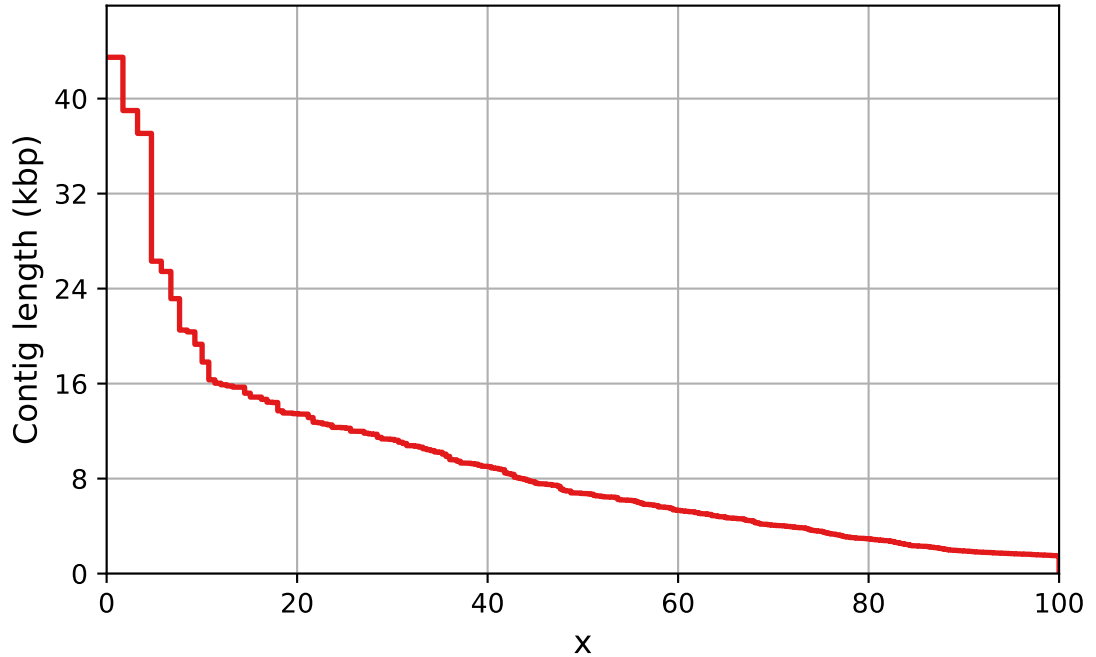


# Report

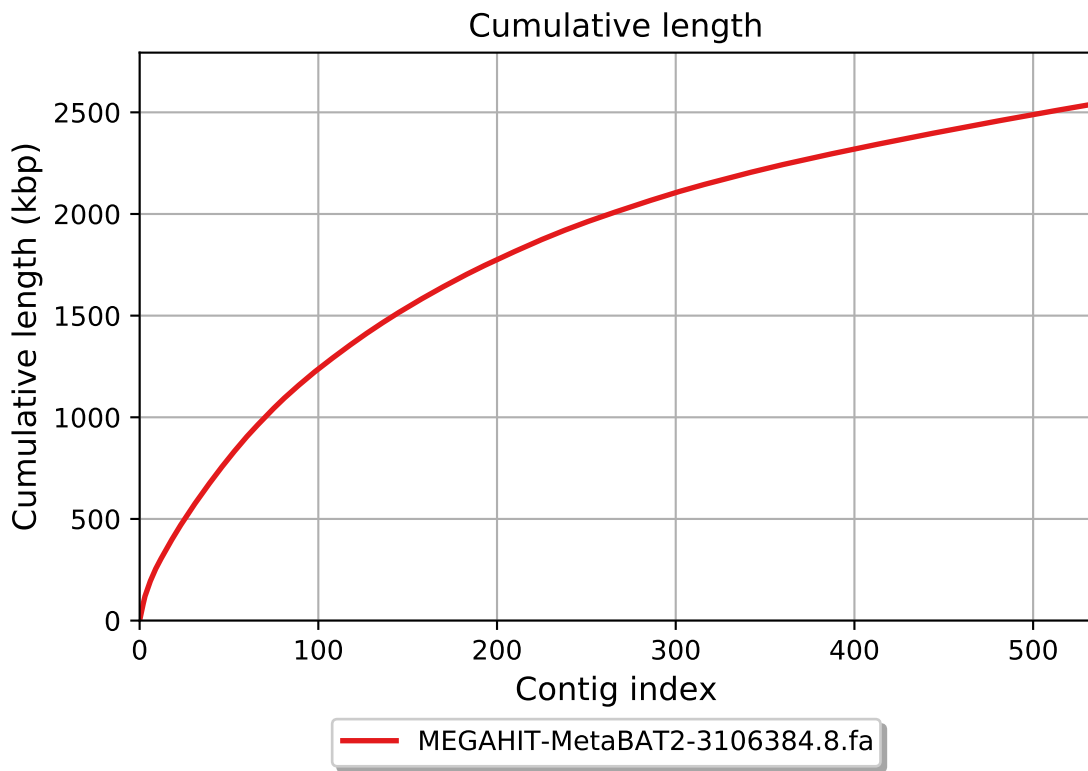
	MEGAHIT-MetaBAT2-3106384.8.fa
# contigs (>= 0 bp)	533
# contigs (>= 1000 bp)	533
# contigs (>= 5000 bp)	162
# contigs (>= 10000 bp)	60
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2539688
Total length (>= 1000 bp)	2539688
Total length (>= 5000 bp)	1603535
Total length (>= 10000 bp)	904923
Total length (>= 25000 bp)	171294
Total length (>= 50000 bp)	0
# contigs	533
Largest contig	43480
Total length	2539688
GC (%)	40.90
N50	6748
N75	3565
L50	105
L75	234
# N's per 100 kbp	0.00
# predicted rRNA genes	1 + 0 part

