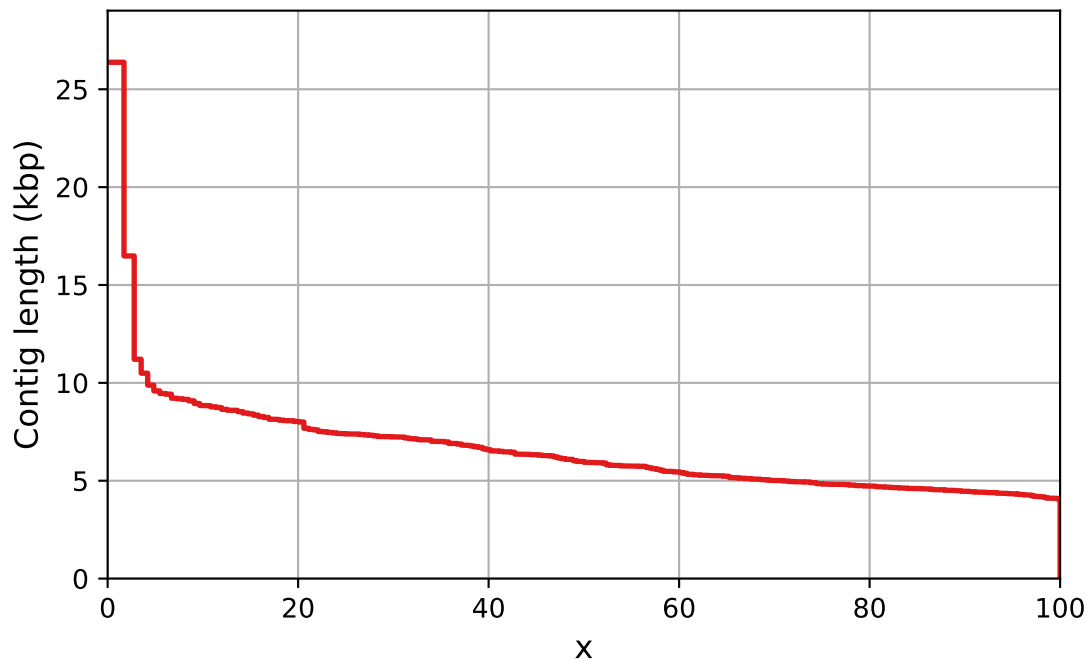


Report

	genome.nextpolish
# contigs (>= 0 bp)	256
# contigs (>= 1000 bp)	256
# contigs (>= 5000 bp)	158
# contigs (>= 10000 bp)	4
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1535387
Total length (>= 1000 bp)	1535387
Total length (>= 5000 bp)	1088291
Total length (>= 10000 bp)	64562
Total length (>= 25000 bp)	26379
Total length (>= 50000 bp)	0
# contigs	256
Largest contig	26379
Total length	1535387
GC (%)	45.83
N50	5980
N90	4457
auN	6704.3
L50	99
L90	221
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 1 part

