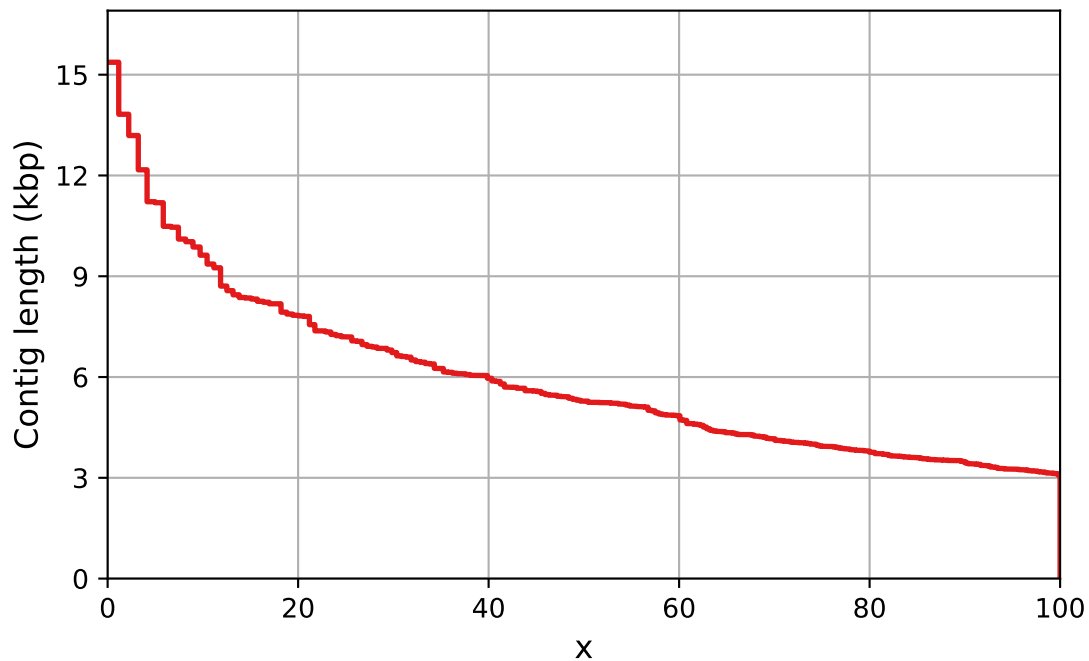


Report

	genome.nextpolish
# contigs (>= 0 bp)	256
# contigs (>= 1000 bp)	256
# contigs (>= 5000 bp)	108
# contigs (>= 10000 bp)	10
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1317006
Total length (>= 1000 bp)	1317006
Total length (>= 5000 bp)	752196
Total length (>= 10000 bp)	118040
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	256
Largest contig	15370
Total length	1317006
GC (%)	49.79
N50	5283
N90	3477
auN	5967.6
L50	90
L90	216
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 1 part

