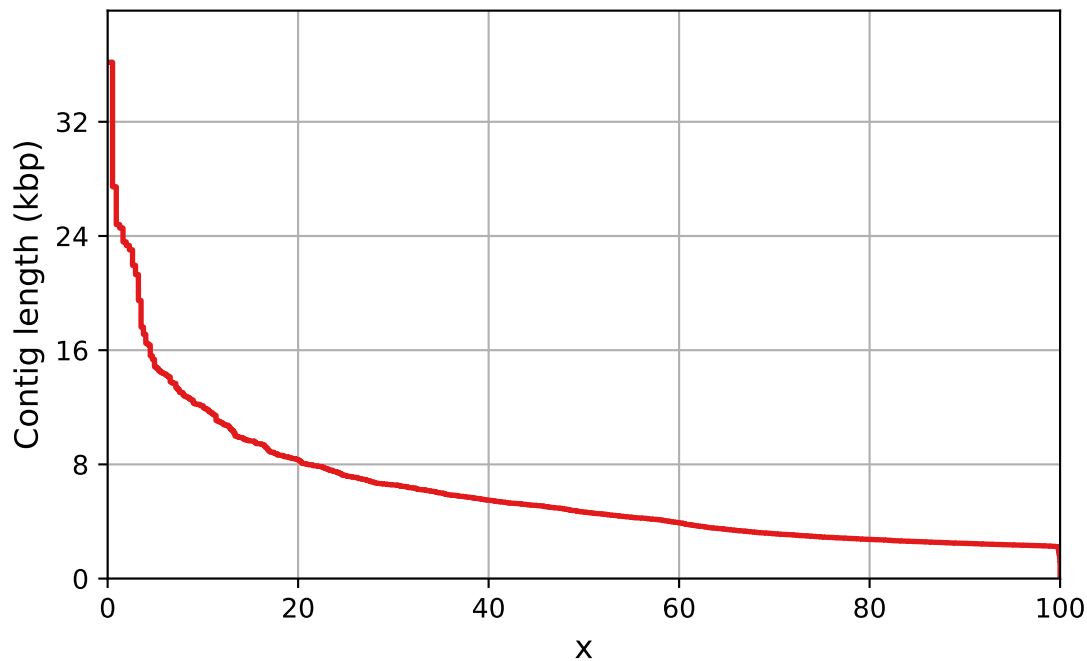


Report

	genome.nextpolish
# contigs (>= 0 bp)	1639
# contigs (>= 1000 bp)	1639
# contigs (>= 5000 bp)	413
# contigs (>= 10000 bp)	64
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	6996834
Total length (>= 1000 bp)	6996834
Total length (>= 5000 bp)	3244558
Total length (>= 10000 bp)	935833
Total length (>= 25000 bp)	63618
Total length (>= 50000 bp)	0
# contigs	1639
Largest contig	36169
Total length	6996834
GC (%)	39.18
N50	4657
N90	2468
auN	6186.9
L50	466
L90	1340
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	22 + 5 part

