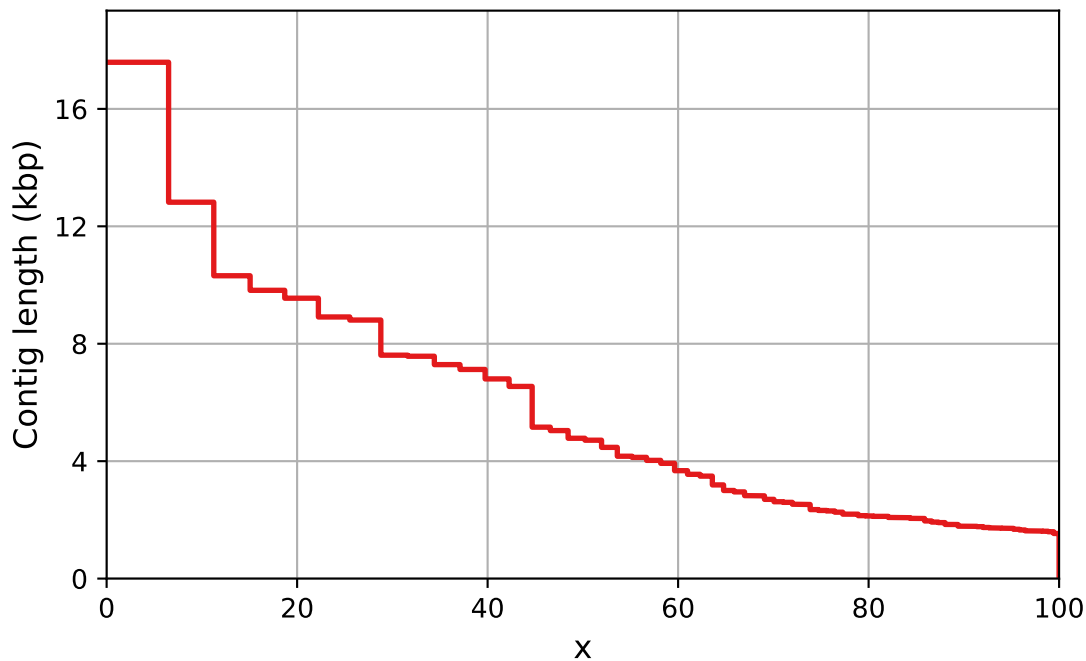


# Report

	MEGAHIT-MetaBAT2-3106237.9.fa
# contigs (>= 0 bp)	72
# contigs (>= 1000 bp)	72
# contigs (>= 5000 bp)	15
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	270296
Total length (>= 1000 bp)	270296
Total length (>= 5000 bp)	130960
Total length (>= 10000 bp)	40725
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	72
Largest contig	17588
Total length	270296
GC (%)	36.51
N50	4780
N75	2321
L50	16
L75	37
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 0 part

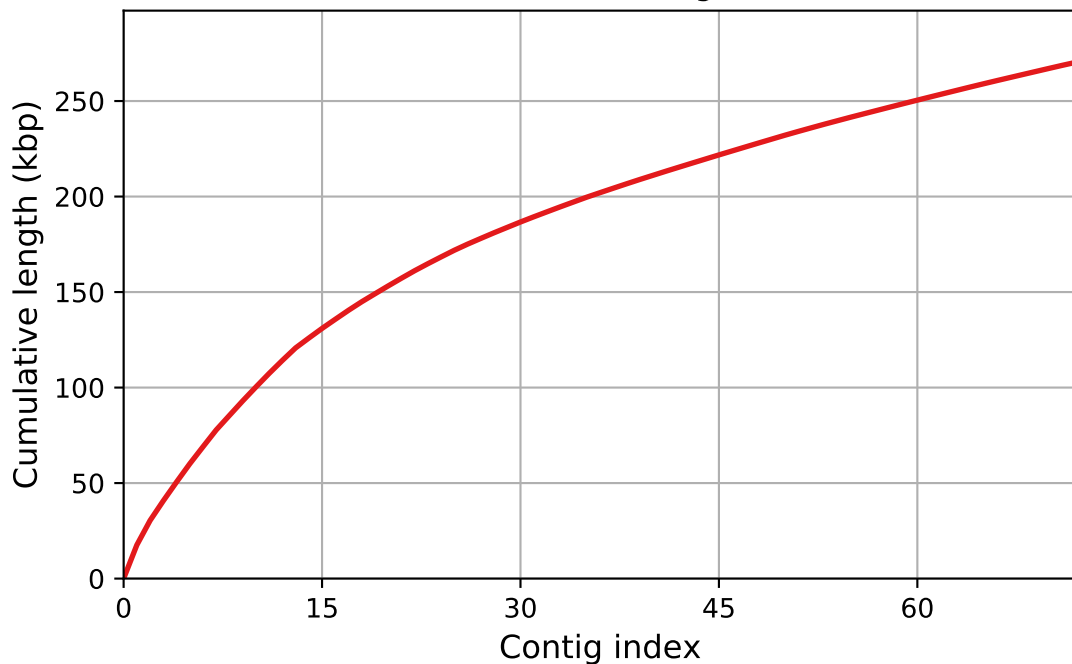
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



