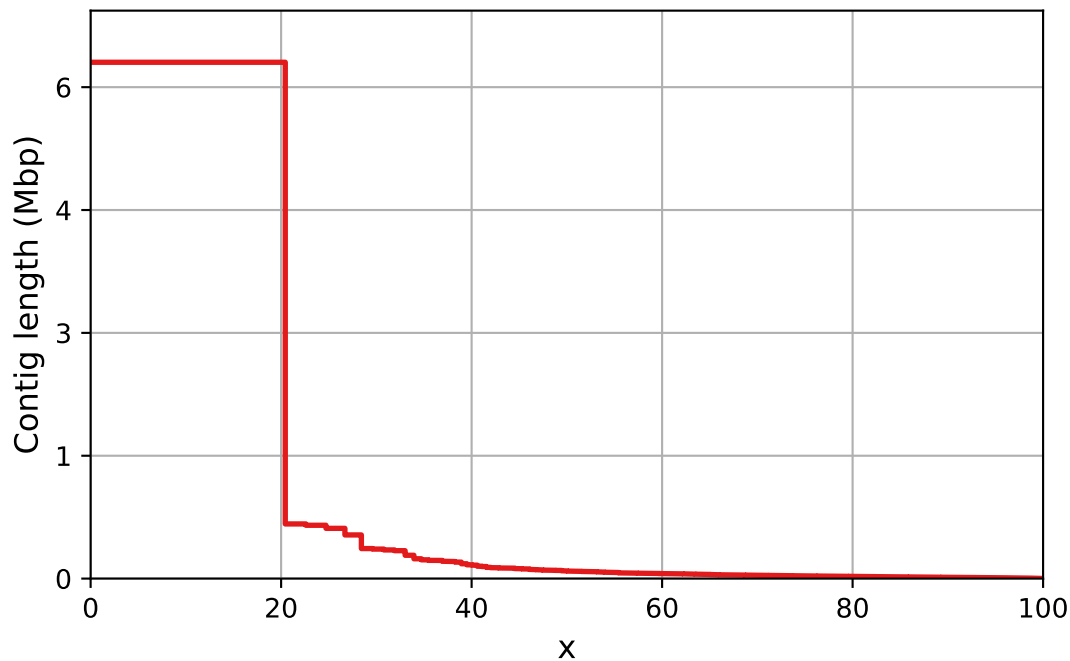


Report

	genome.nextpolish
# contigs (>= 0 bp)	766
# contigs (>= 1000 bp)	762
# contigs (>= 5000 bp)	671
# contigs (>= 10000 bp)	517
# contigs (>= 25000 bp)	260
# contigs (>= 50000 bp)	114
Total length (>= 0 bp)	30847384
Total length (>= 1000 bp)	30844571
Total length (>= 5000 bp)	30516515
Total length (>= 10000 bp)	29364924
Total length (>= 25000 bp)	25127091
Total length (>= 50000 bp)	20056116
# contigs	766
Largest contig	6303985
Total length	30847384
GC (%)	53.65
N50	92499
N90	15871
auN	1400859.2
L50	45
L90	393
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	75 + 1 part

