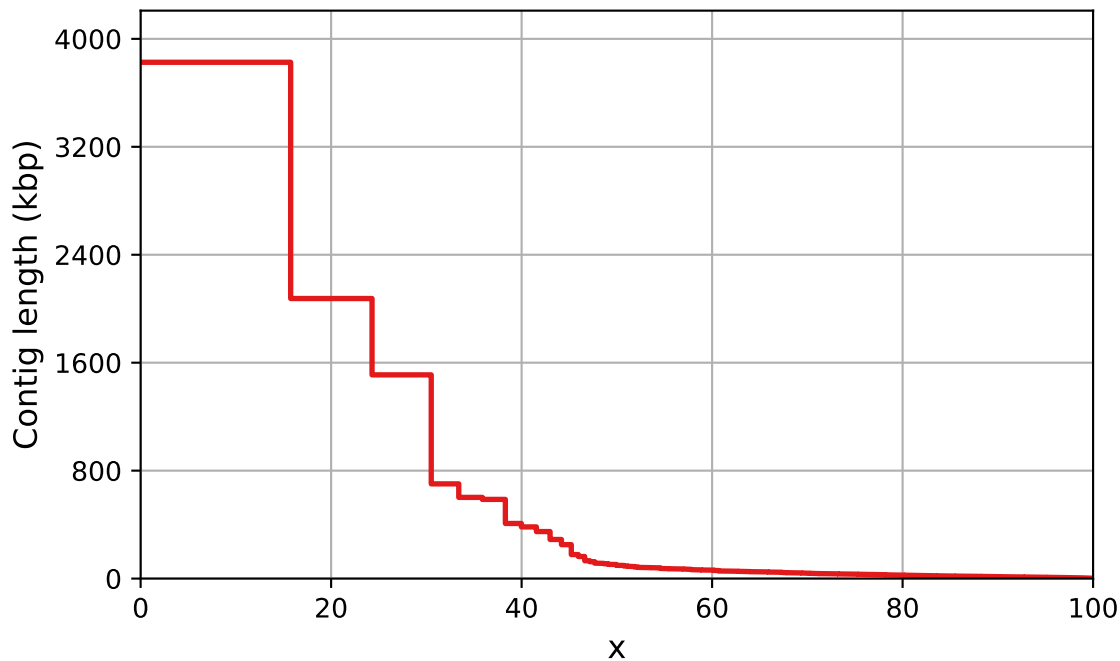


Report

	genome.nextpolish
# contigs (>= 0 bp)	611
# contigs (>= 1000 bp)	608
# contigs (>= 5000 bp)	541
# contigs (>= 10000 bp)	401
# contigs (>= 25000 bp)	180
# contigs (>= 50000 bp)	77
Total length (>= 0 bp)	24292398
Total length (>= 1000 bp)	24290047
Total length (>= 5000 bp)	24052121
Total length (>= 10000 bp)	23007175
Total length (>= 25000 bp)	19492821
Total length (>= 50000 bp)	15907437
# contigs	611
Largest contig	3826473
Total length	24292398
GC (%)	57.43
N50	98201
N90	14424
auN	972390.0
L50	21
L90	306
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	49 + 3 part