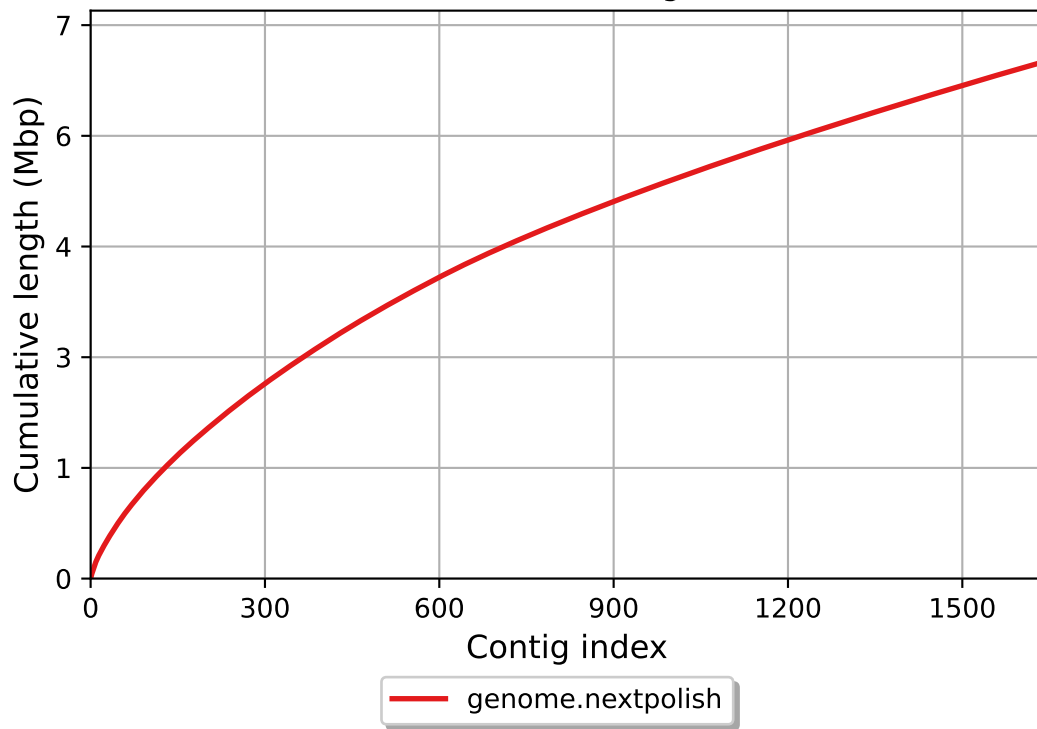
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

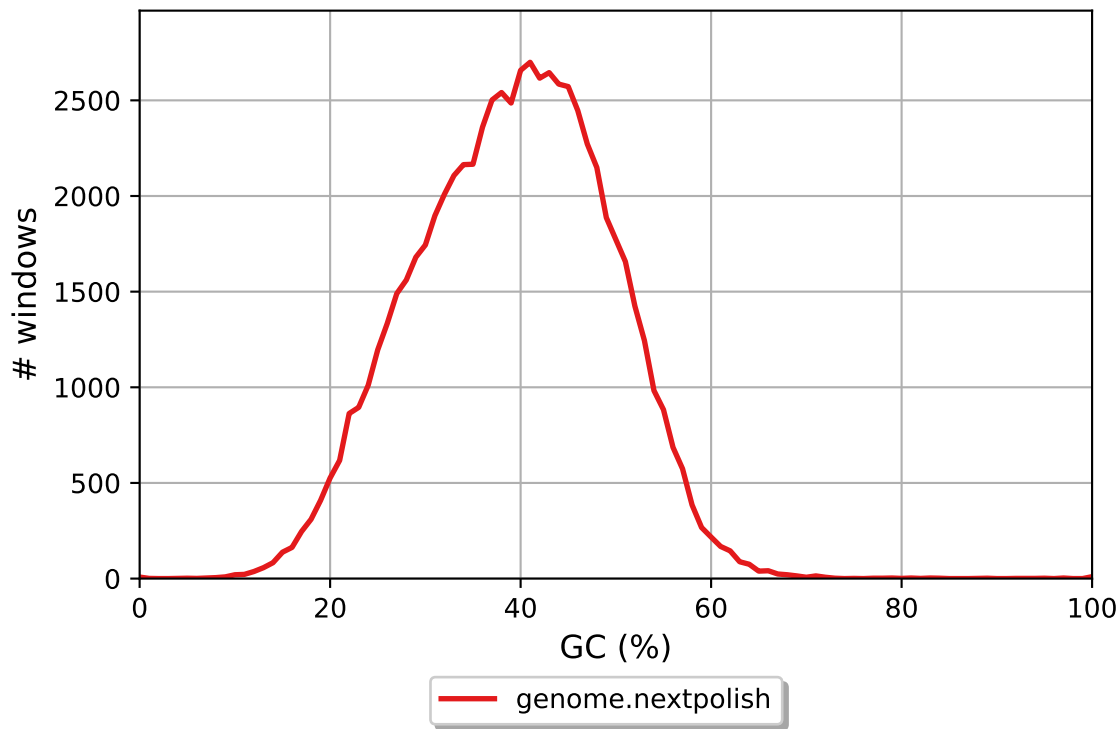


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

