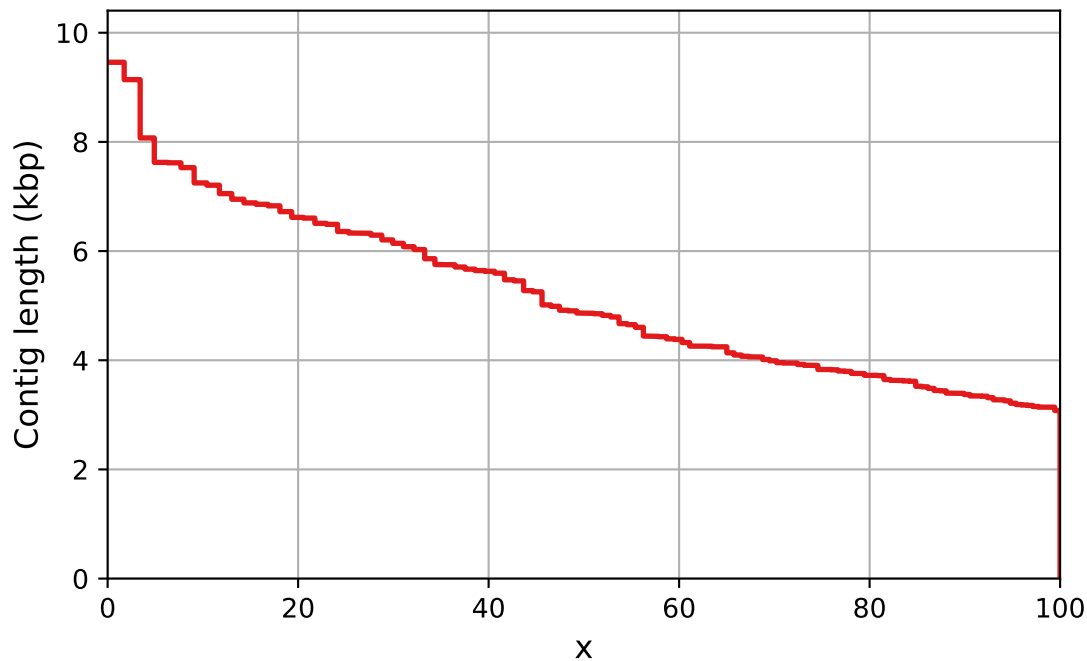


# Report

	genome.nextpolish
# contigs (>= 0 bp)	114
# contigs (>= 1000 bp)	114
# contigs (>= 5000 bp)	39
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	544318
Total length (>= 1000 bp)	544318
Total length (>= 5000 bp)	253258
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	114
Largest contig	9459
Total length	544318
GC (%)	47.18
N50	4865
N90	3373
auN	5204.3
L50	43
L90	98
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 0 part

