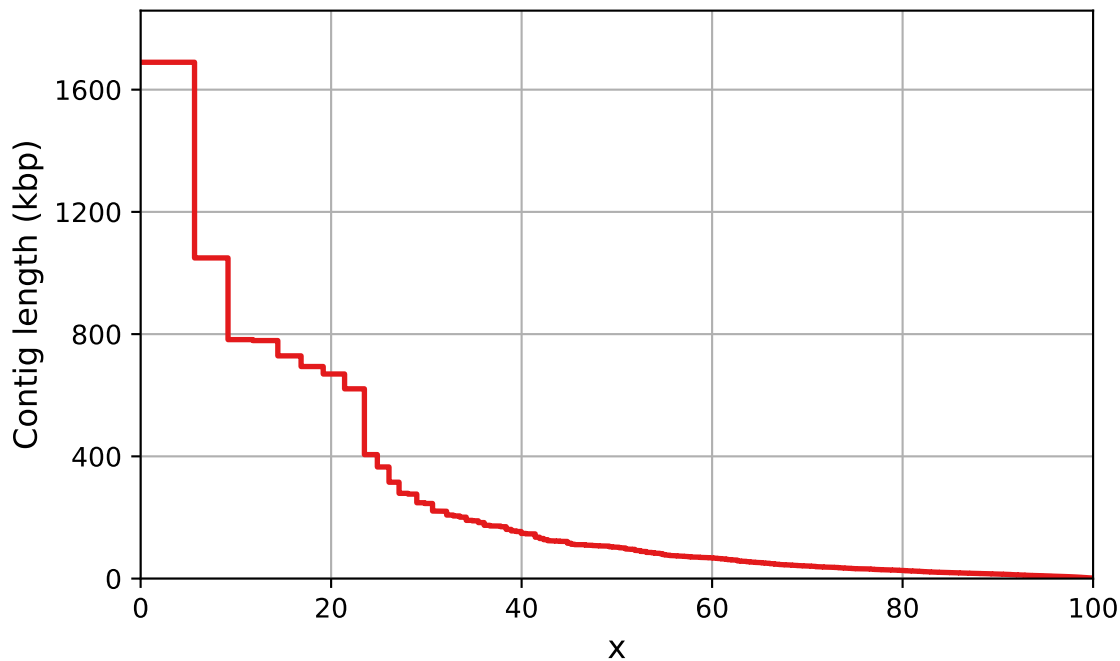


Report

	genome.nextpolish
# contigs (>= 0 bp)	841
# contigs (>= 1000 bp)	833
# contigs (>= 5000 bp)	667
# contigs (>= 10000 bp)	490
# contigs (>= 25000 bp)	251
# contigs (>= 50000 bp)	123
Total length (>= 0 bp)	29851371
Total length (>= 1000 bp)	29846138
Total length (>= 5000 bp)	29342434
Total length (>= 10000 bp)	28021399
Total length (>= 25000 bp)	24125677
Total length (>= 50000 bp)	19671209
# contigs	840
Largest contig	1689625
Total length	29850969
GC (%)	50.38
N50	102542
N90	14946
auN	306909.2
L50	56
L90	396
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	63 + 6 part

