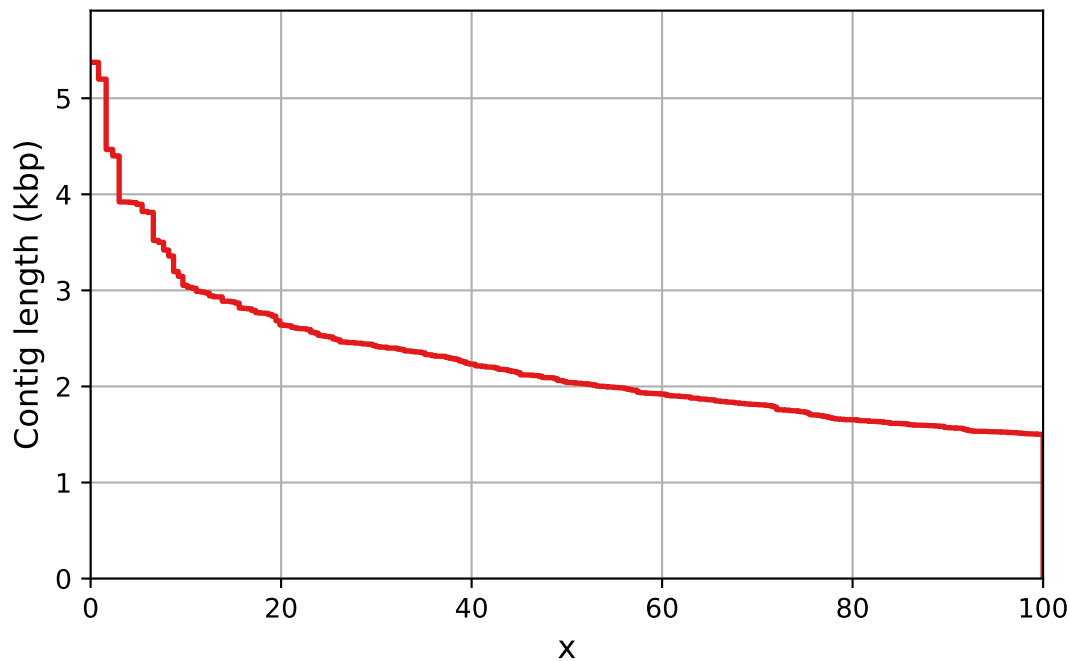


Report

	MEGAHIT-MetaBAT2-3106384.2.fa
# contigs (>= 0 bp)	311
# contigs (>= 1000 bp)	311
# contigs (>= 5000 bp)	2
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	649059
Total length (>= 1000 bp)	649059
Total length (>= 5000 bp)	10573
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	311
Largest contig	5374
Total length	649059
GC (%)	43.32
N50	2042
N75	1732
L50	124
L75	210
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 0 part

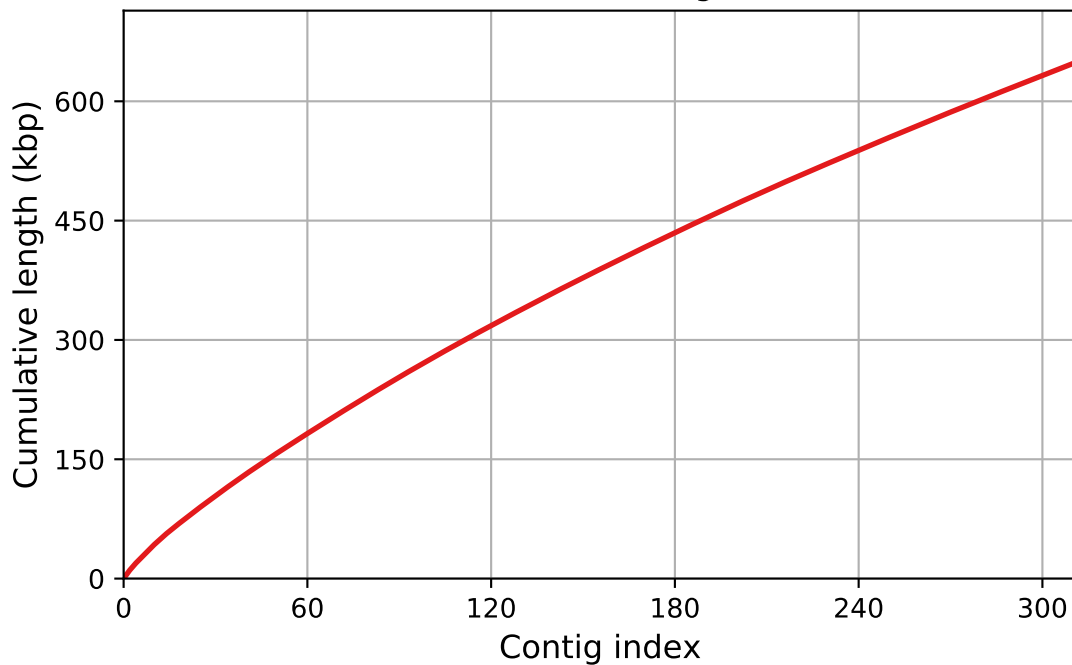
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