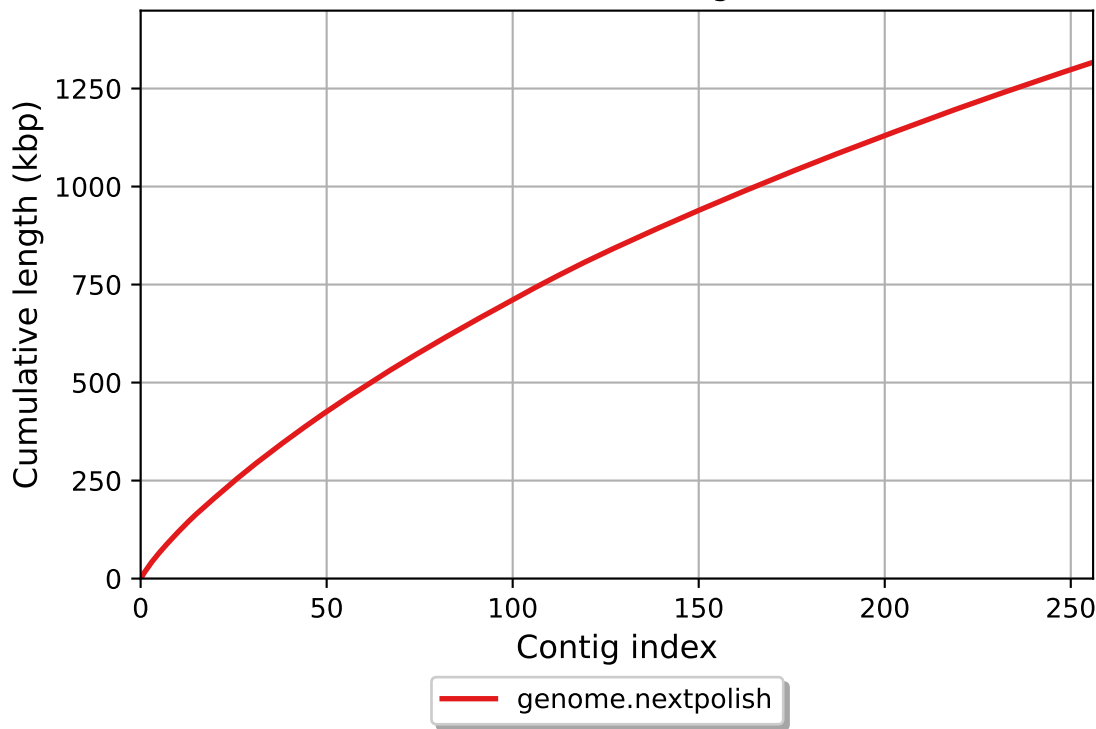
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

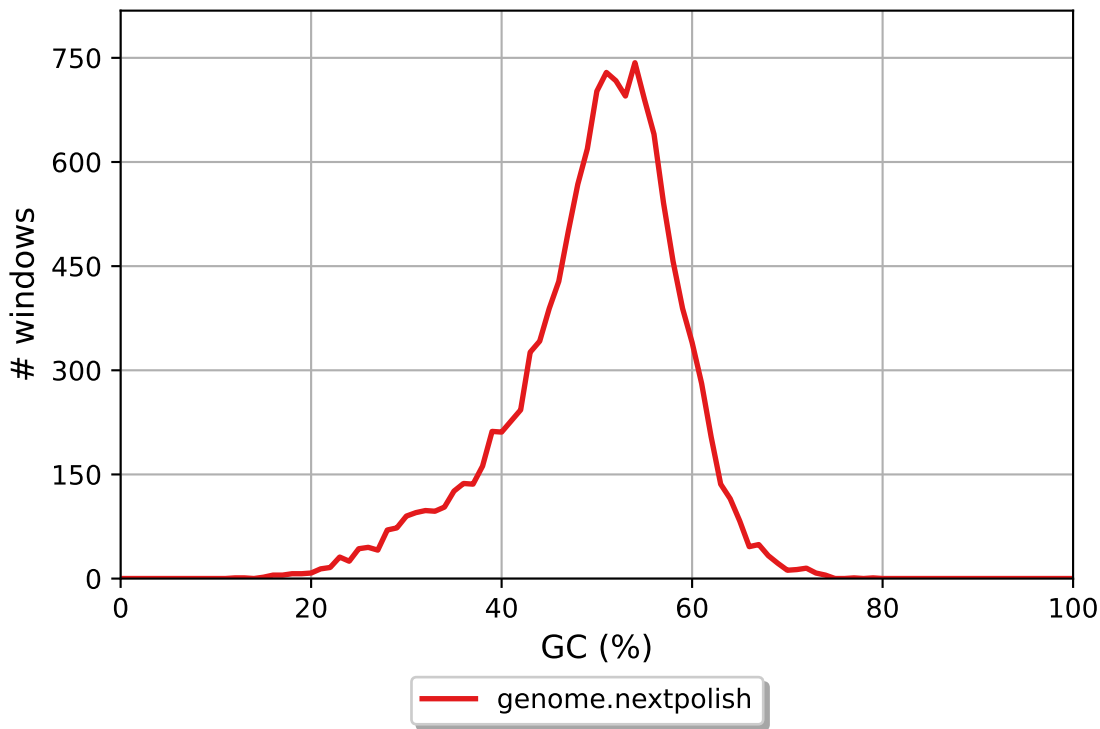


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

