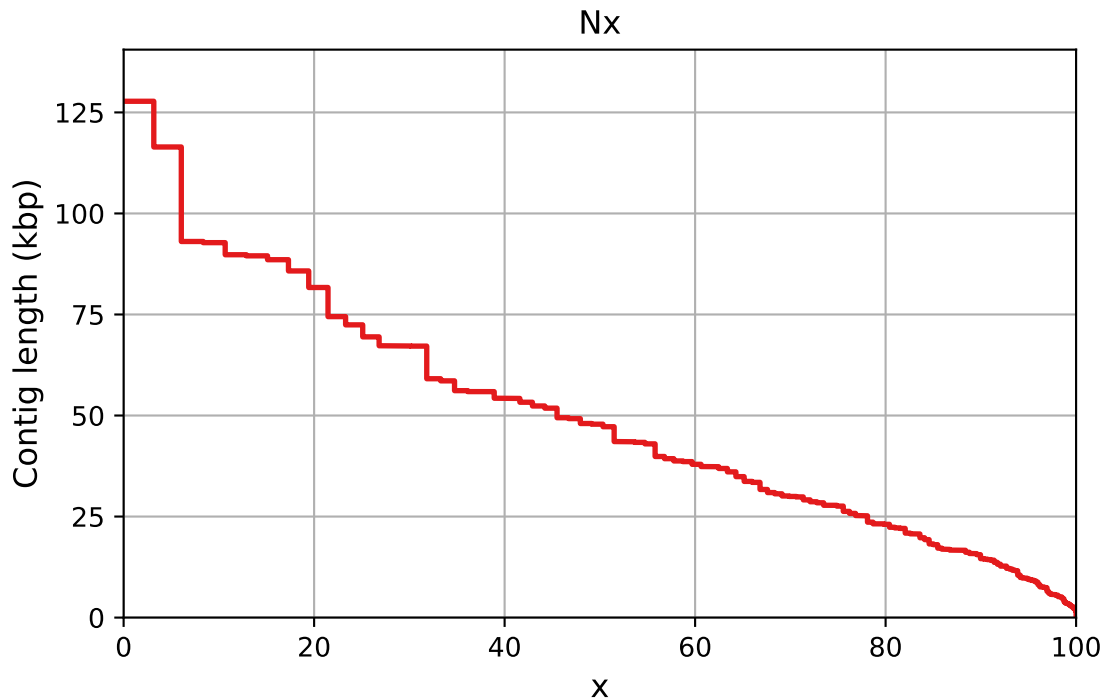


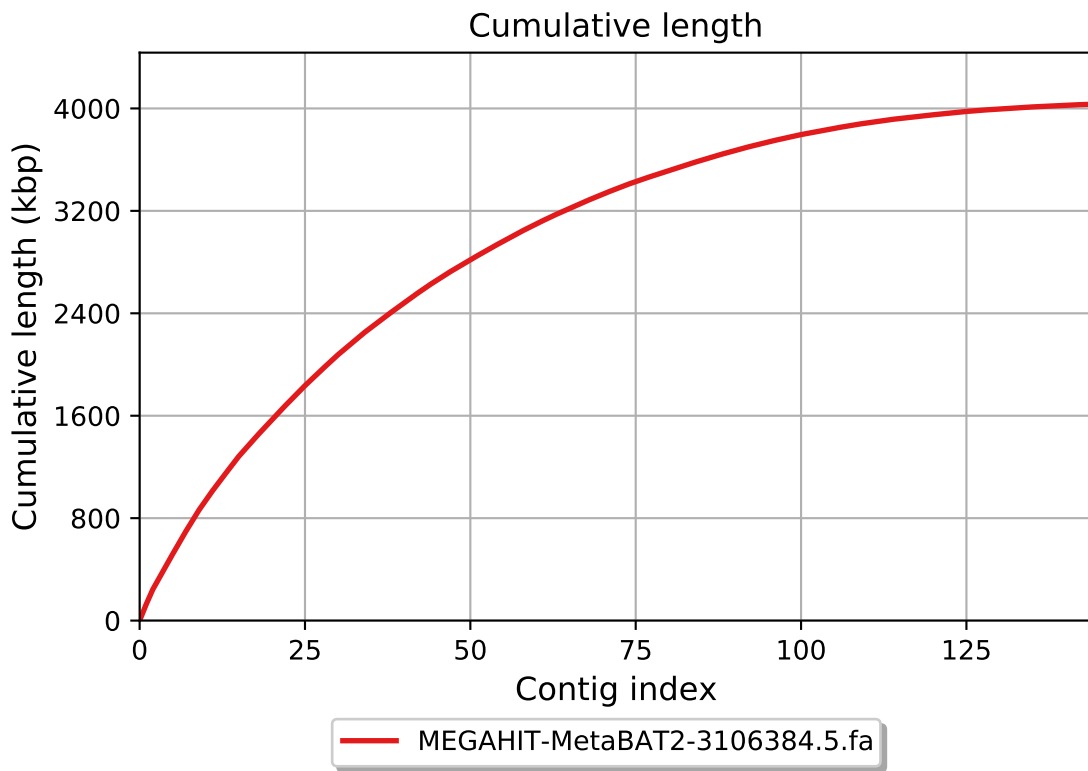
# Report

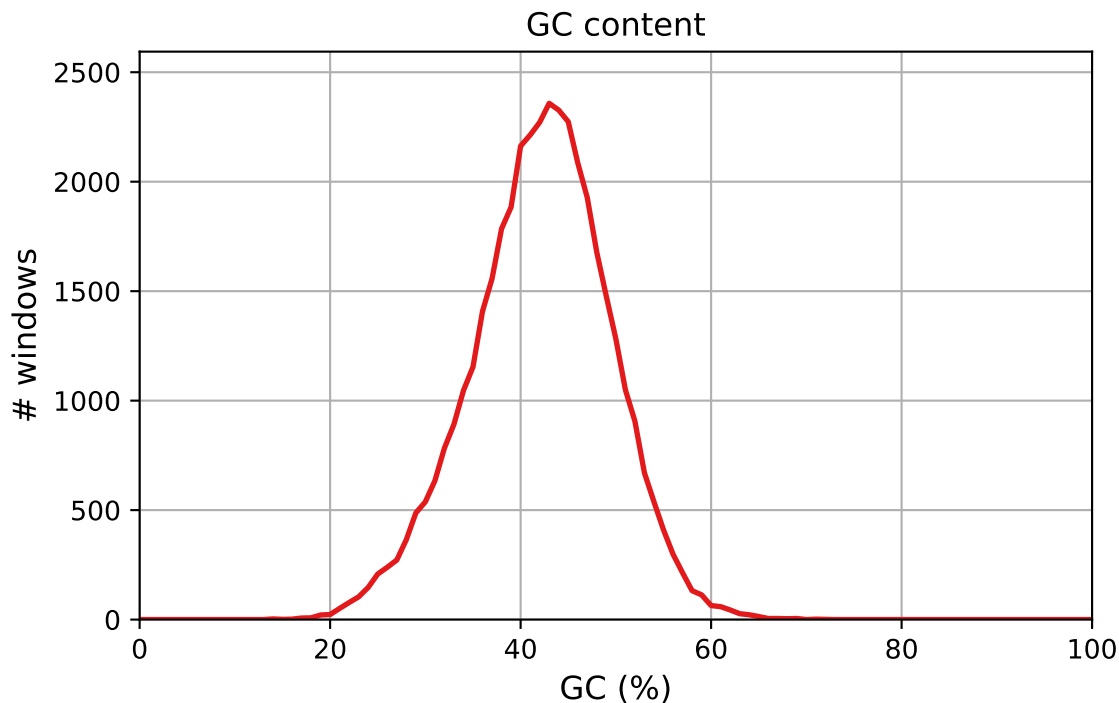
	MEGAHIT-MetaBAT2-3106384.5.fa
# contigs (>= 0 bp)	144
# contigs (>= 1000 bp)	144
# contigs (>= 5000 bp)	124
# contigs (>= 10000 bp)	100
# contigs (>= 25000 bp)	62
# contigs (>= 50000 bp)	25
Total length (>= 0 bp)	4032579
Total length (>= 1000 bp)	4032579
Total length (>= 5000 bp)	3971547
Total length (>= 10000 bp)	3795802