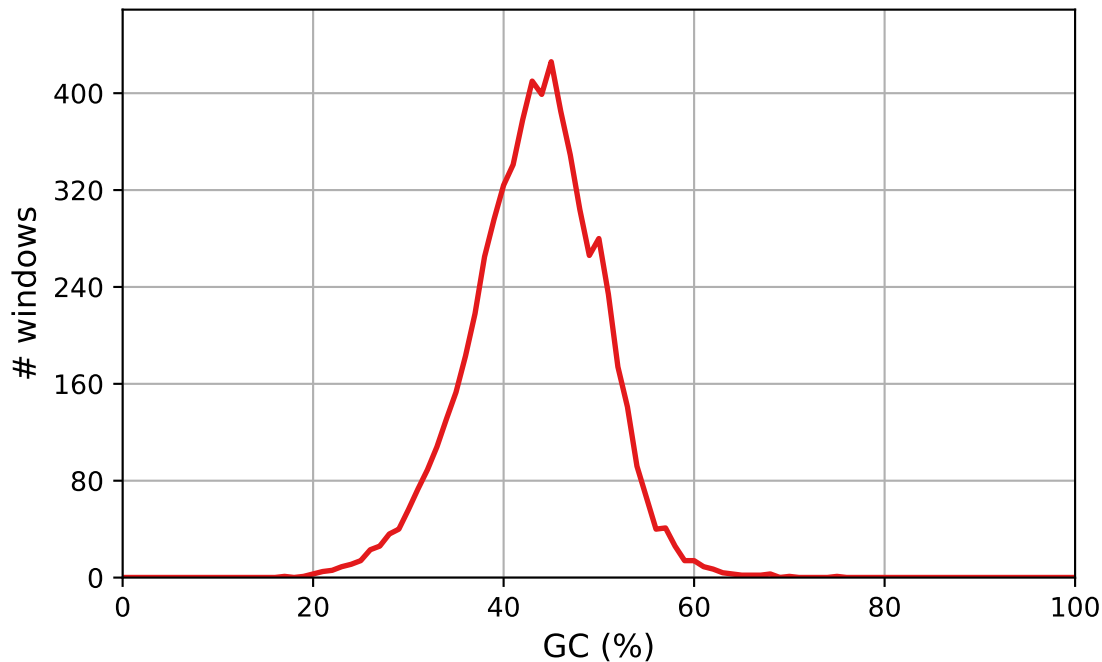
MEGAHIT-MetaBAT2-3106384.2.fa

Cumulative length



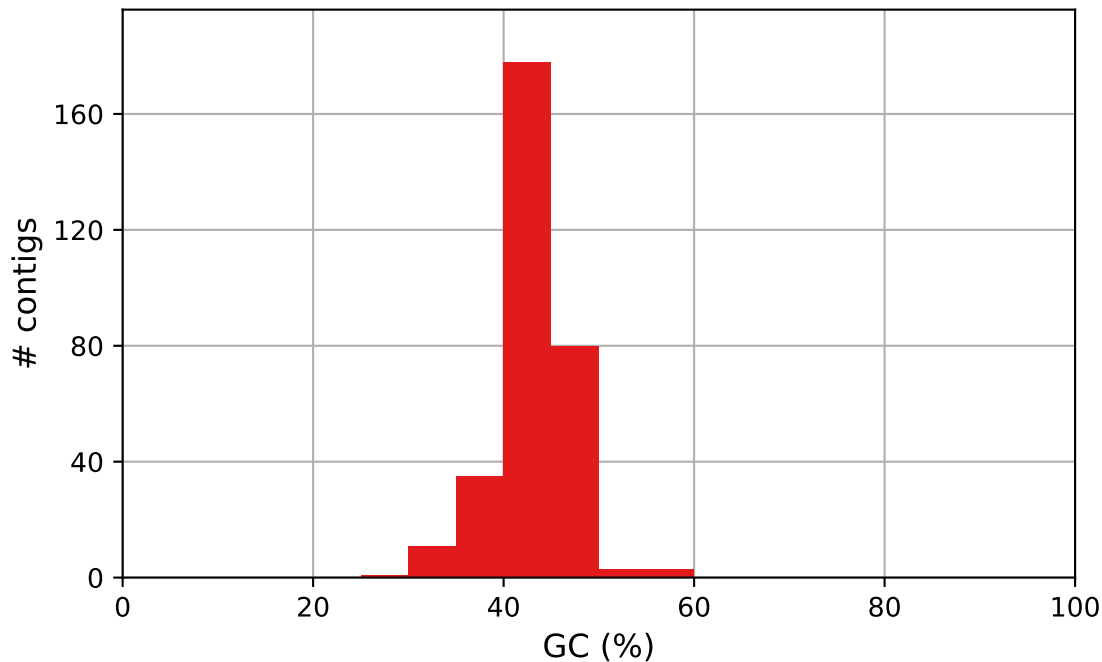
— MEGAHIT-MetaBAT2-3106384.2.fa

GC content



— MEGAHIT-MetaBAT2-3106384.2.fa

MEGAHIT-MetaBAT2-3106384.2.fa GC content



MEGAhit-MetaBAT2-3106384.2.fa