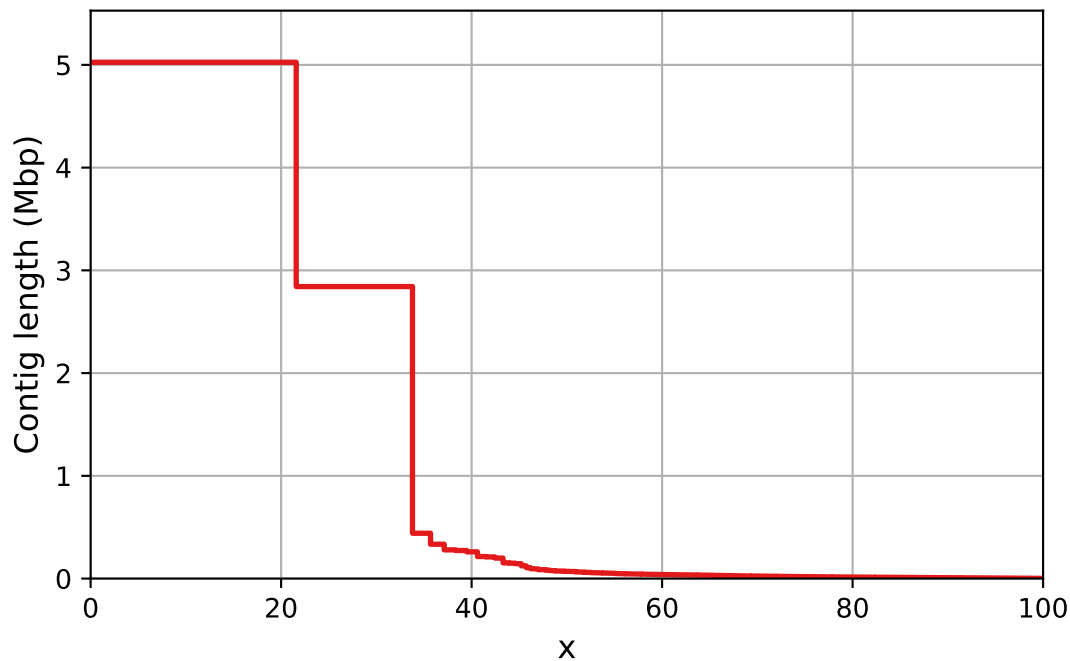


Report

	genome.nextpolish
# contigs (>= 0 bp)	966
# contigs (>= 1000 bp)	961
# contigs (>= 5000 bp)	689
# contigs (>= 10000 bp)	410
# contigs (>= 25000 bp)	144
# contigs (>= 50000 bp)	46
Total length (>= 0 bp)	23282398
Total length (>= 1000 bp)	23278920
Total length (>= 5000 bp)	22379790
Total length (>= 10000 bp)	20367854
Total length (>= 25000 bp)	16247086
Total length (>= 50000 bp)	12829745
# contigs	966
Largest contig	5025286
Total length	23282398
GC (%)	47.13
N50	70418
N90	8442
auN	1479060.2
L50	26
L90	474
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	86 + 6 part

