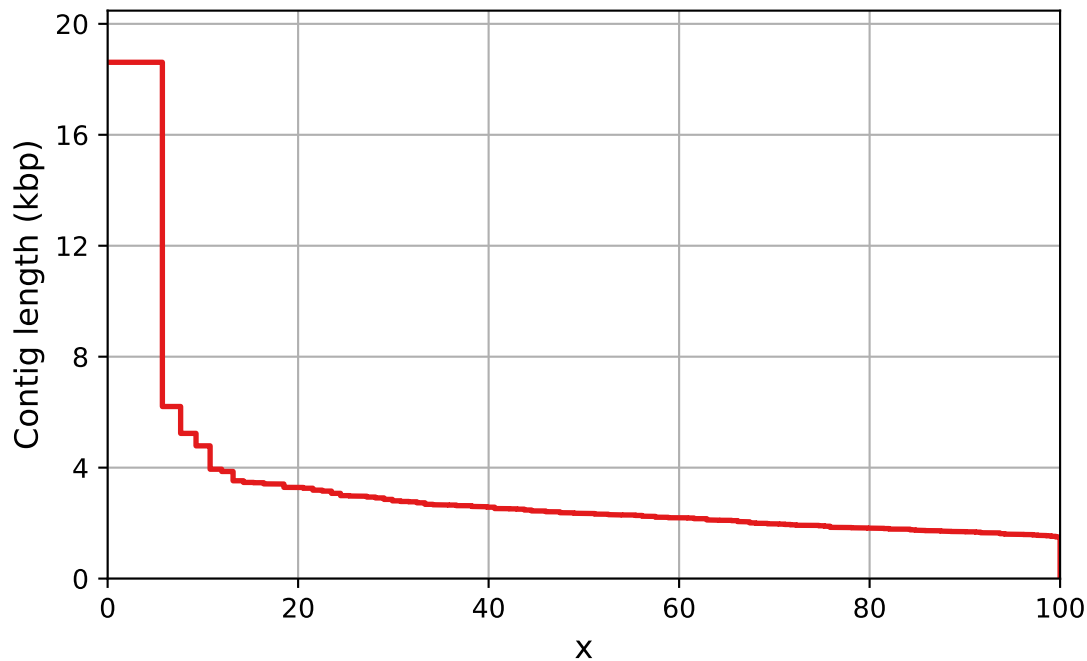


Report

	MEGAHIT-MetaBAT2-3106237.2.fa
# contigs (>= 0 bp)	134
# contigs (>= 1000 bp)	134
# contigs (>= 5000 bp)	3
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	323770
Total length (>= 1000 bp)	323770
Total length (>= 5000 bp)	30053
Total length (>= 10000 bp)	18614
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	134
Largest contig	18614
Total length	323770
GC (%)	44.13
N50	2349
N75	1904
L50	49
L75	87
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 0 part

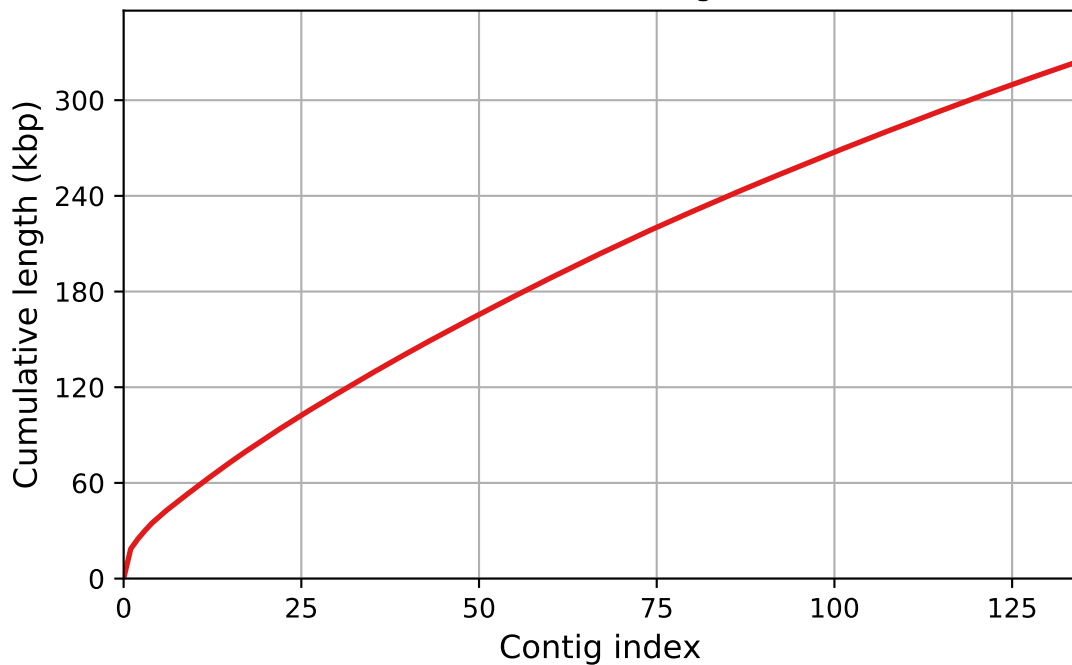
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



