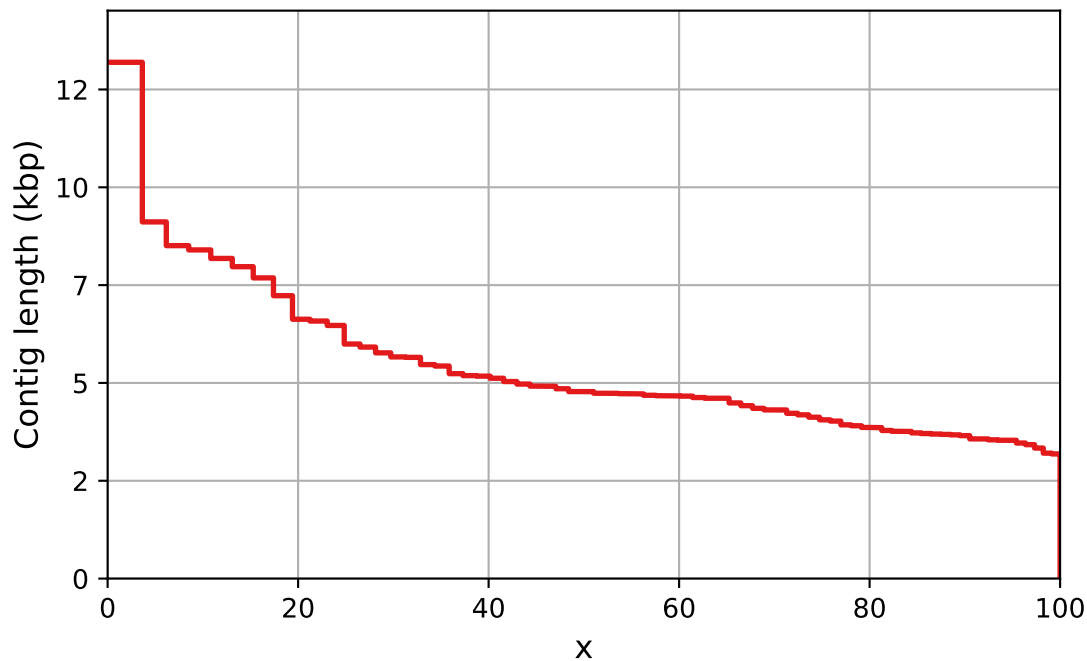


Report

	genome.nextpolish
# contigs (>= 0 bp)	73
# contigs (>= 1000 bp)	73
# contigs (>= 5000 bp)	23
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	362153
Total length (>= 1000 bp)	362153
Total length (>= 5000 bp)	155622
Total length (>= 10000 bp)	13195
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	73
Largest contig	13195
Total length	362153
GC (%)	48.82
N50	4778
N90	3653
auN	5518.0
L50	29
L90	63
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 0 part

