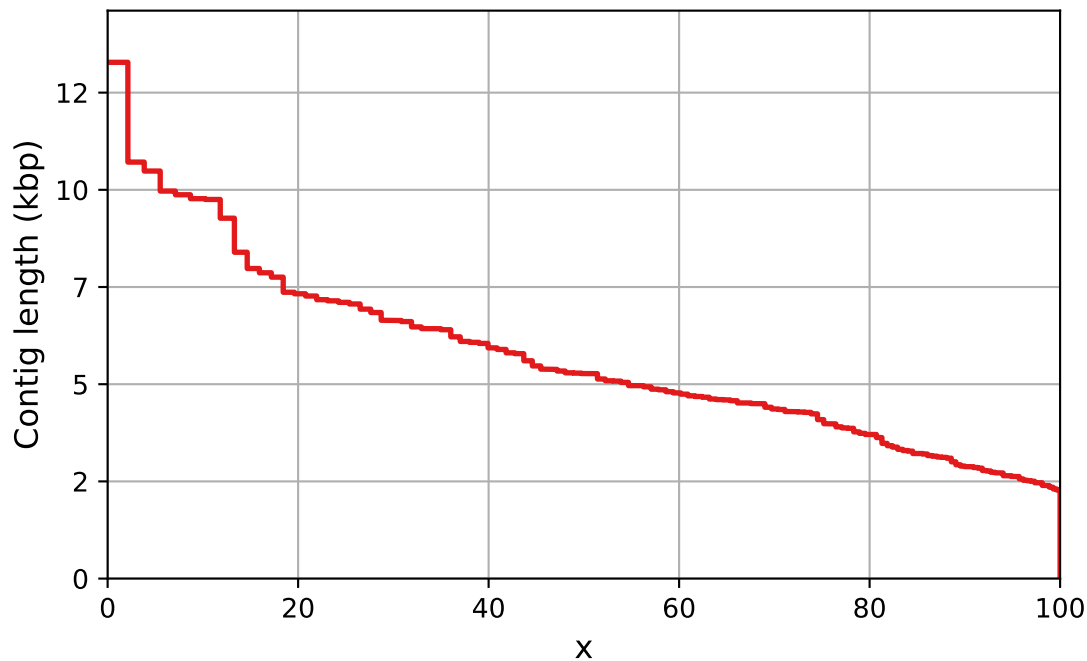


Report

	genome.nextpolish
# contigs (>= 0 bp)	128
# contigs (>= 1000 bp)	128
# contigs (>= 5000 bp)	49
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	624400
Total length (>= 1000 bp)	624400
Total length (>= 5000 bp)	341469
Total length (>= 10000 bp)	34476
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	128
Largest contig	13280
Total length	624400
GC (%)	38.52
N50	5274
N90	2880
auN	5774.4
L50	44
L90	105
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 0 part