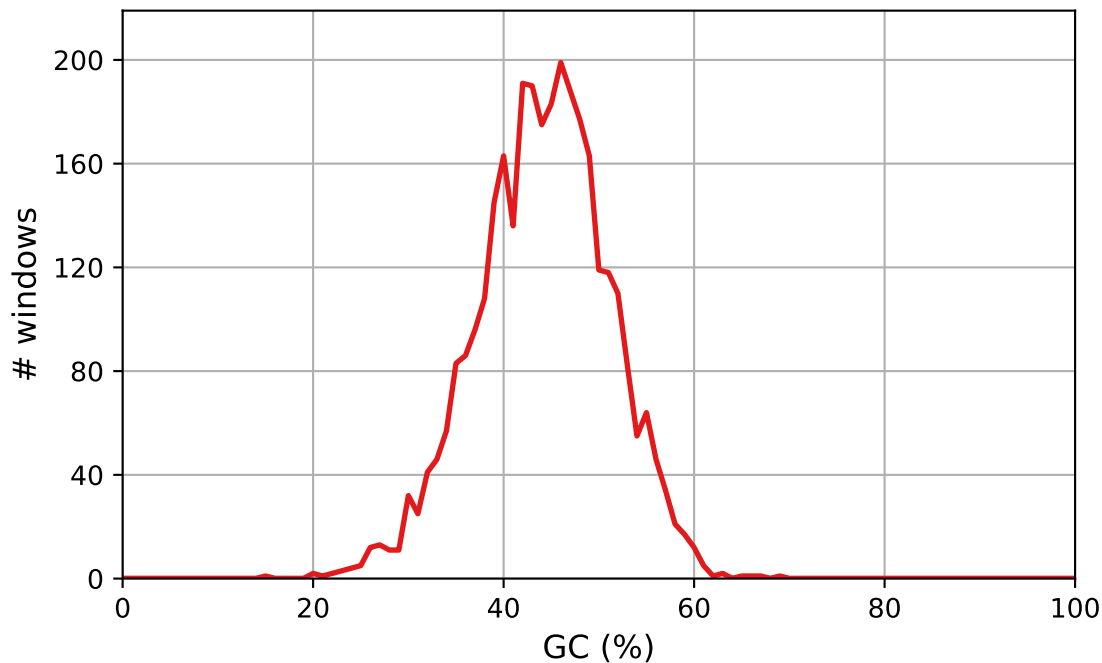
— MEGAHIT-MetaBAT2-3106237.2.fa

Cumulative length



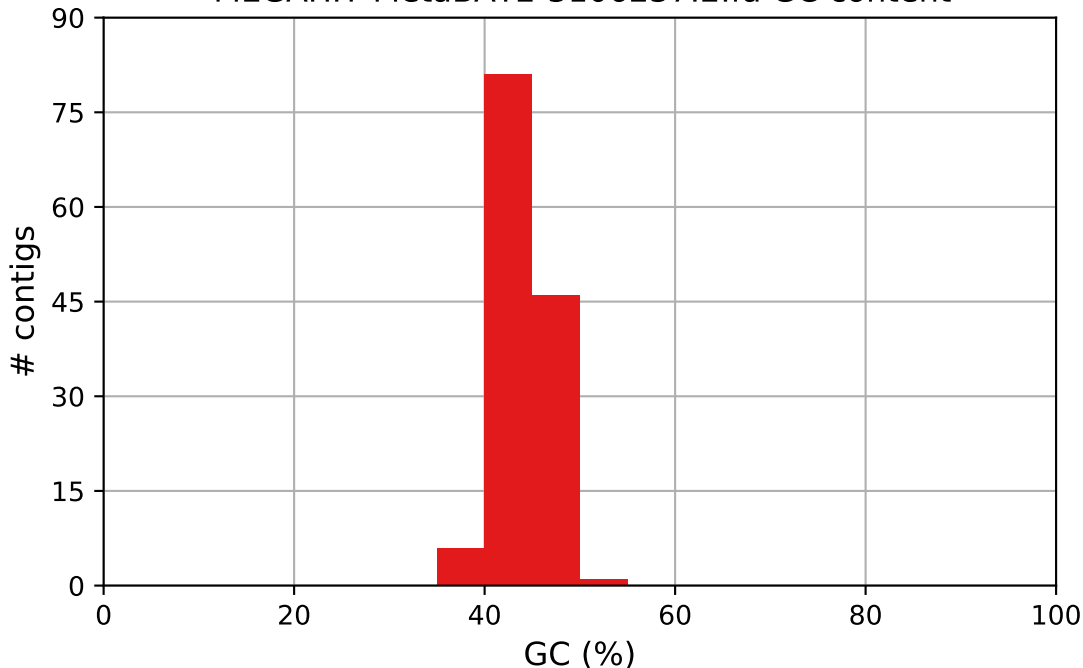
— MEGAHIT-MetaBAT2-3106237.2.fa

GC content



— MEGAHit-MetaBAT2-3106237.2.fa

MEGAHIT-MetaBAT2-3106237.2.fa GC content



MEGAHit-MetaBAT2-3106237.2.fa