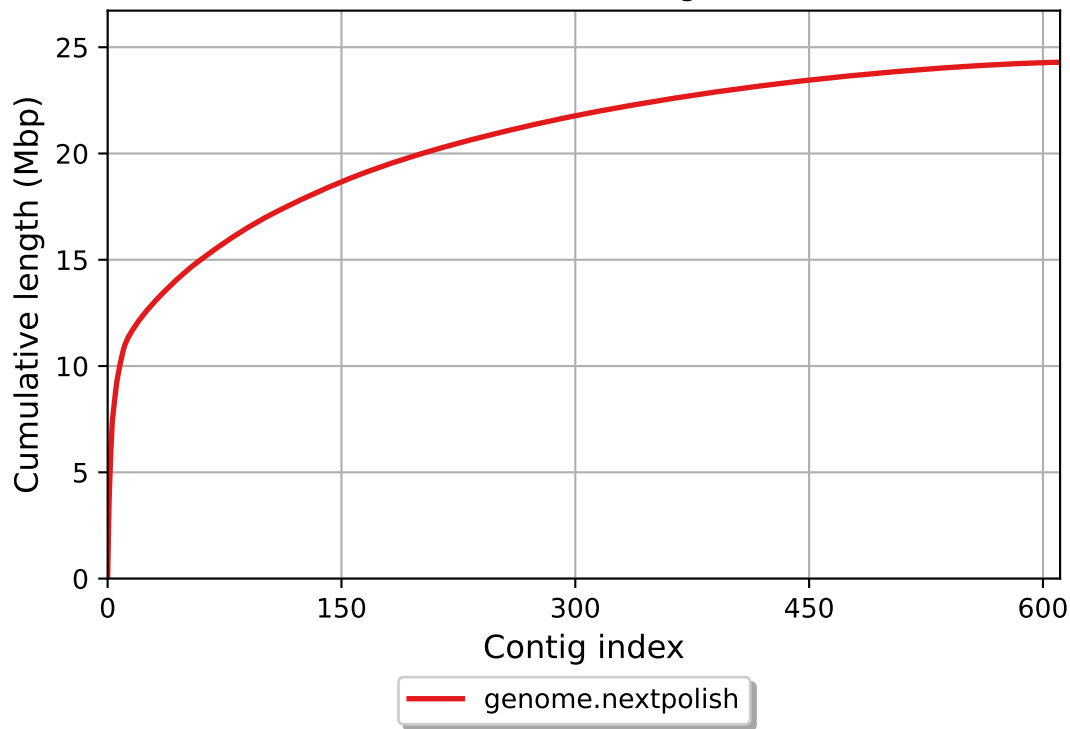
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

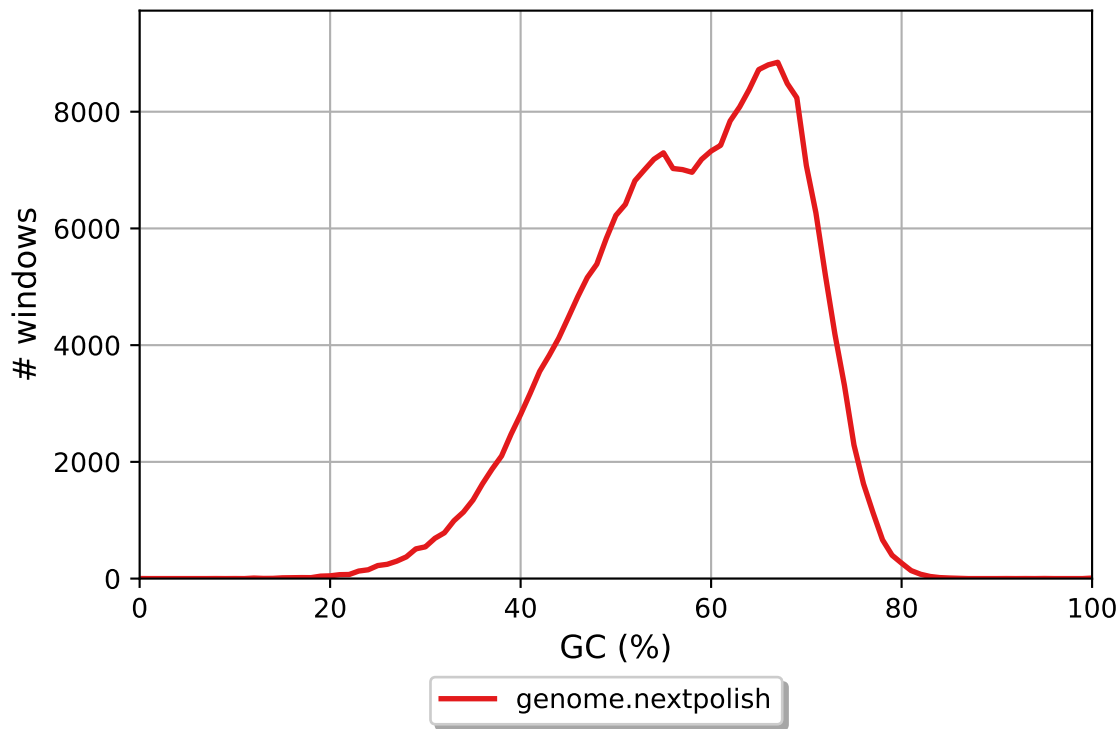


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

