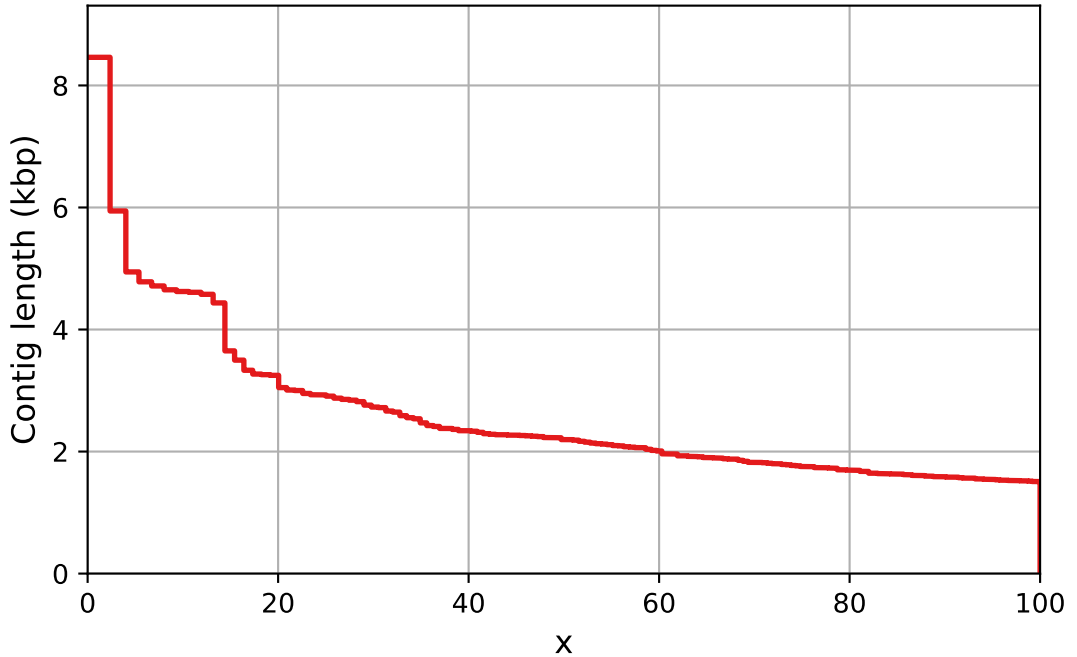


# Report

	MEGAHIT-MetaBAT2-3106237.1.fa
# contigs (>= 0 bp)	160
# contigs (>= 1000 bp)	160
# contigs (>= 5000 bp)	2
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	359000
Total length (>= 1000 bp)	359000
Total length (>= 5000 bp)	14404
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	160
Largest contig	8461
Total length	359000
GC (%)	40.07
N50	2196
N75	1755
L50	59
L75	105
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 0 part

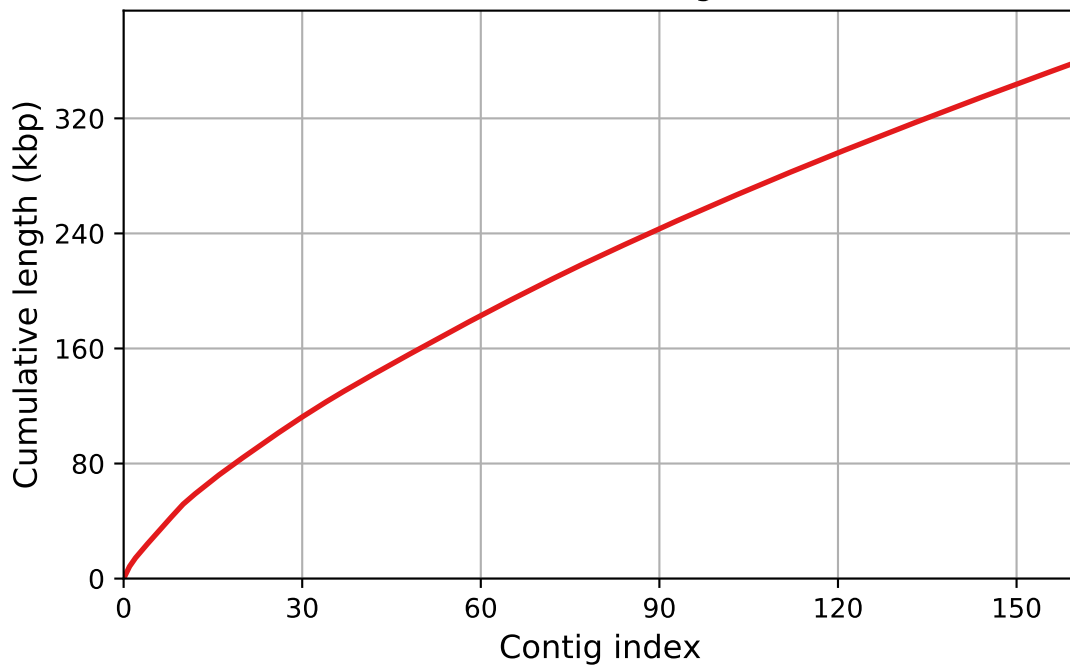
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



