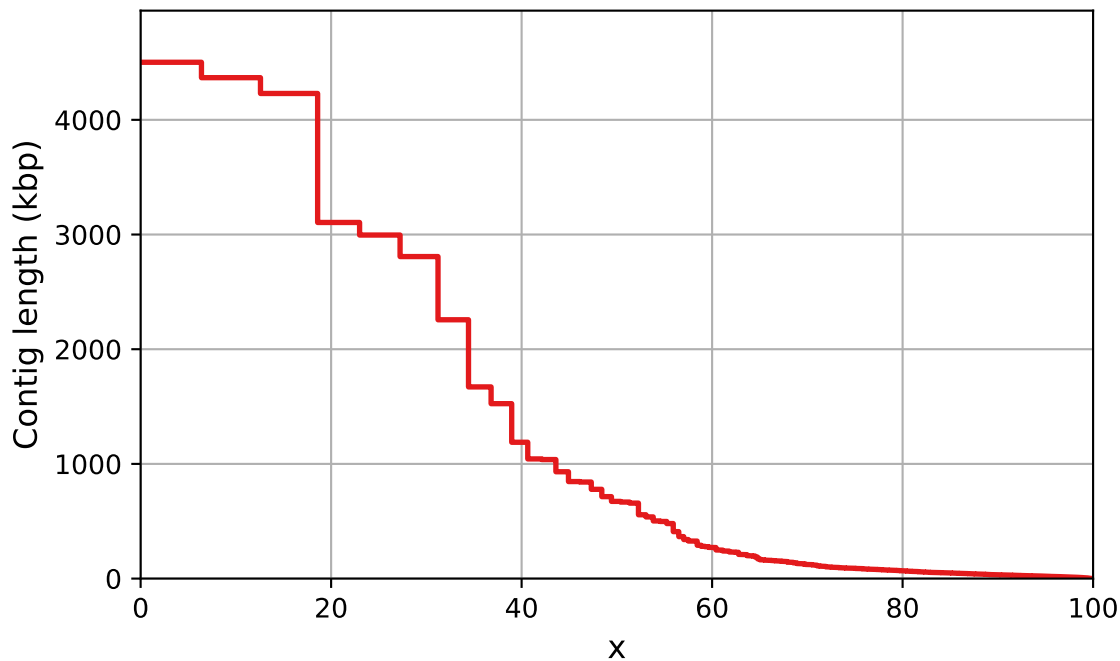


# Report

	genome.nextpolish
# contigs (>= 0 bp)	922
# contigs (>= 1000 bp)	833
# contigs (>= 5000 bp)	687
# contigs (>= 10000 bp)	603
# contigs (>= 25000 bp)	386
# contigs (>= 50000 bp)	210
Total length (>= 0 bp)	70489212
Total length (>= 1000 bp)	70428701
Total length (>= 5000 bp)	70130967
Total length (>= 10000 bp)	69491840
Total length (>= 25000 bp)	65814557
Total length (>= 50000 bp)	59616304
# contigs	915
Largest contig	4501950
Total length	70486327
GC (%)	45.86
N50	673142
N90	33171
auN	1517568.3
L50	18
L90	304
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	166 + 7 part