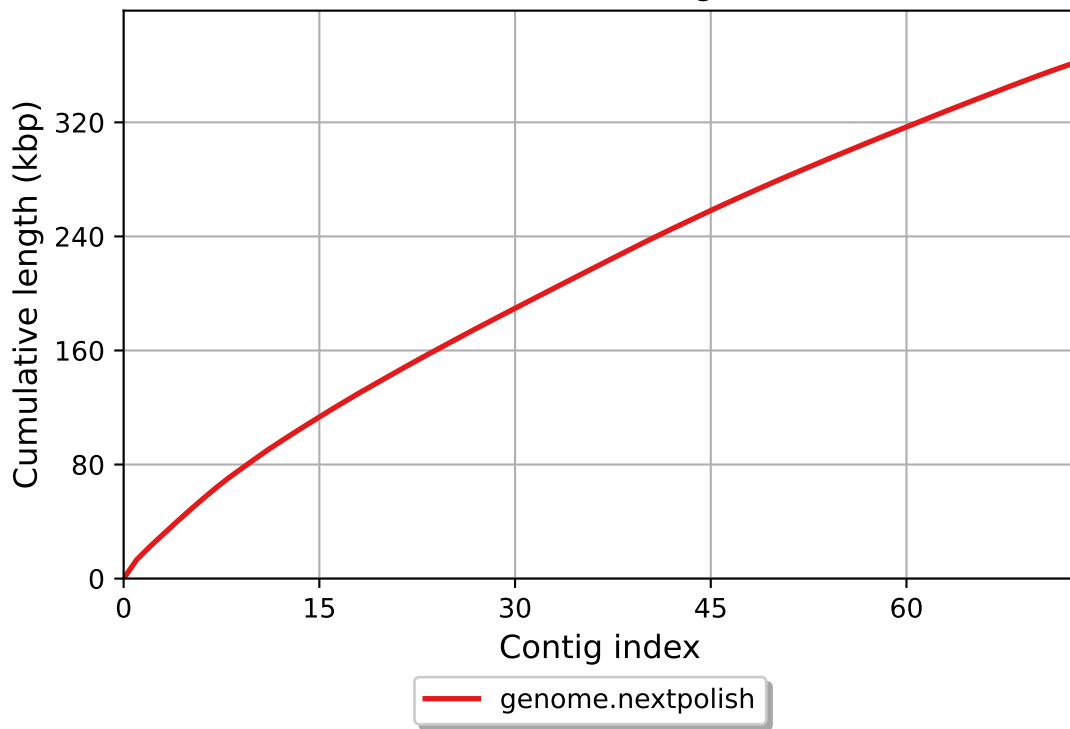
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

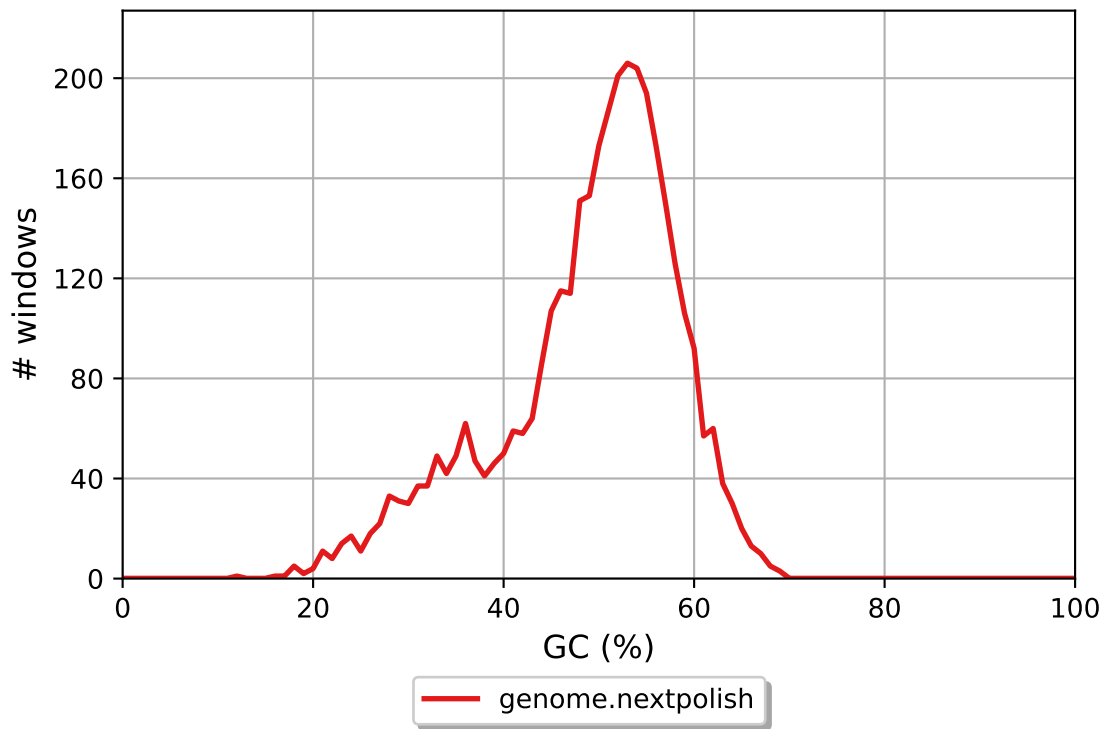


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

