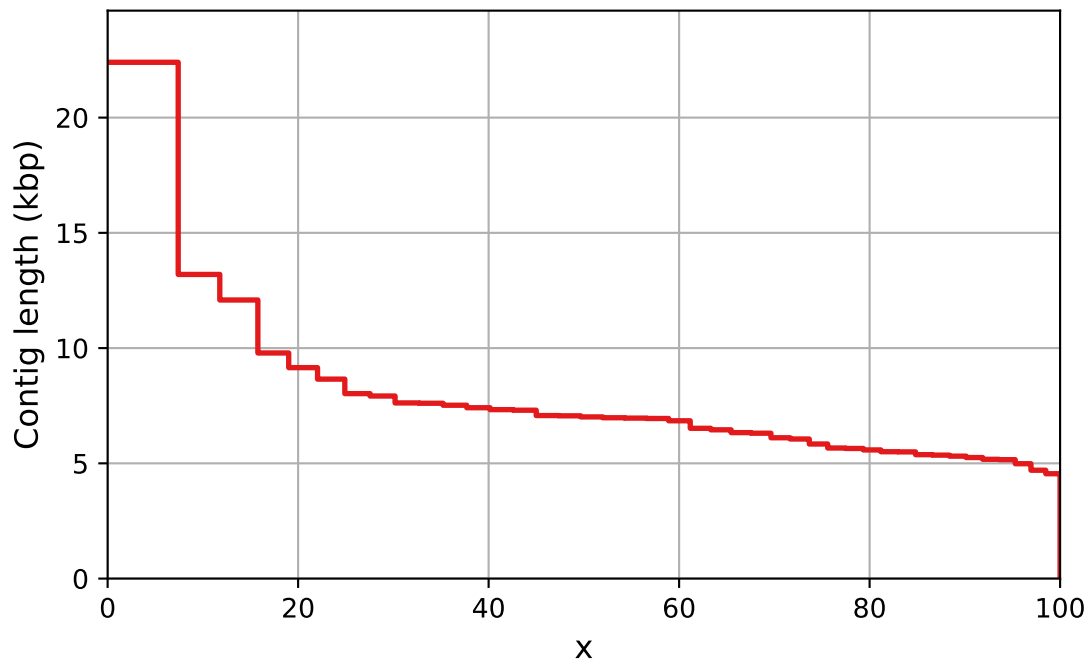


Report

	genome.nextpolish
# contigs (>= 0 bp)	42
# contigs (>= 1000 bp)	42
# contigs (>= 5000 bp)	39
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	302282
Total length (>= 1000 bp)	302282
Total length (>= 5000 bp)	288047
Total length (>= 10000 bp)	47683
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	42
Largest contig	22400
Total length	302282
GC (%)	50.04
N50	7015
N90	5312
auN	8408.6
L50	17
L90	36
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

