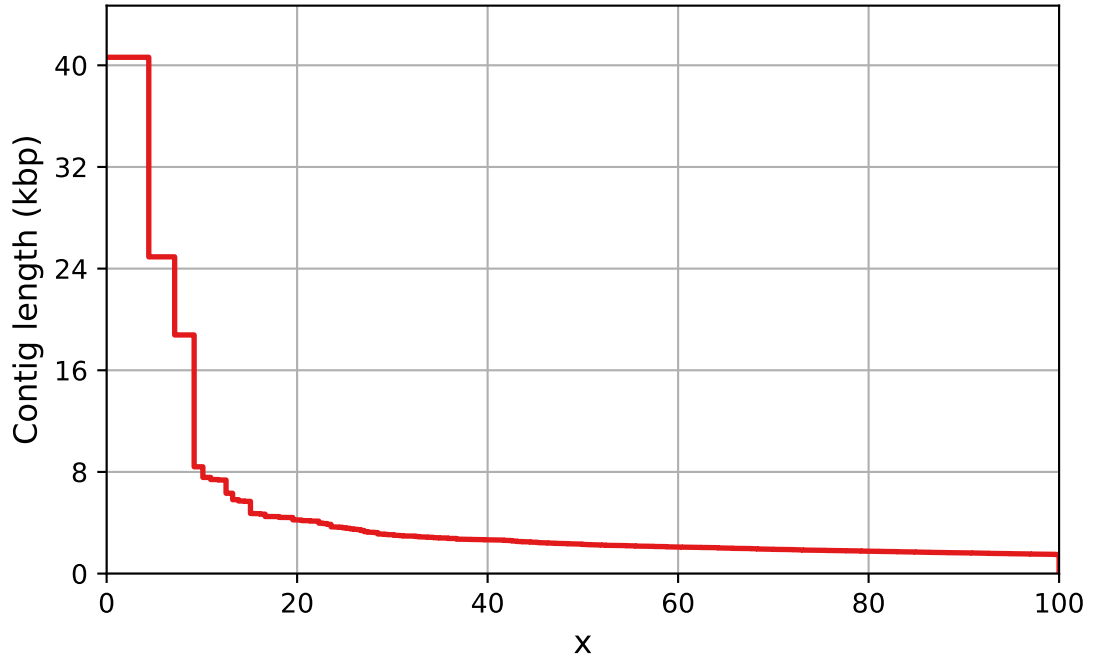


# Report

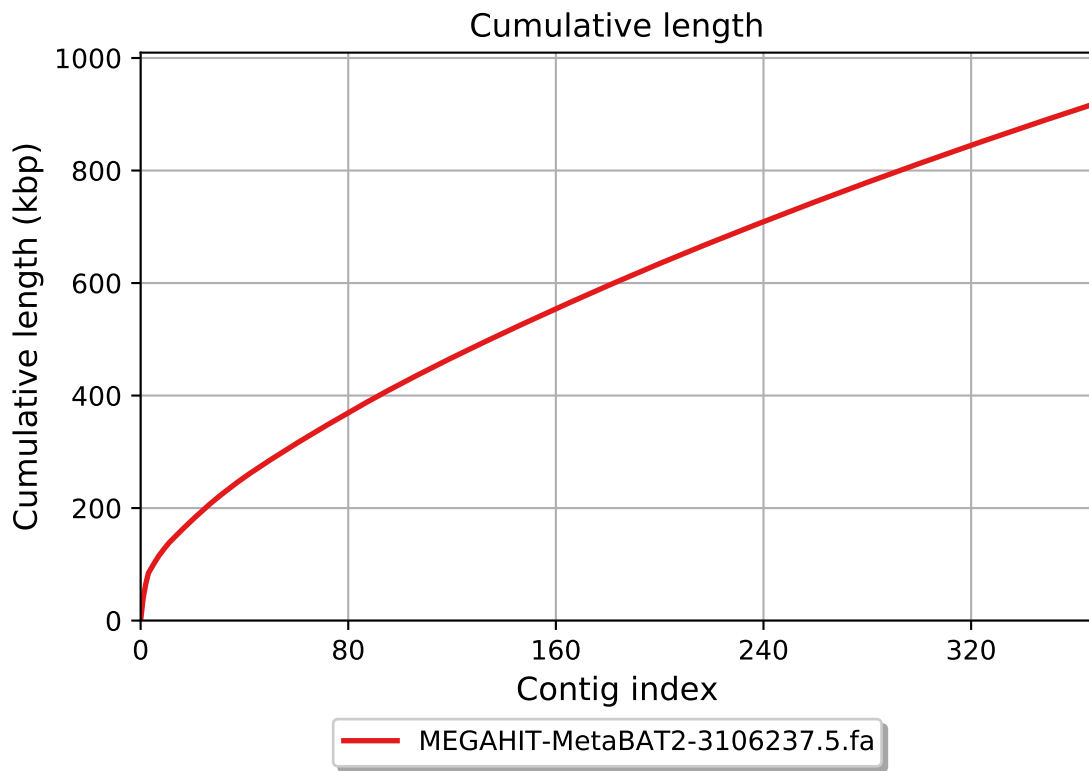
	MEGAHIT-MetaBAT2-3106237.5.fa
# contigs (>= 0 bp)	367
# contigs (>= 1000 bp)	367
# contigs (>= 5000 bp)	11
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	917780
Total length (>= 1000 bp)	917780
Total length (>= 5000 bp)	138607
Total length (>= 10000 bp)	84336
Total length (>= 25000 bp)	40631
Total length (>= 50000 bp)	0
# contigs	367
Largest contig	40631
Total length	917780
GC (%)	42.27
N50	2299
N75	1829
L50	117
L75	229
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 0 part

