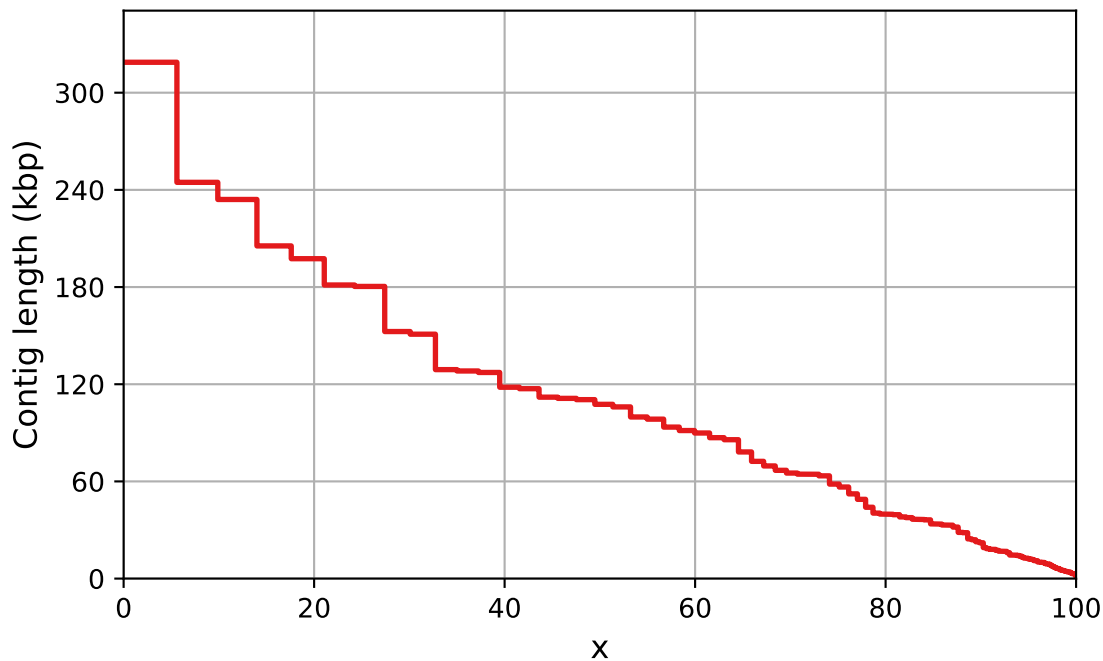


Report

	genome.nextpolish
# contigs (>= 0 bp)	123
# contigs (>= 1000 bp)	122
# contigs (>= 5000 bp)	101
# contigs (>= 10000 bp)	84
# contigs (>= 25000 bp)	55
# contigs (>= 50000 bp)	37
Total length (>= 0 bp)	5698410
Total length (>= 1000 bp)	5697833
Total length (>= 5000 bp)	5621439
Total length (>= 10000 bp)	5499120
Total length (>= 25000 bp)	5049726
Total length (>= 50000 bp)	4390138
# contigs	123
Largest contig	318785
Total length	5698410
GC (%)	59.31
N50	107590
N90	22113
auN	120283.8
L50	18
L90	59
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	18 + 9 part