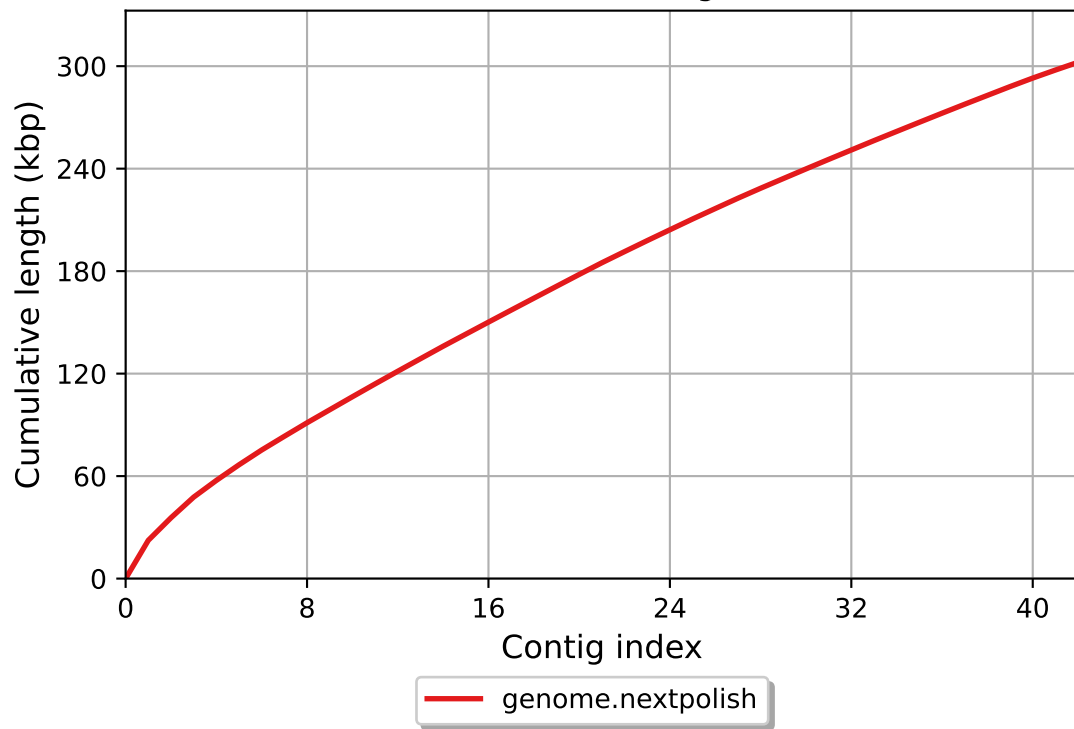
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

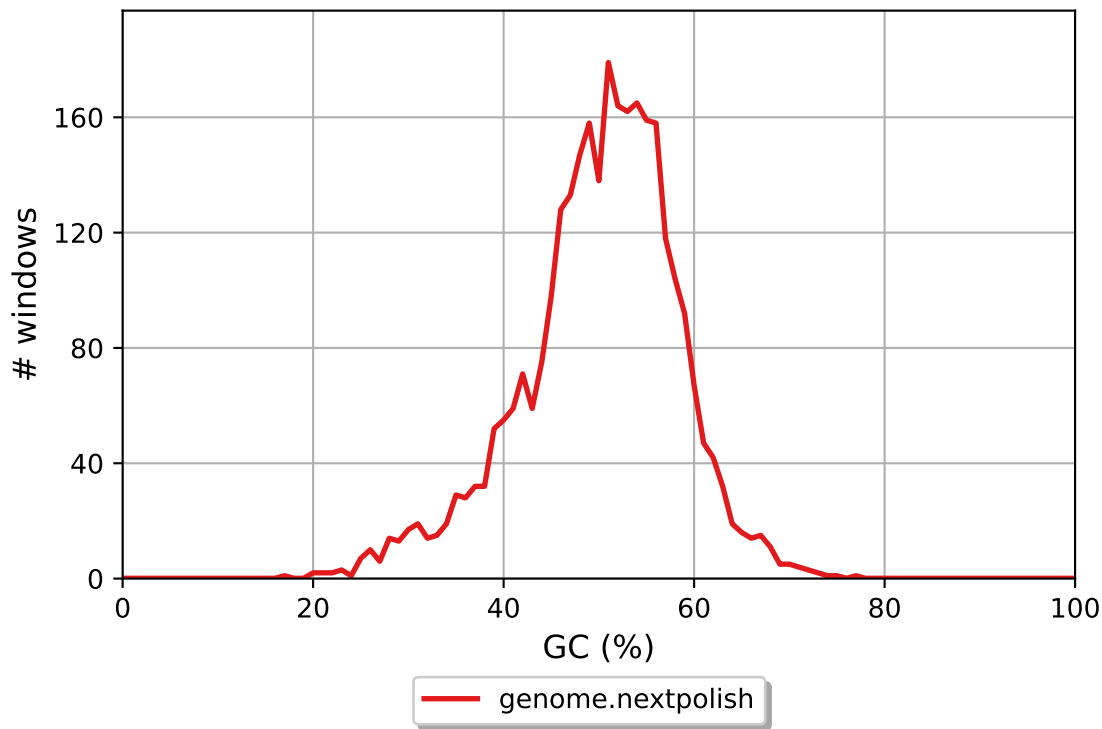


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

