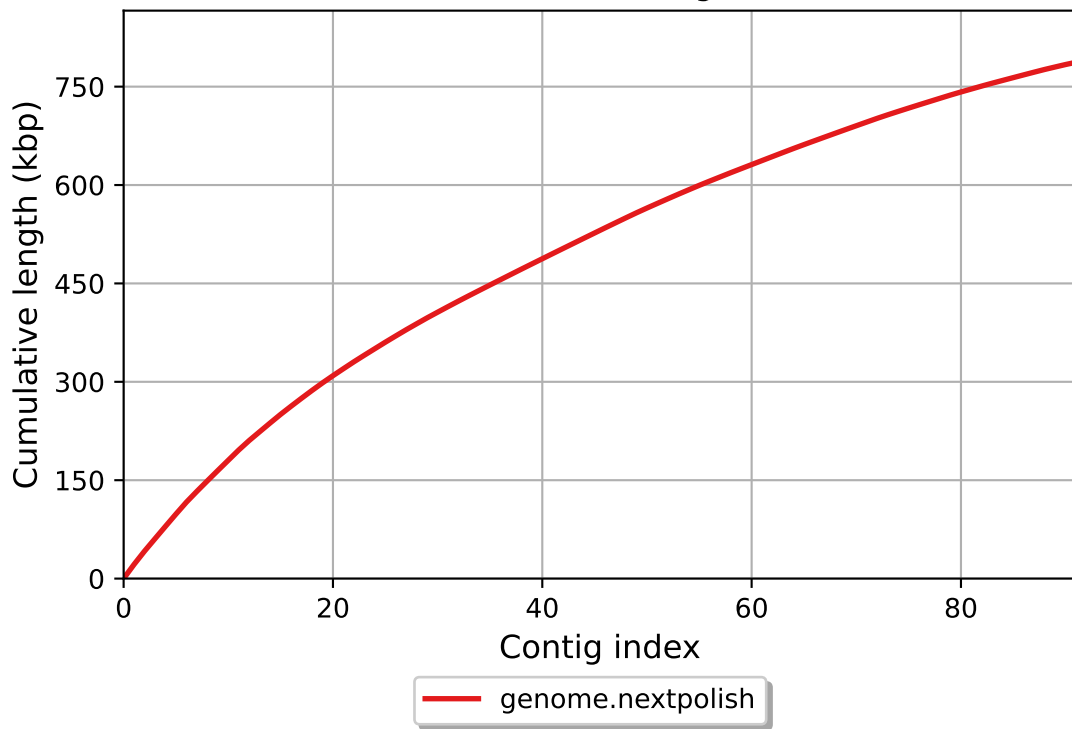
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

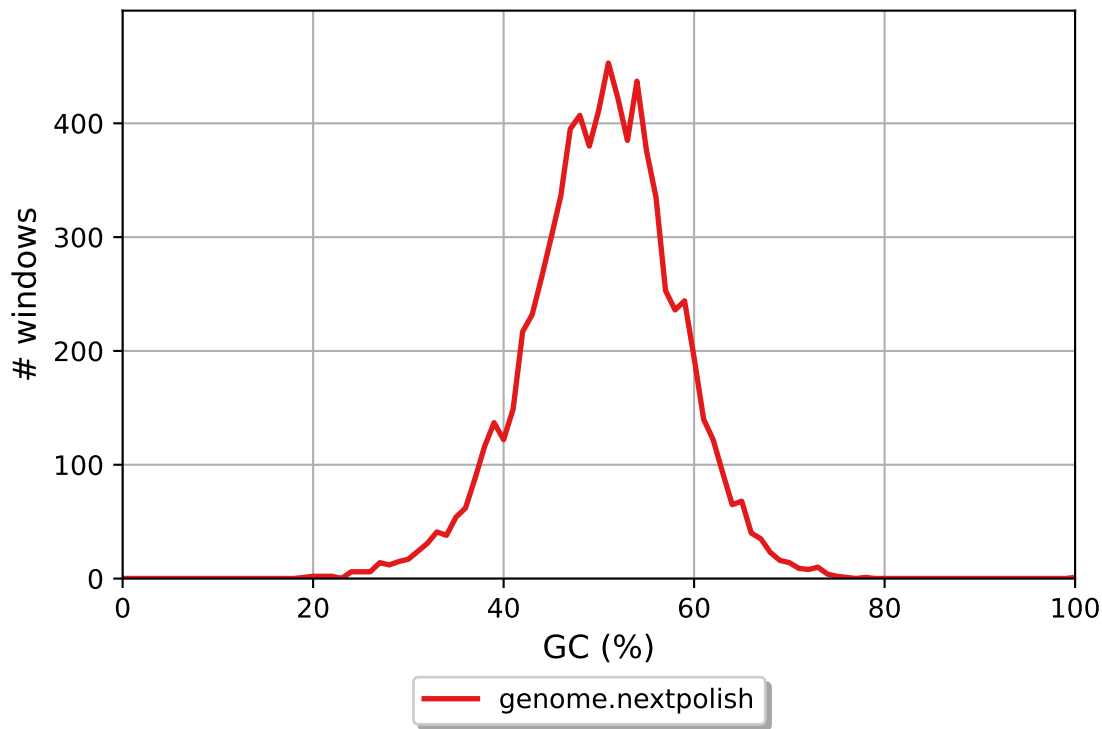


genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

