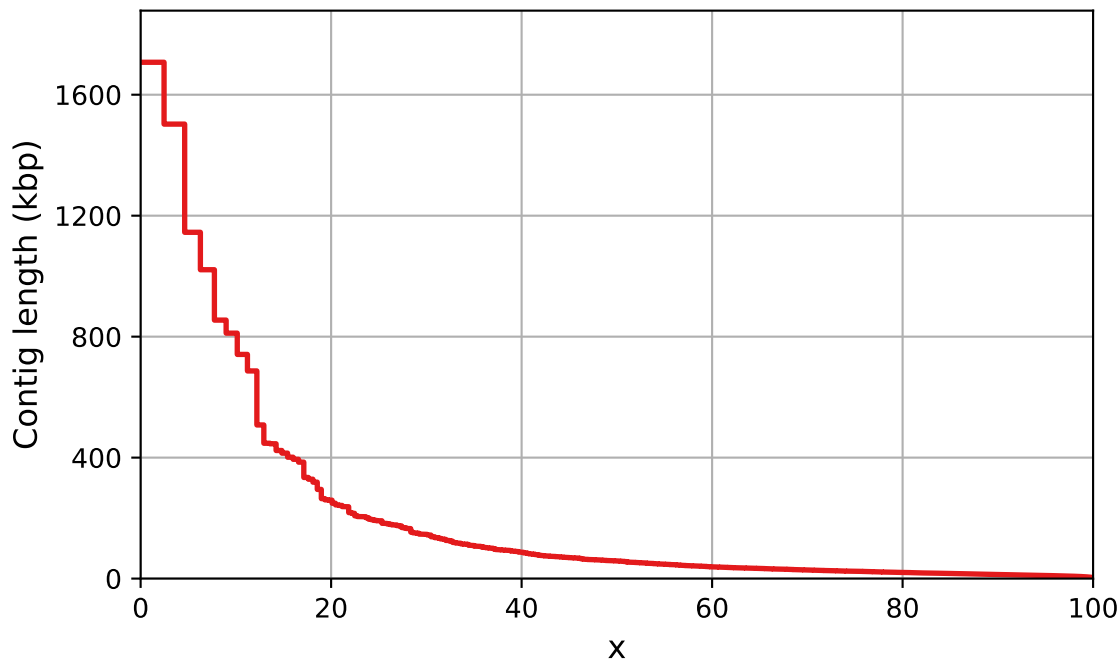


# Report

	genome.nextpolish
# contigs (>= 0 bp)	2060
# contigs (>= 1000 bp)	2053
# contigs (>= 5000 bp)	1990
# contigs (>= 10000 bp)	1604
# contigs (>= 25000 bp)	679
# contigs (>= 50000 bp)	272
Total length (>= 0 bp)	69407423
Total length (>= 1000 bp)	69402140
Total length (>= 5000 bp)	69152776
Total length (>= 10000 bp)	66173567
Total length (>= 25000 bp)	51223843
Total length (>= 50000 bp)	37278631
# contigs	2060
Largest contig	1707288
Total length	69407423
GC (%)	49.06
N50	58167
N90	13433
auN	223583.9
L50	225
L90	1286
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	169 + 13 part