Total length (>= 25000 bp)	3149388
Total length (>= 50000 bp)	1834988
# contigs	144
Largest contig	127758
Total length	4032579
GC (%)	42.09
N50	47856
N75	27585
L50	29
L75	58
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 1 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).



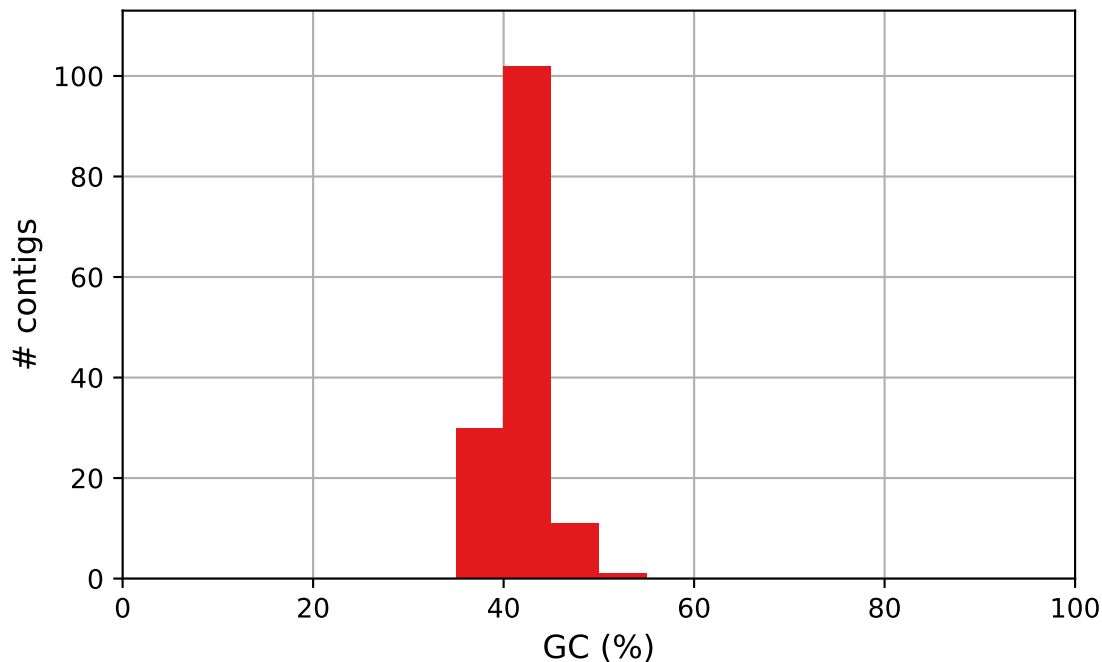
— MEGAHIT-MetaBAT2-3106384.5.fa





— MEGAHIT-MetaBAT2-3106384.5.fa

MEGAHIT-MetaBAT2-3106384.5.fa GC content



MEGAHit-MetaBAT2-3106384.5.fa