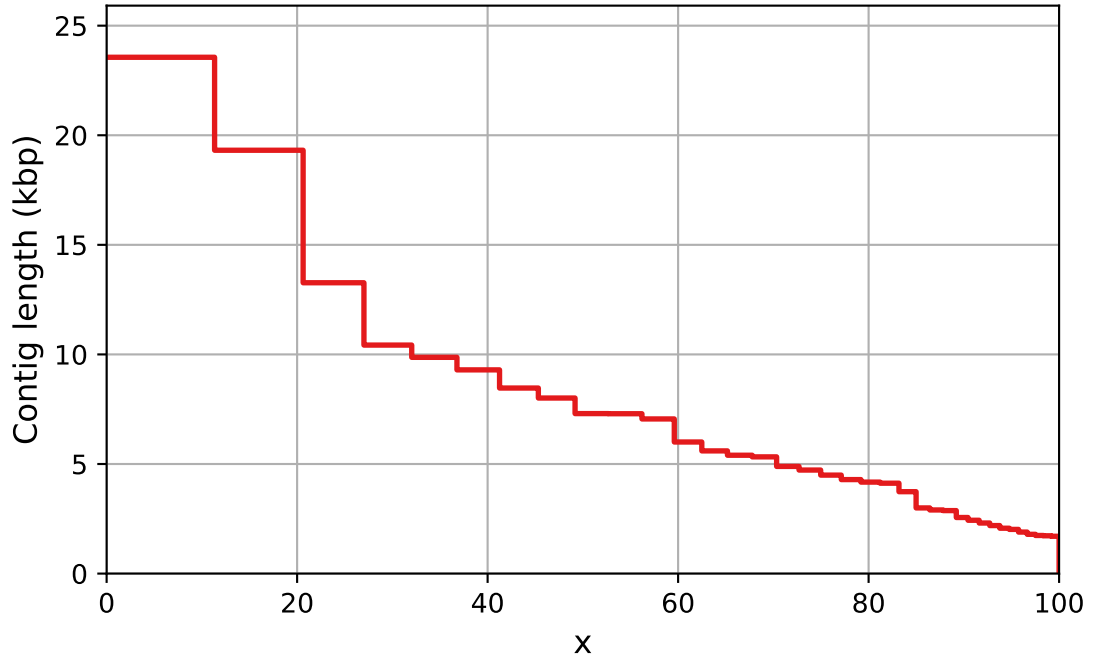


Report

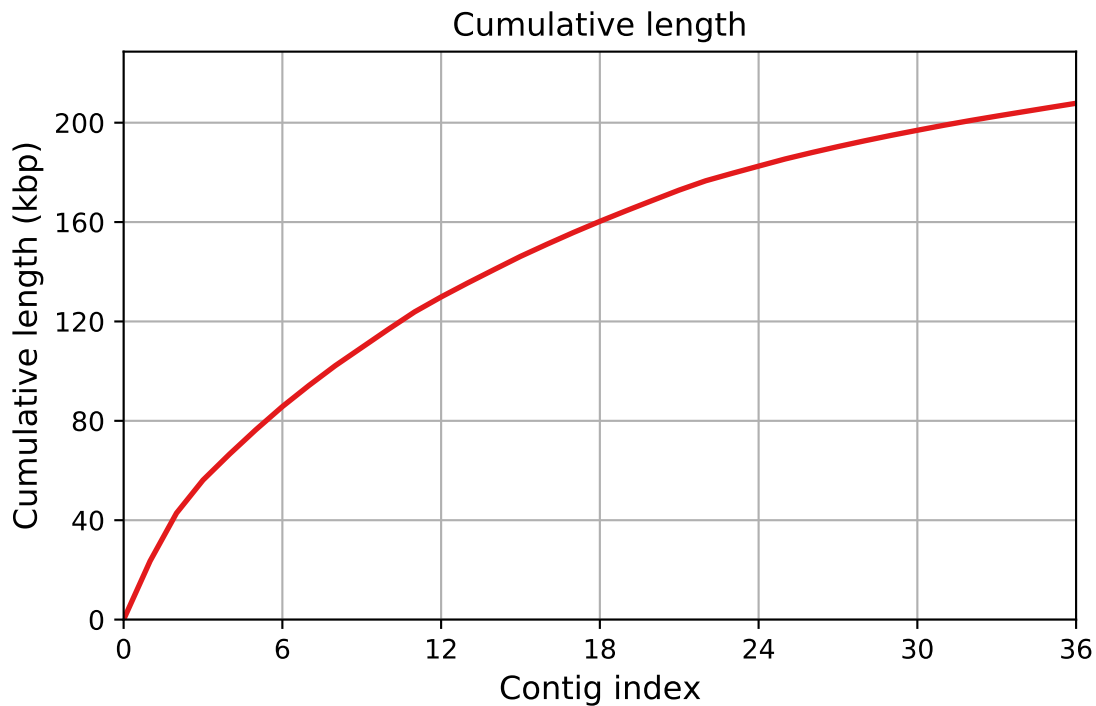
	MEGAHIT-MetaBAT2-3108596.10.fa
# contigs (>= 0 bp)	36
# contigs (>= 1000 bp)	36
# contigs (>= 5000 bp)	15
# contigs (>= 10000 bp)	4
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	207819
Total length (>= 1000 bp)	207819
Total length (>= 5000 bp)	146186
Total length (>= 10000 bp)	66570
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	36
Largest contig	23558
Total length	207819
GC (%)	47.58
N50	7300
N75	4493
L50	9
L75	18
# 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