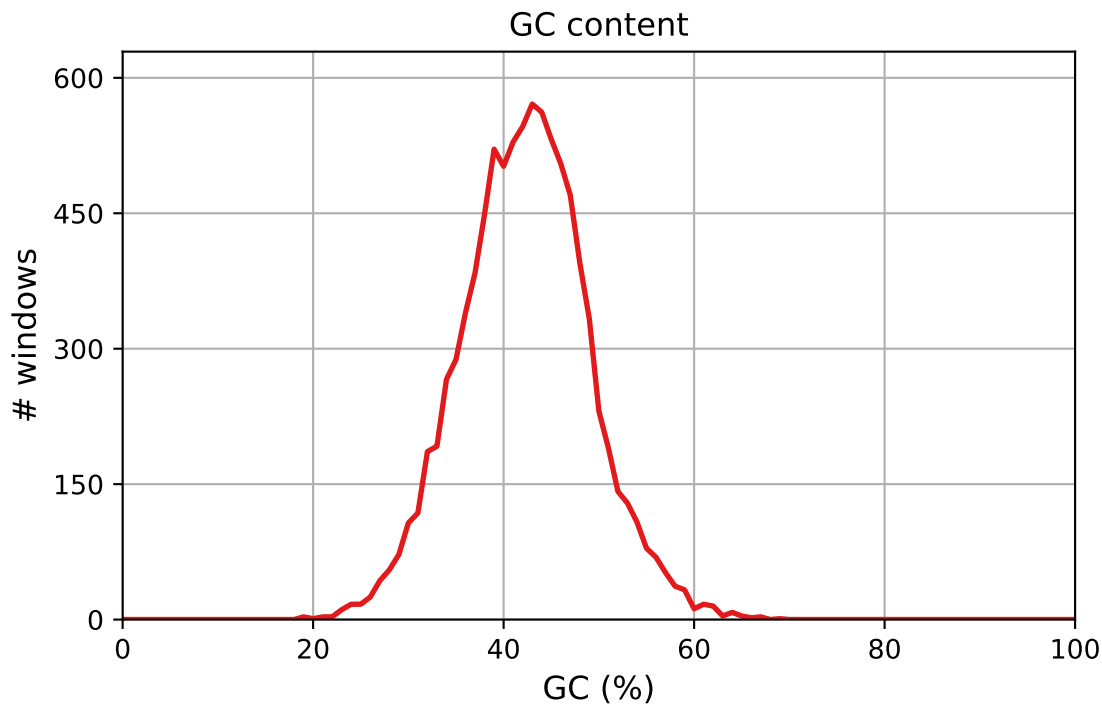
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



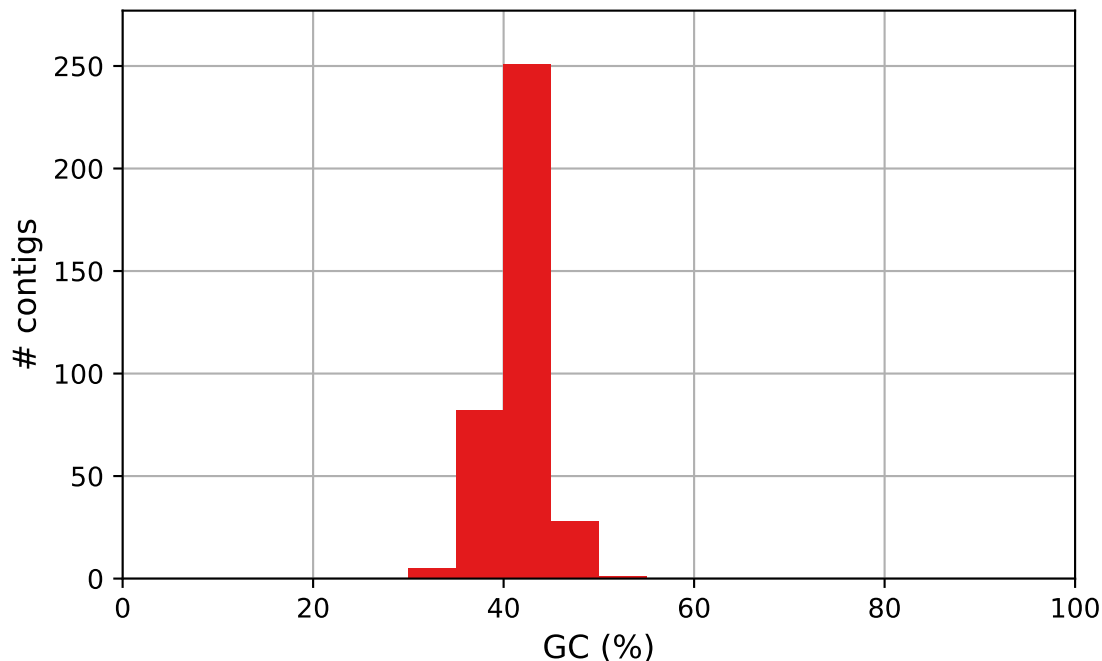
MEGAHIT-MetaBAT2-3106237.5.fa





— MEGAHit-MetaBAT2-3106237.5.fa

MEGAHIT-MetaBAT2-3106237.5.fa GC content



MEGAHit-MetaBAT2-3106237.5.fa