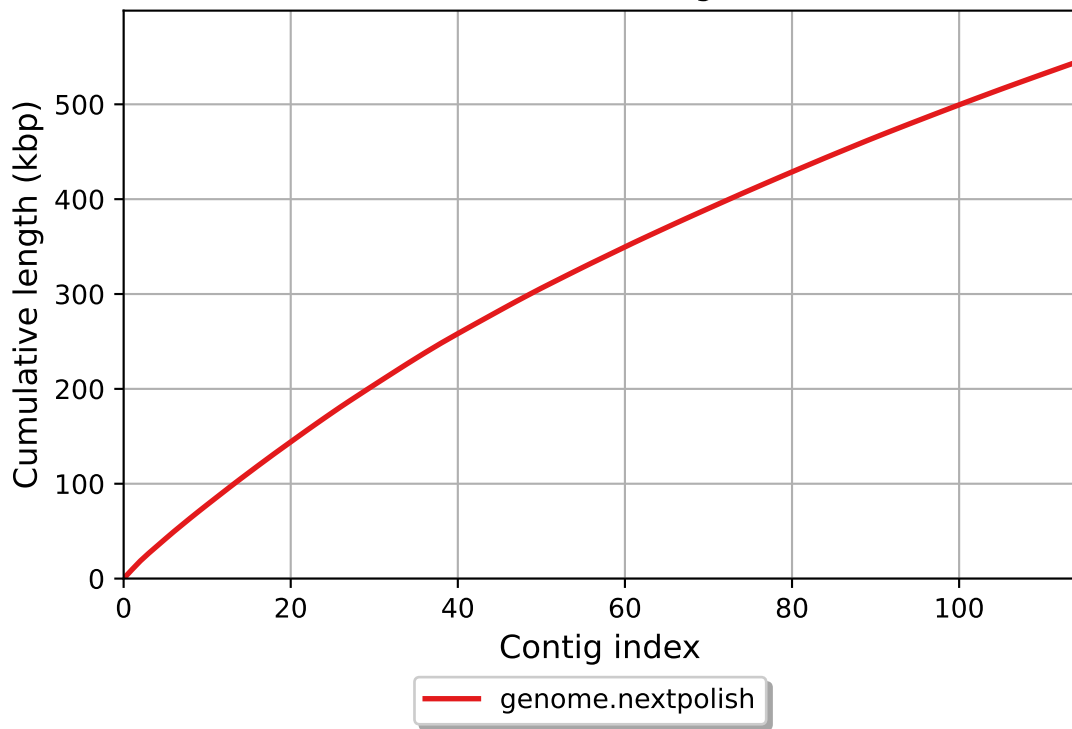
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

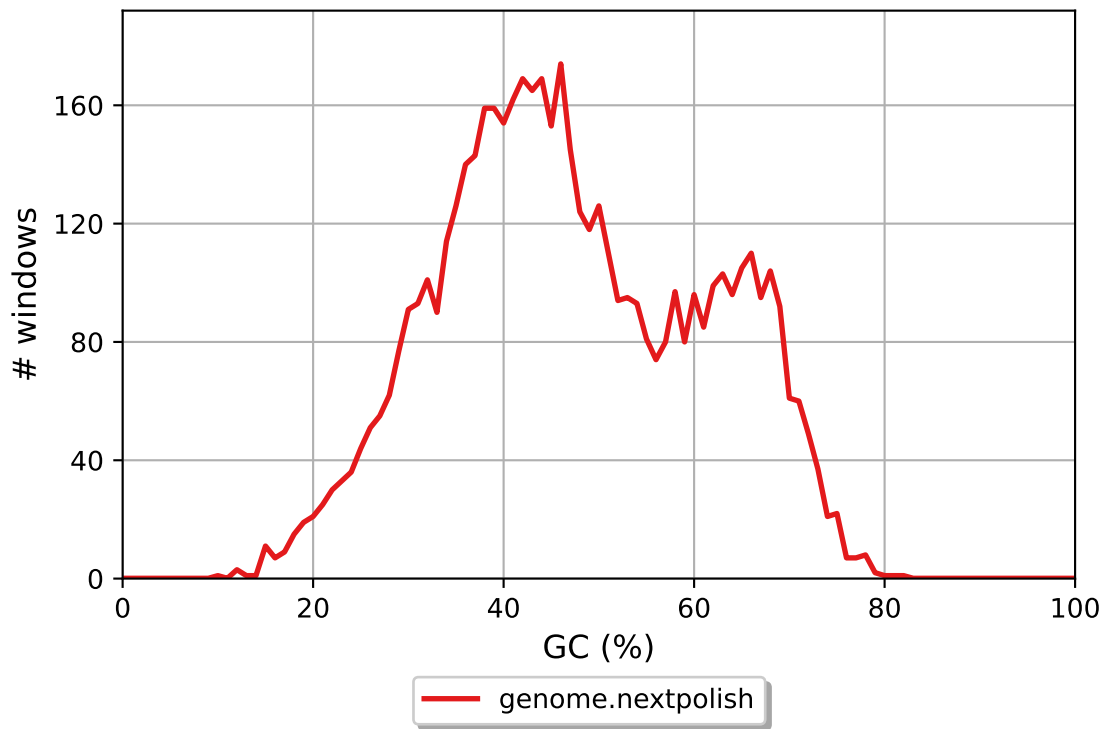


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

