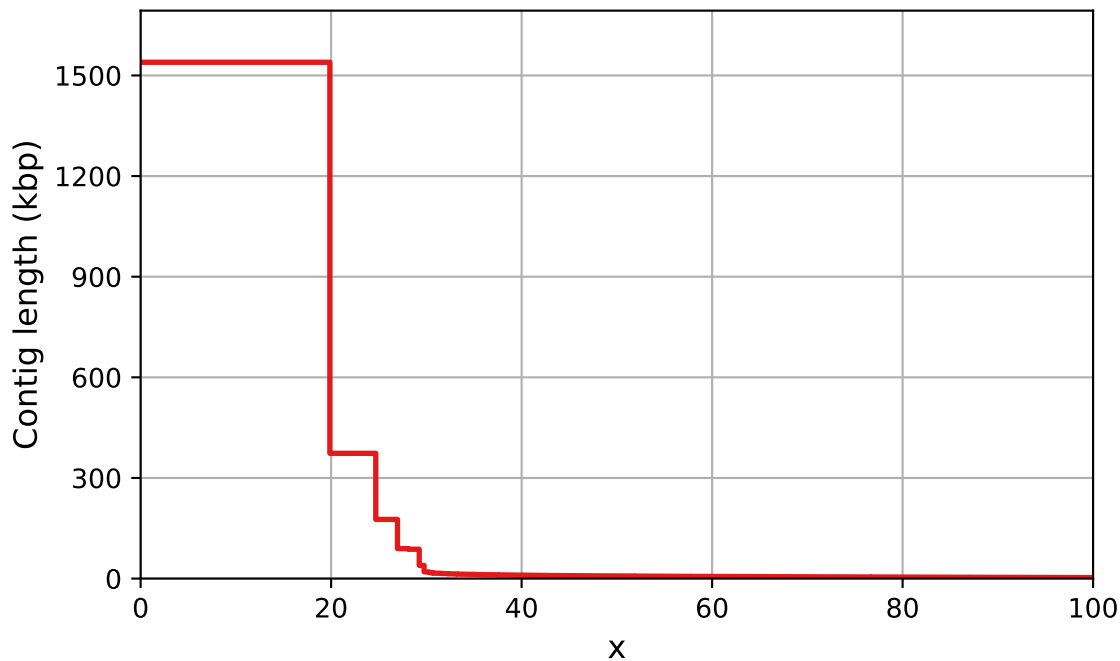


Report

	genome.nextpolish
# contigs (>= 0 bp)	976
# contigs (>= 1000 bp)	976
# contigs (>= 5000 bp)	471
# contigs (>= 10000 bp)	73
# contigs (>= 25000 bp)	6
# contigs (>= 50000 bp)	5
Total length (>= 0 bp)	7747277
Total length (>= 1000 bp)	7747277
Total length (>= 5000 bp)	5790914
Total length (>= 10000 bp)	3145492
Total length (>= 25000 bp)	2305530
Total length (>= 50000 bp)	2266179
# contigs	976
Largest contig	1539310
Total length	7747277
GC (%)	36.45
N50	7833
N90	3695
auN	334850.4
L50	157
L90	748
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 6 part