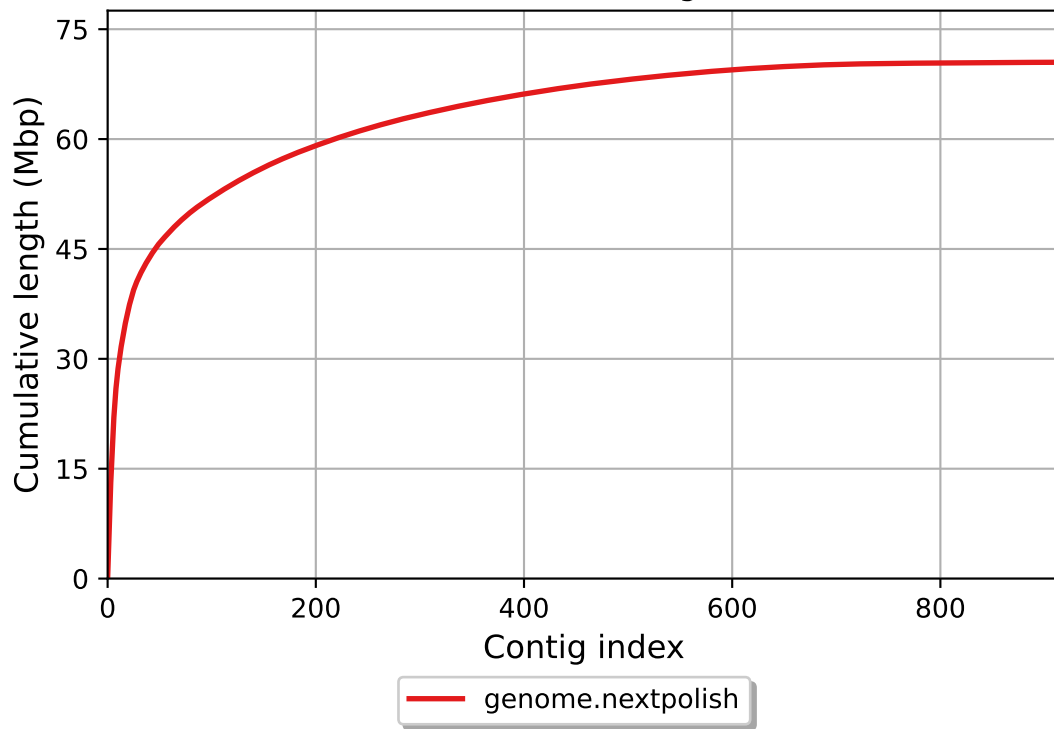
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

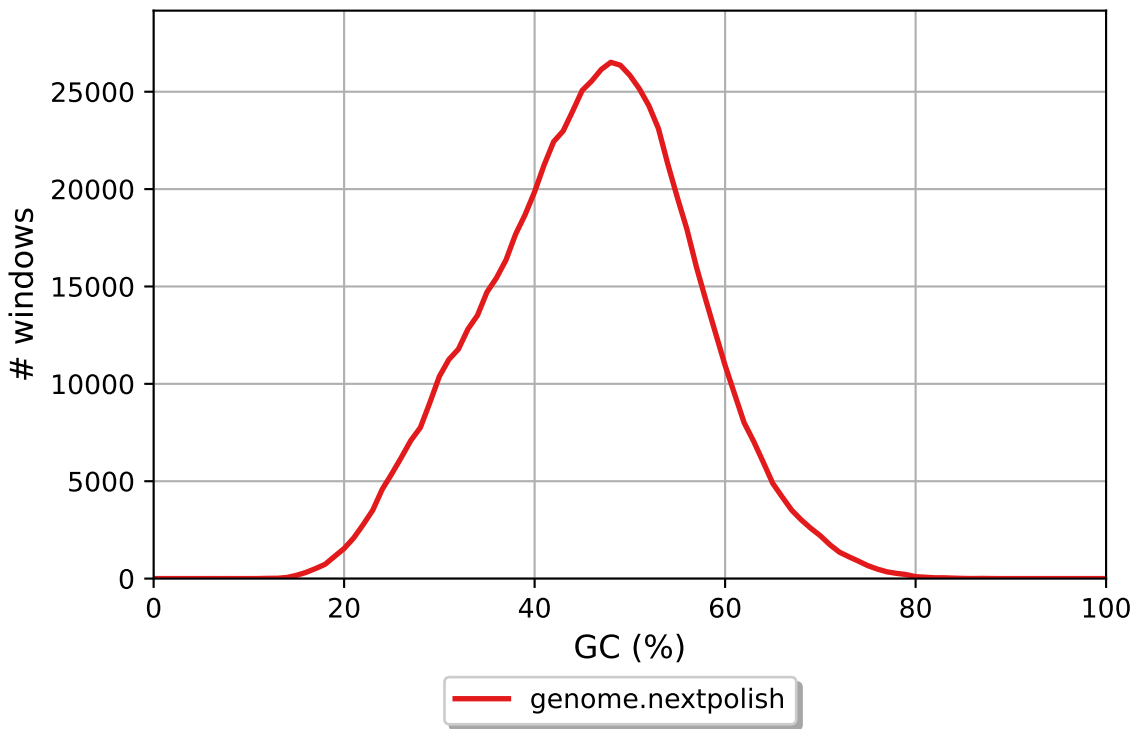


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