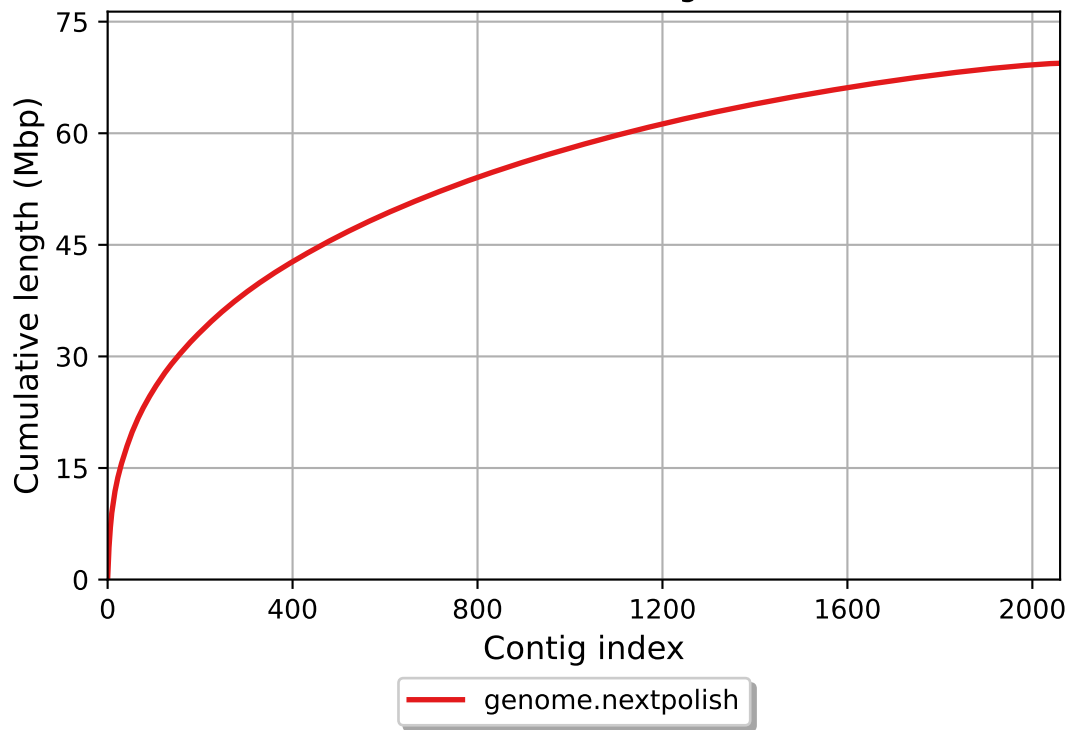
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

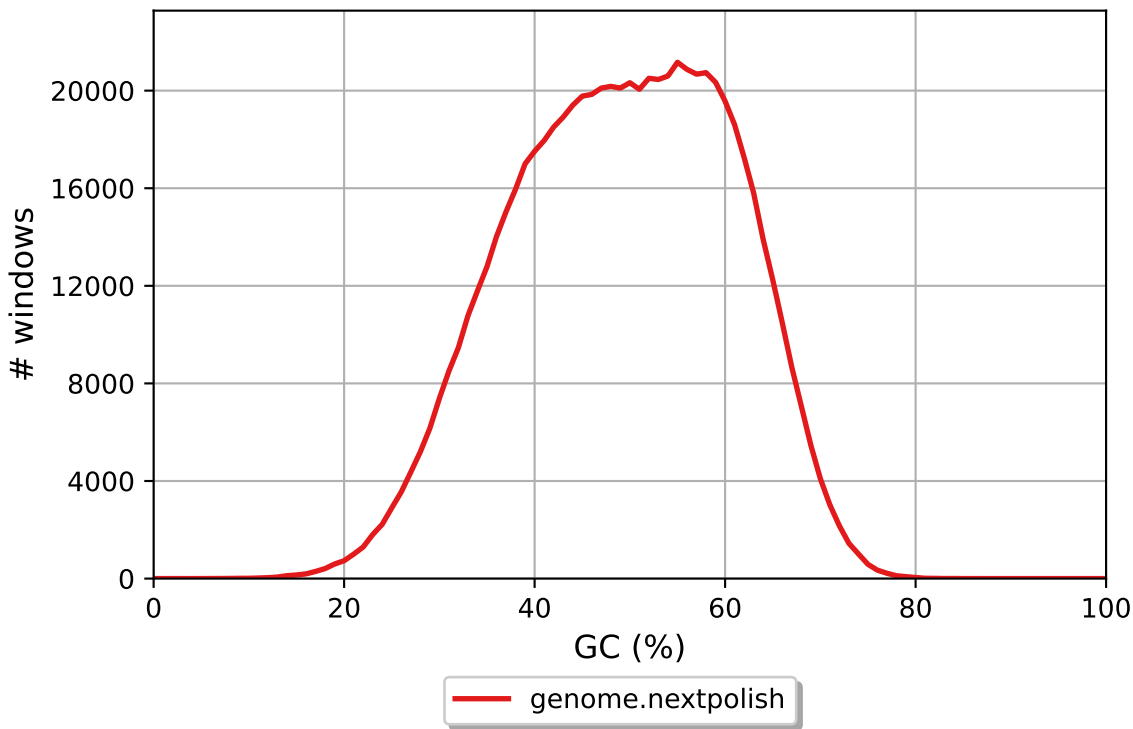


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