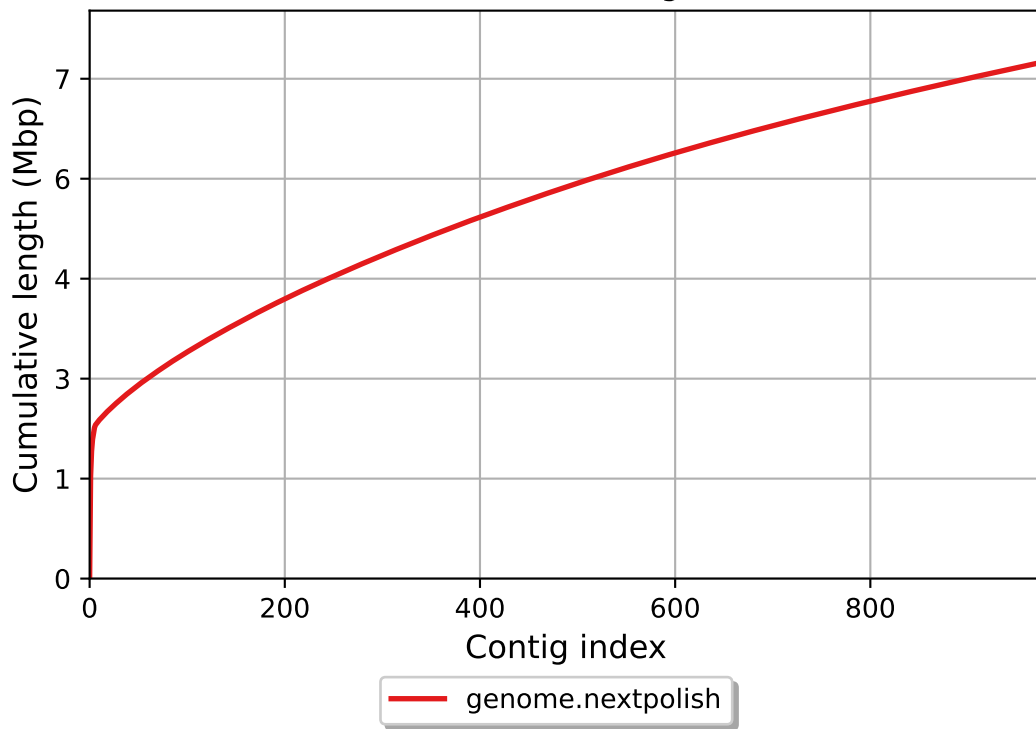
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

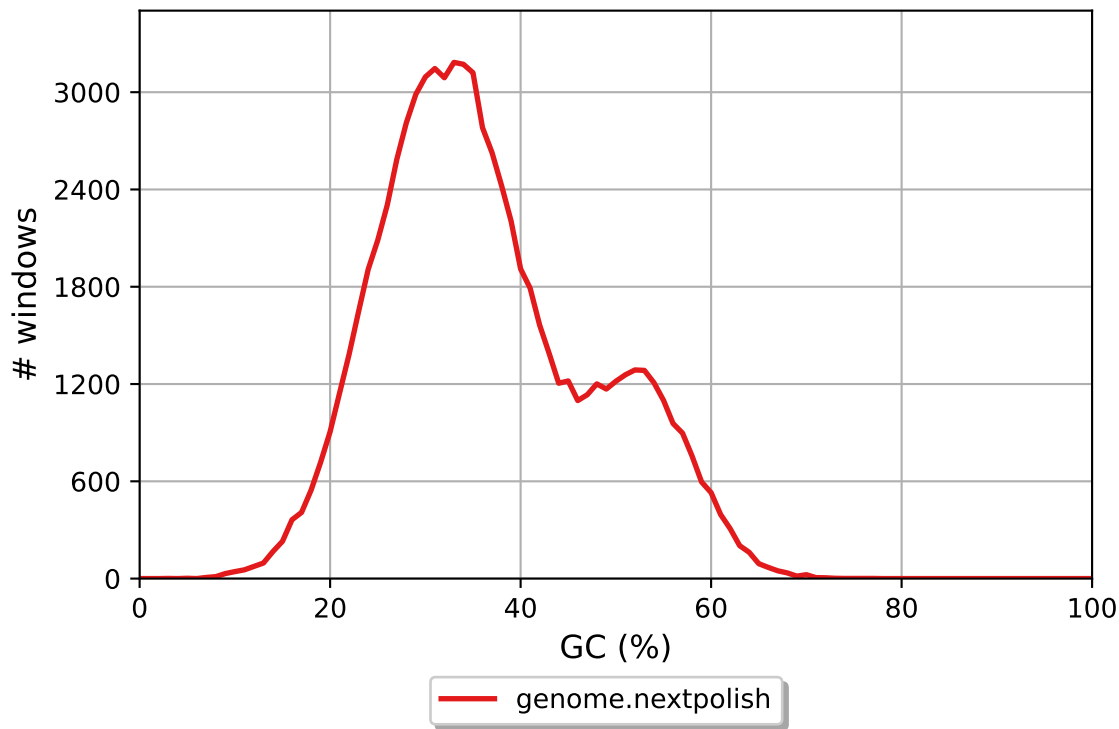


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