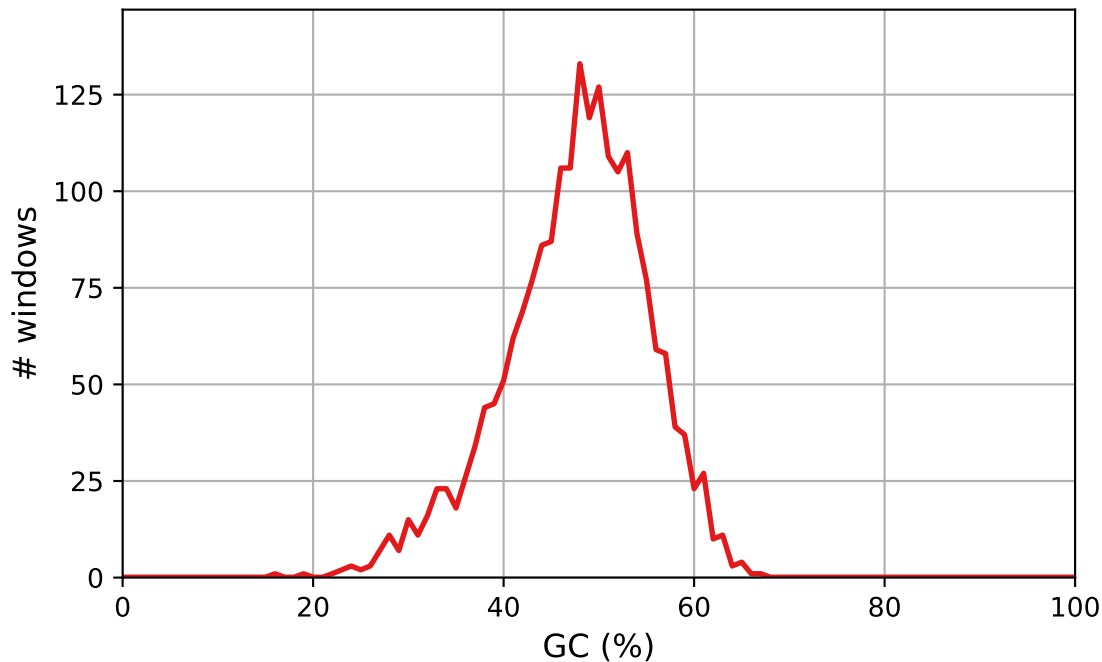


MEGAHIT-MetaBAT2-3108596.10.fa



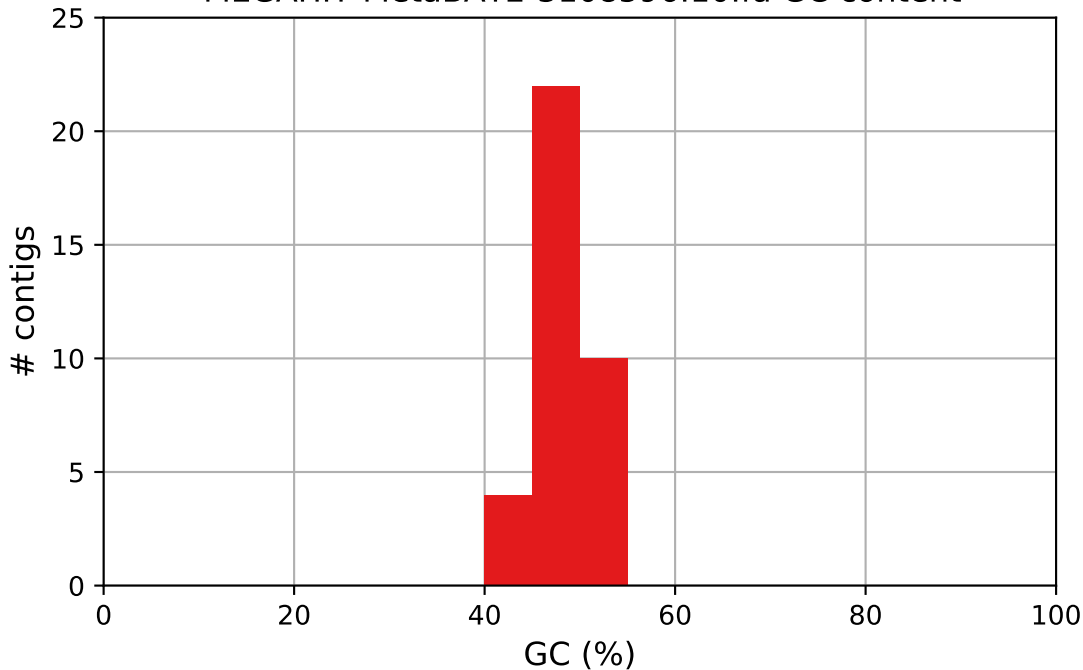
MEGAHIT-MetaBAT2-3108596.10.fa

GC content



MEGAHit-MetaBAT2-3108596.10.fa

MEGAHIT-MetaBAT2-3108596.10.fa GC content



MEGAhit-MetaBAT2-3108596.10.fa