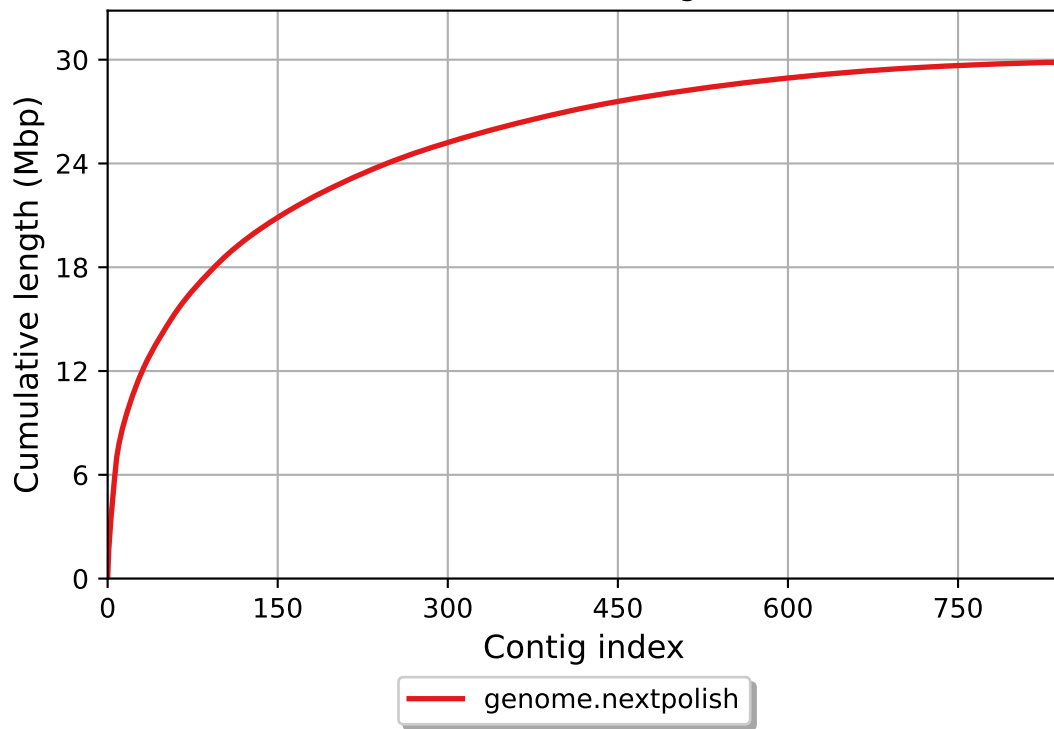
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

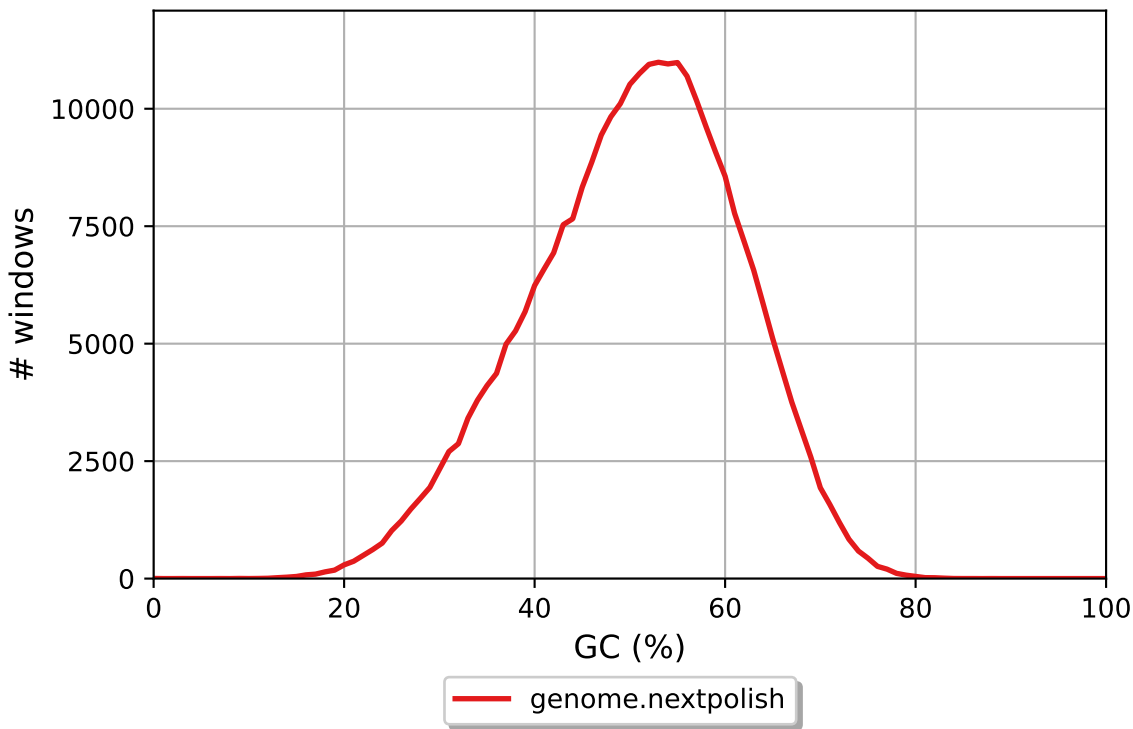


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