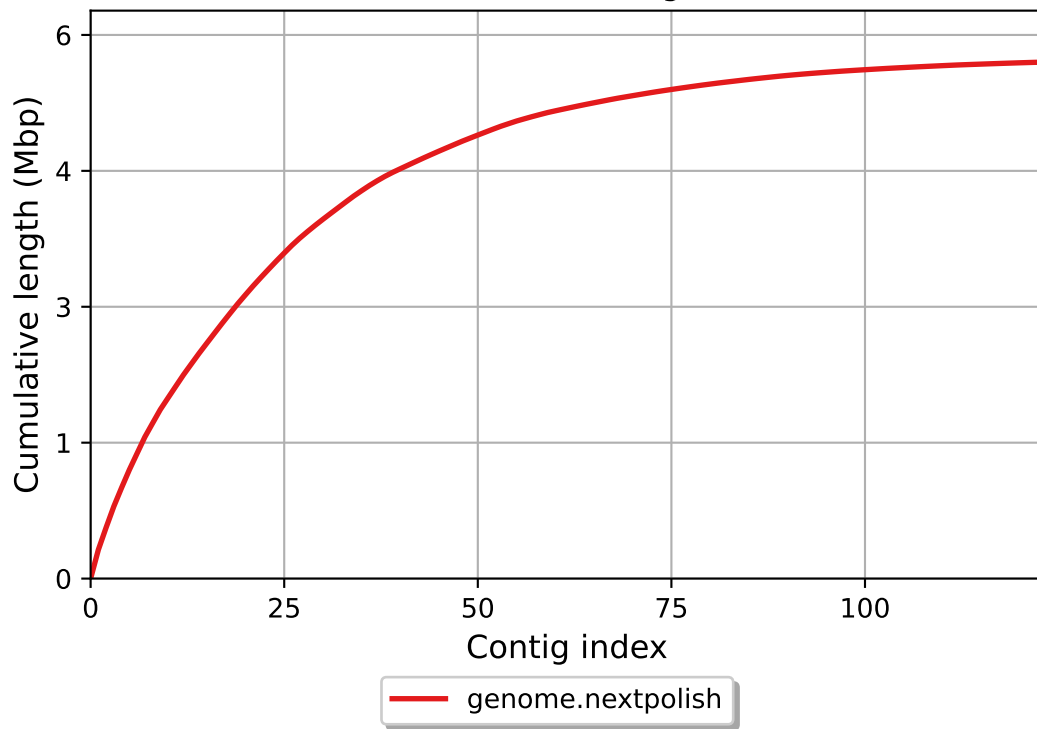
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

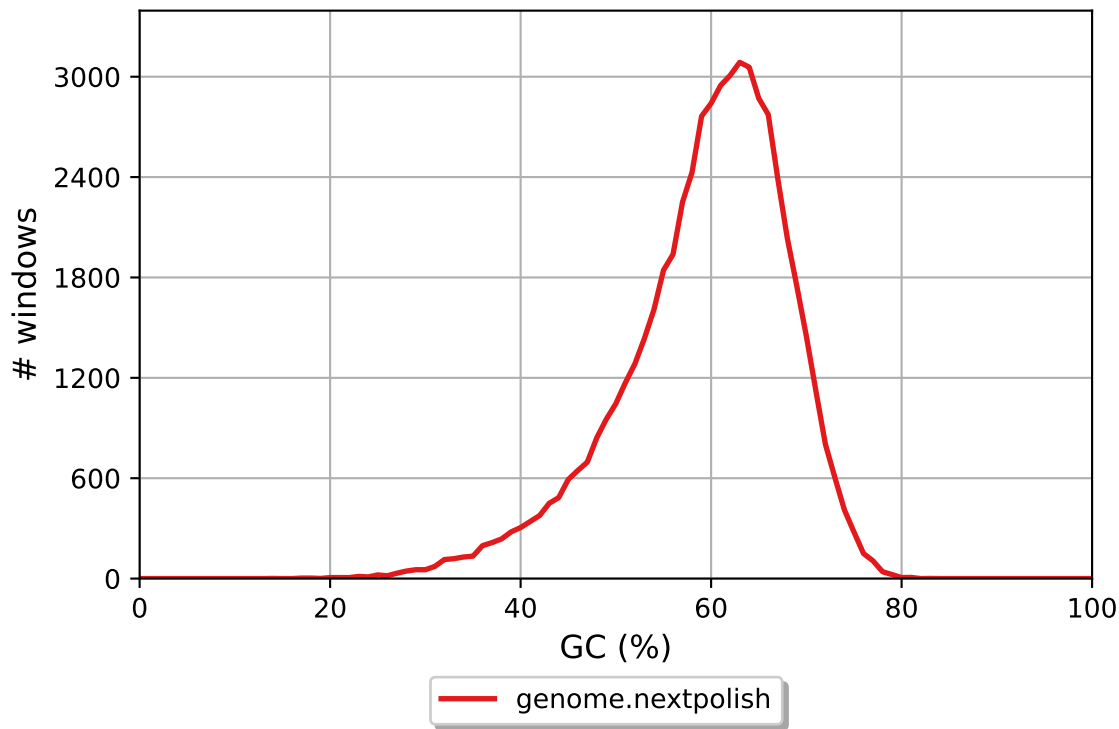


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