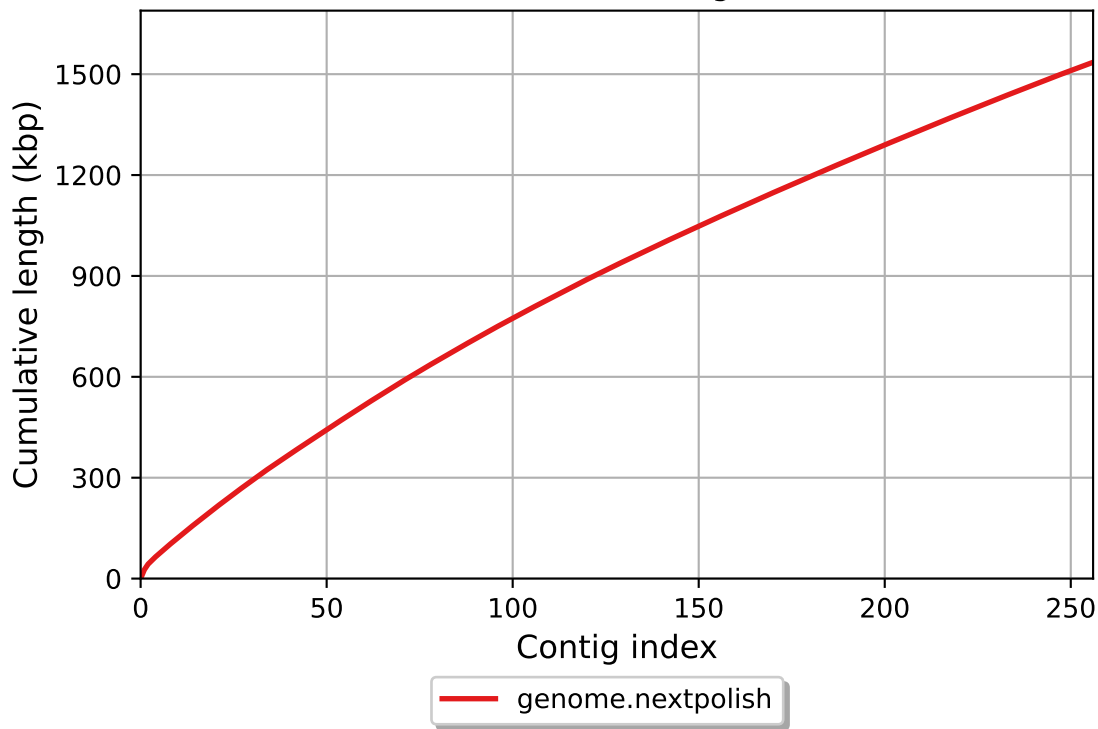
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

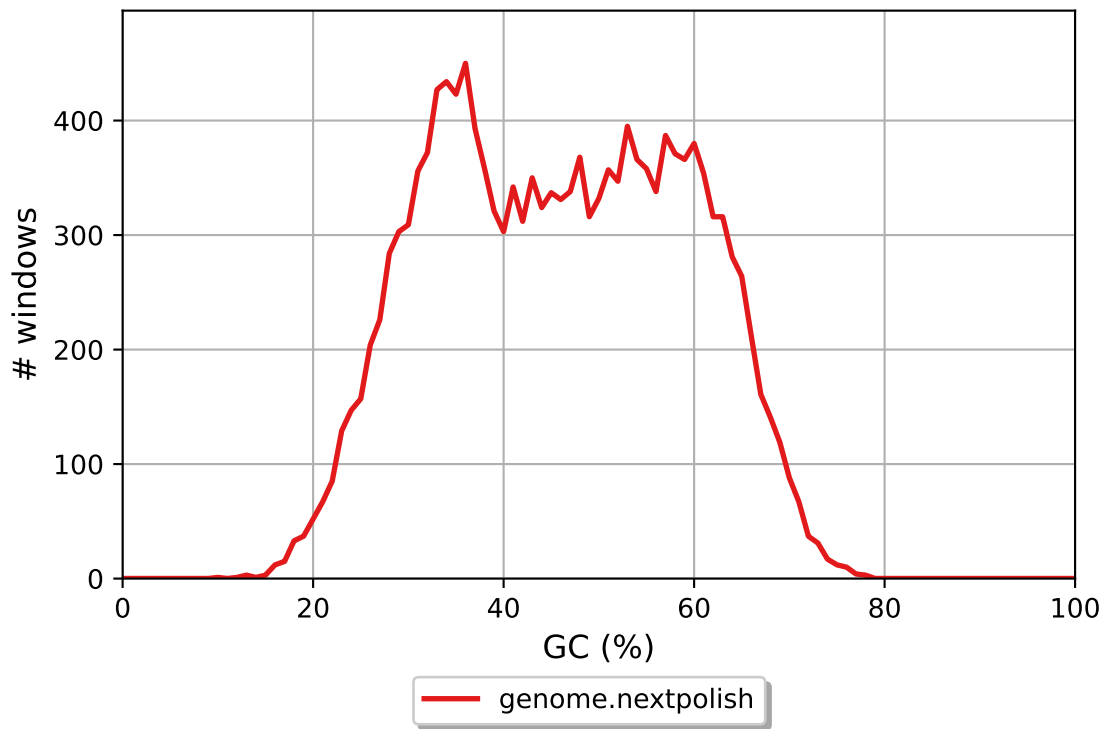


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