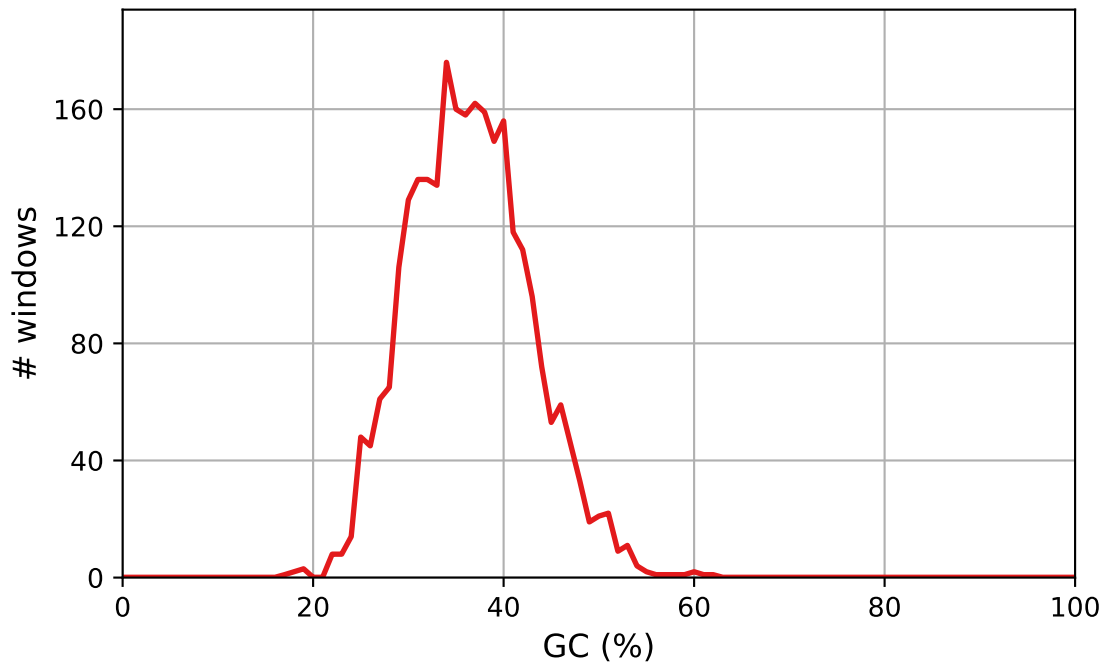
MEGAHIT-MetaBAT2-3106237.9.fa

Cumulative length



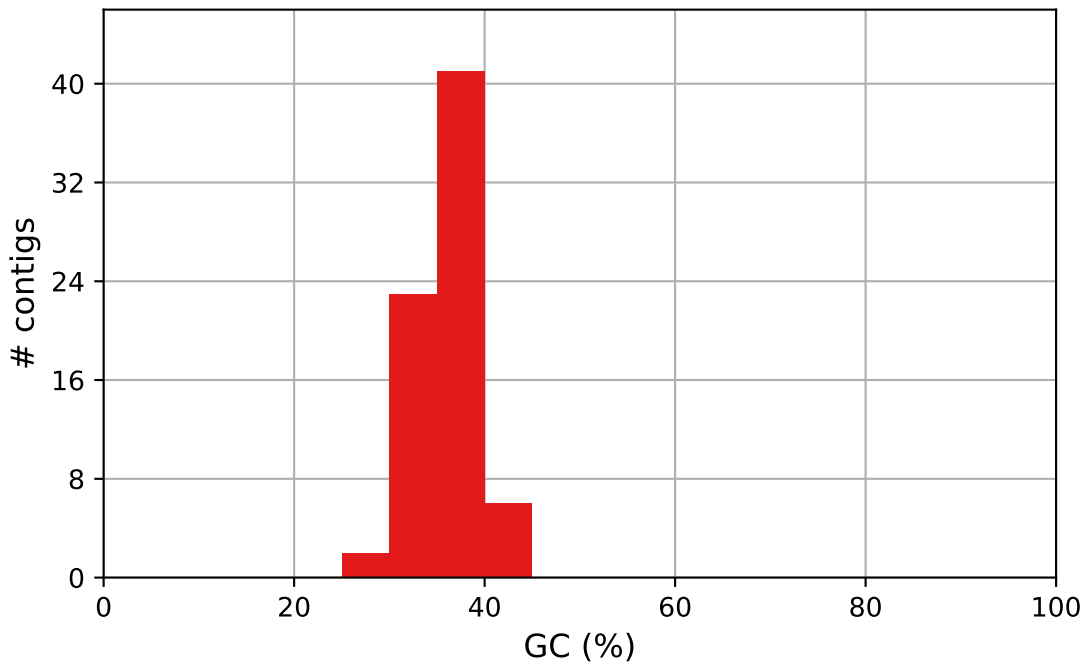
— MEGAHIT-MetaBAT2-3106237.9.fa

## GC content



— MEGAHIT-MetaBAT2-3106237.9.fa

MEGAHIT-MetaBAT2-3106237.9.fa GC content



MEGAhit-MetaBAT2-3106237.9.fa