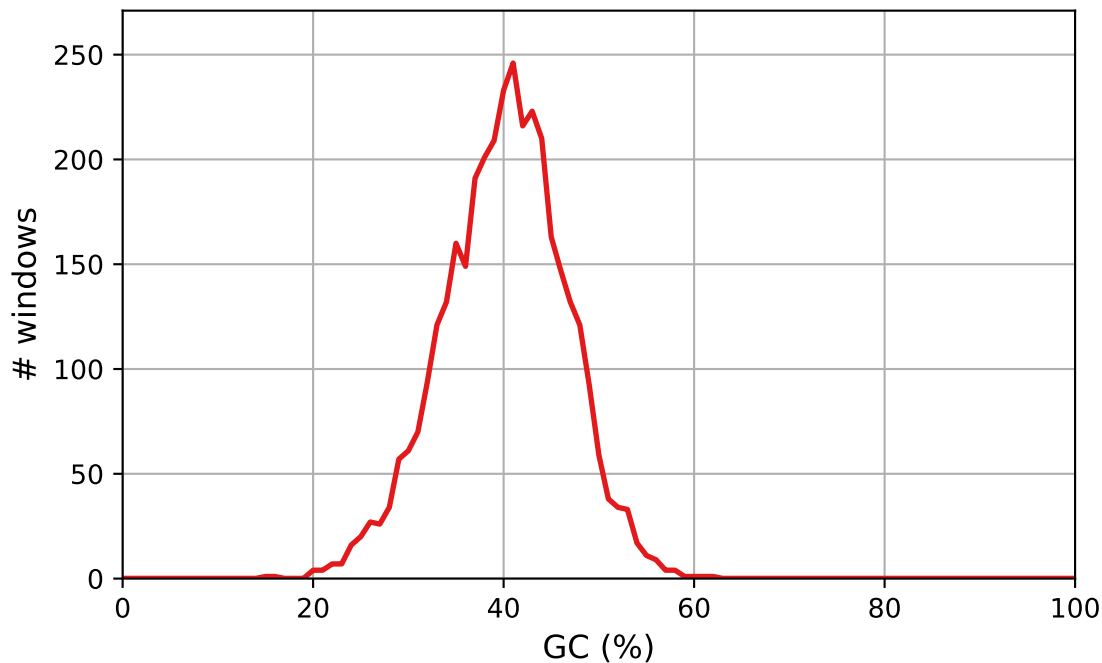
— MEGAHIT-MetaBAT2-3106237.1.fa

Cumulative length



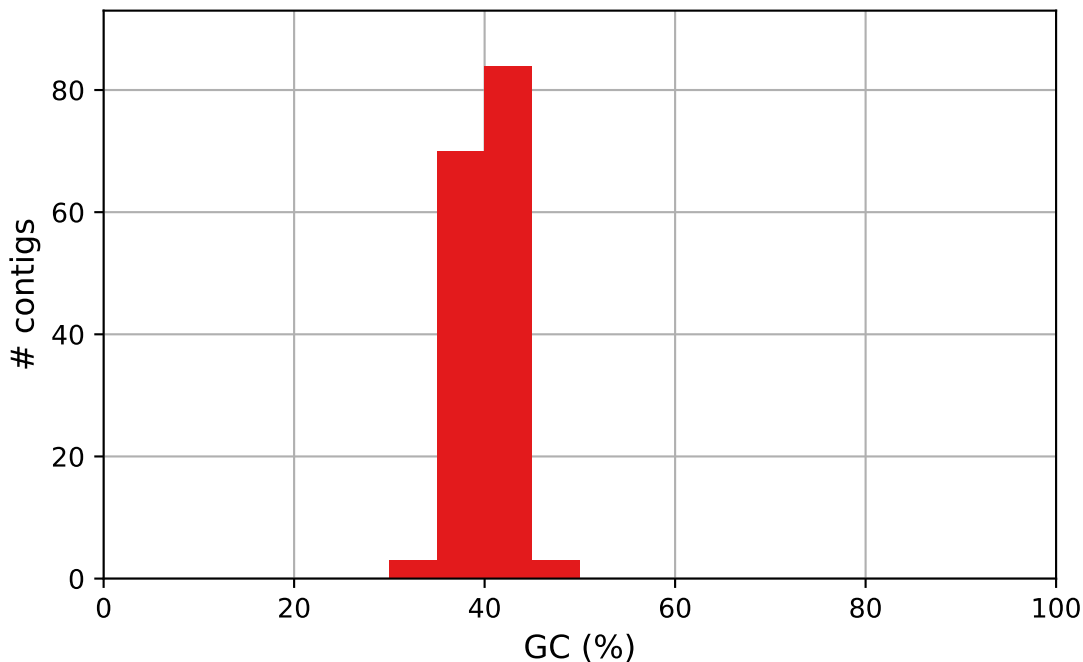
— MEGAHIT-MetaBAT2-3106237.1.fa

## GC content



— MEGAHIT-MetaBAT2-3106237.1.fa

MEGAHIT-MetaBAT2-3106237.1.fa GC content



MEGAHit-MetaBAT2-3106237.1.fa