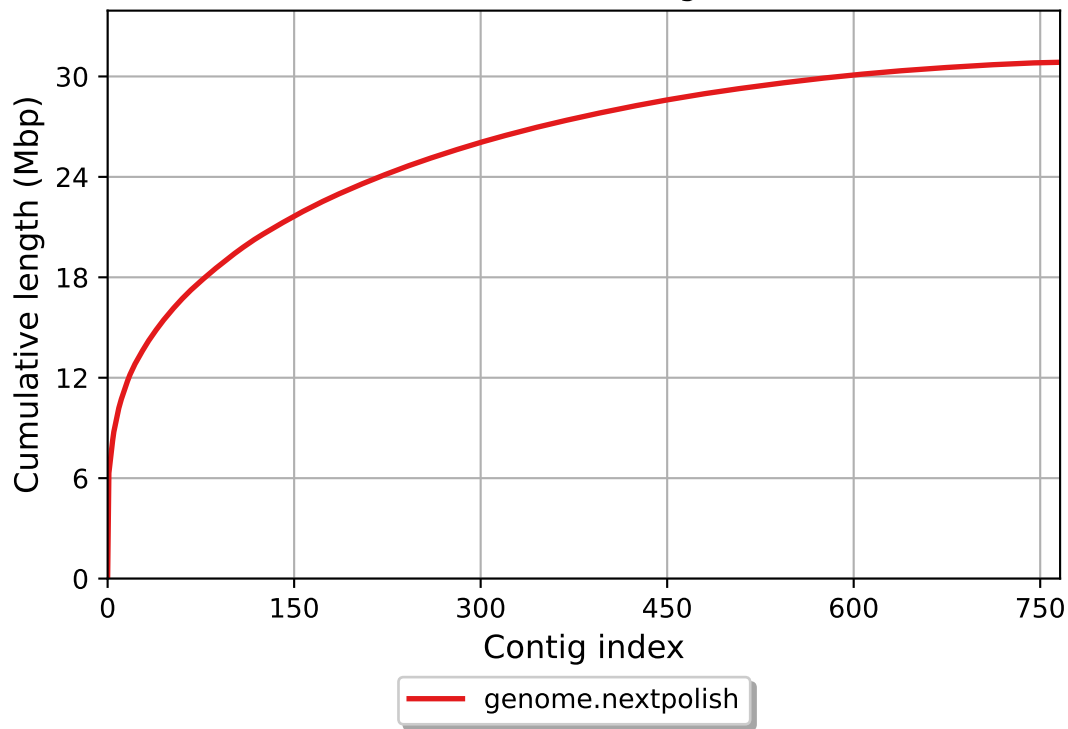
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

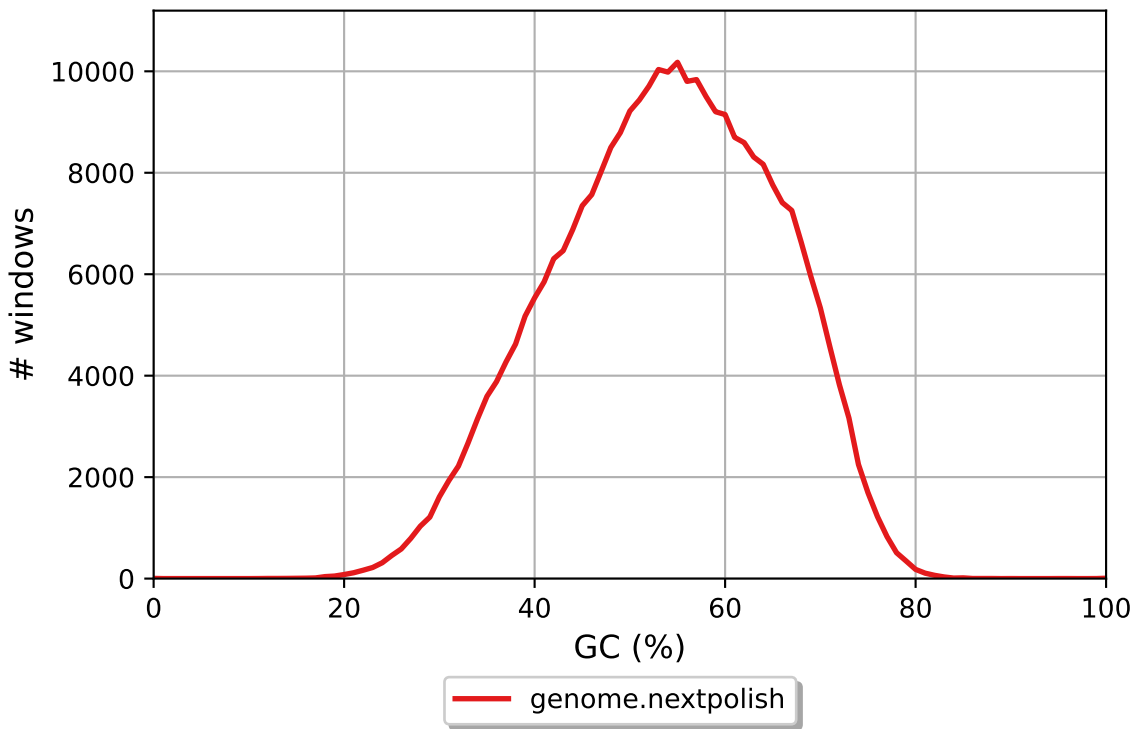


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