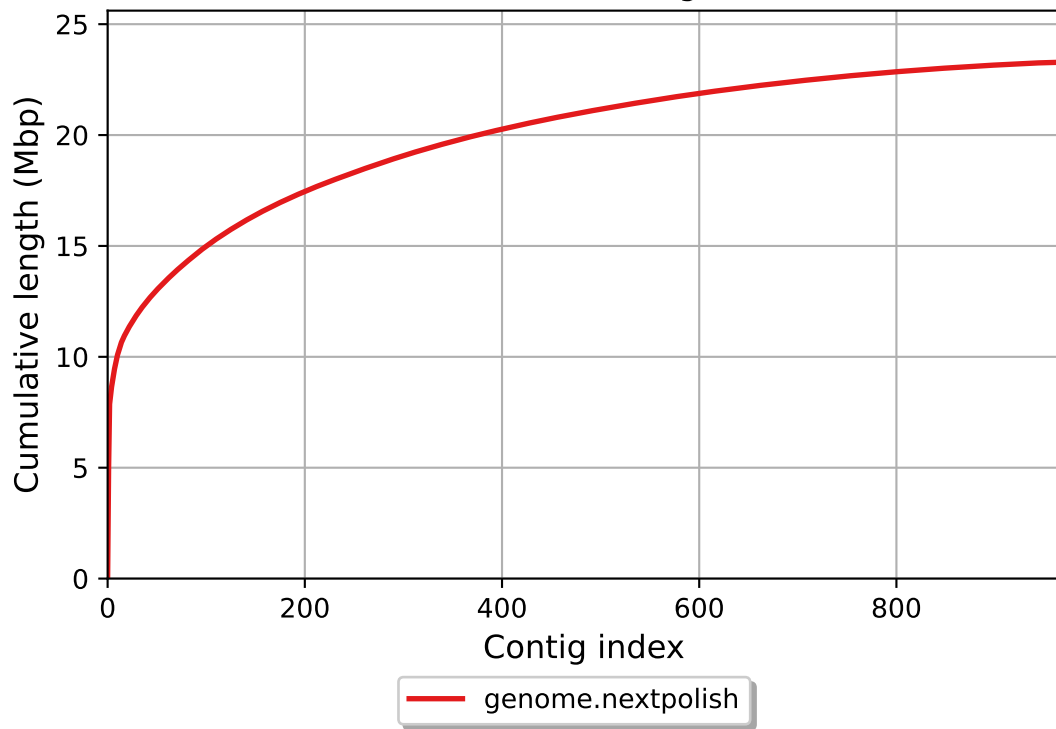
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

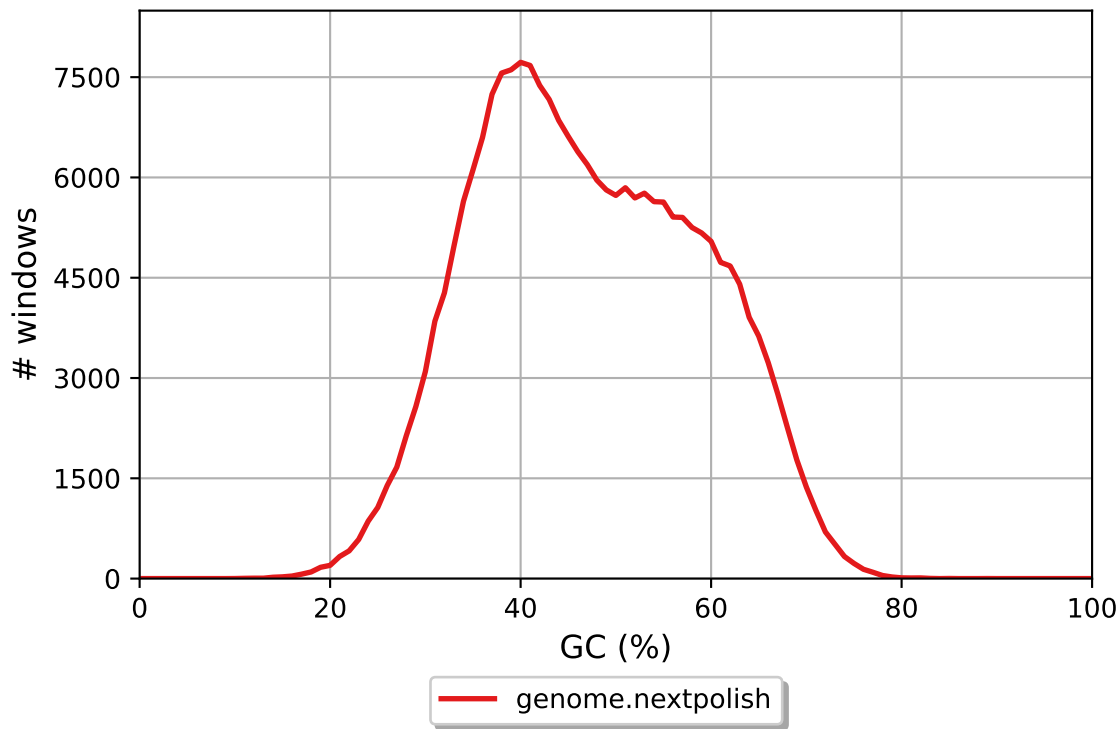


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