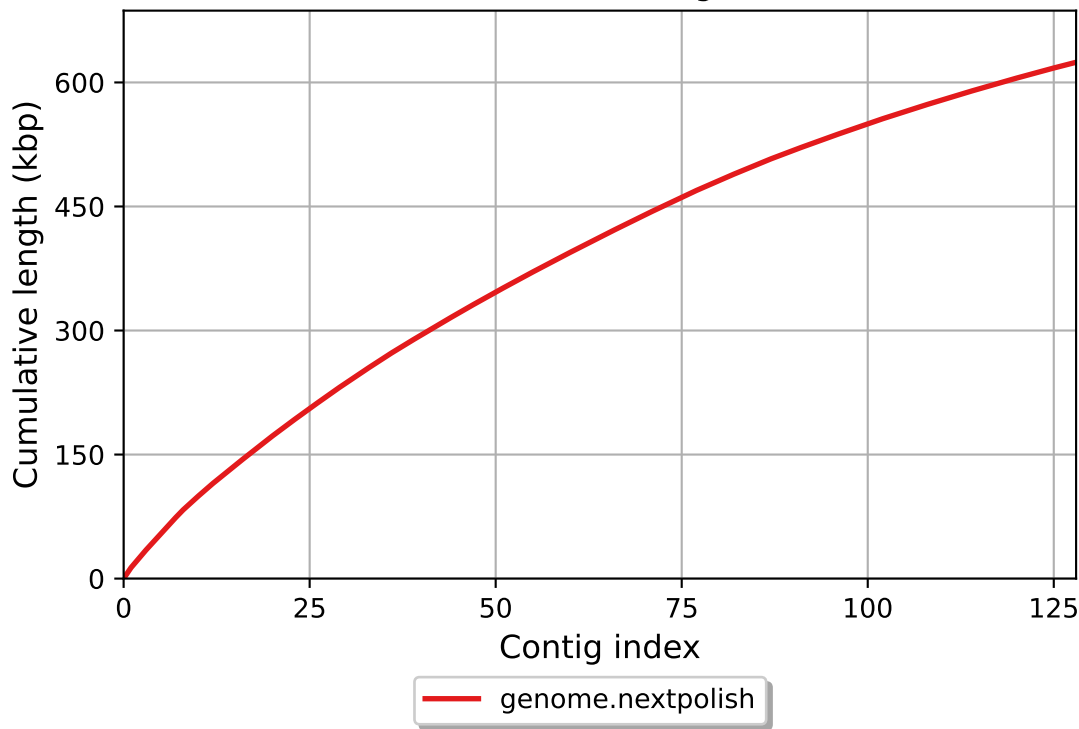
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

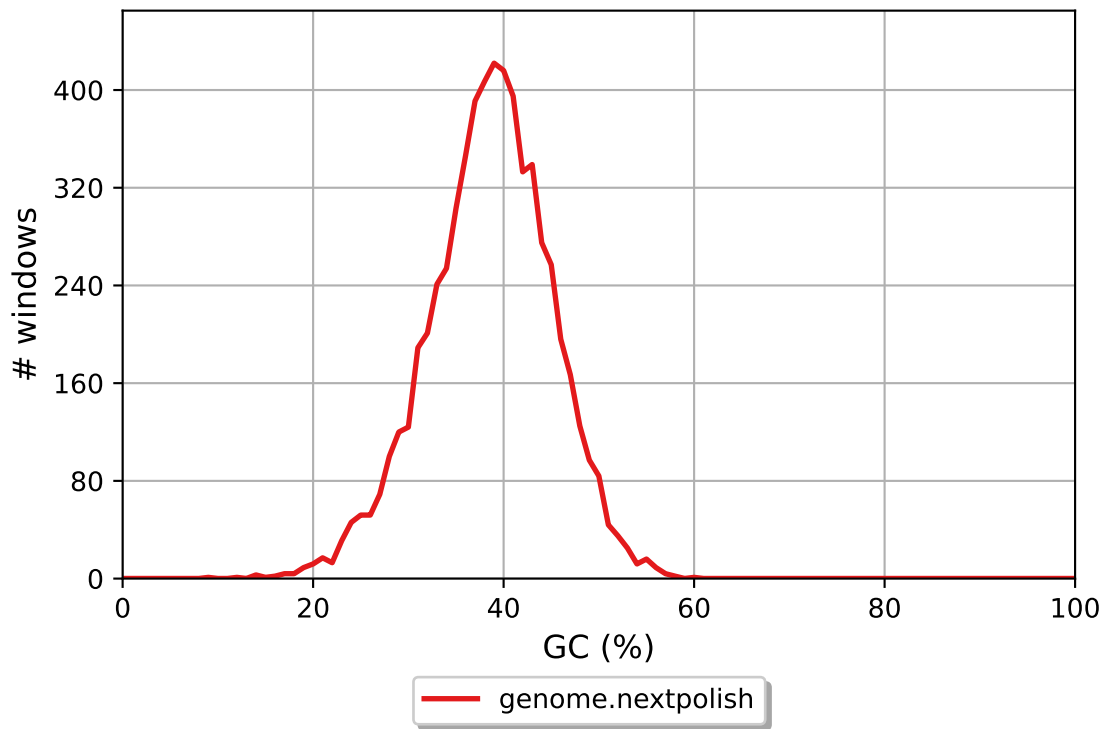


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