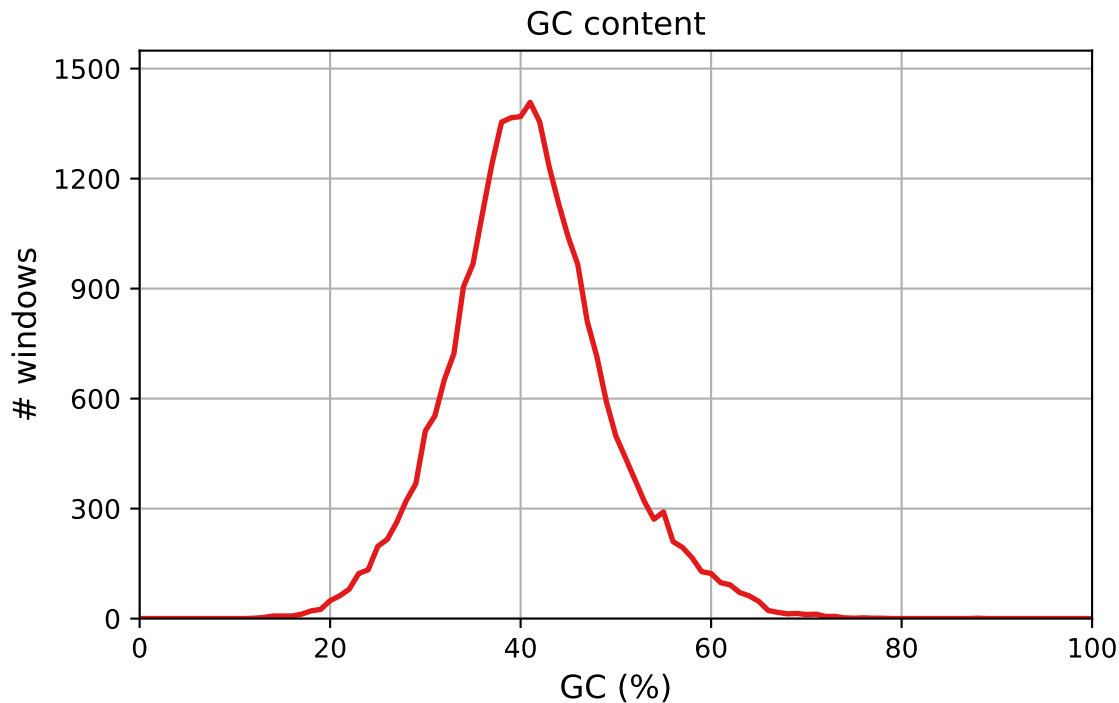
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



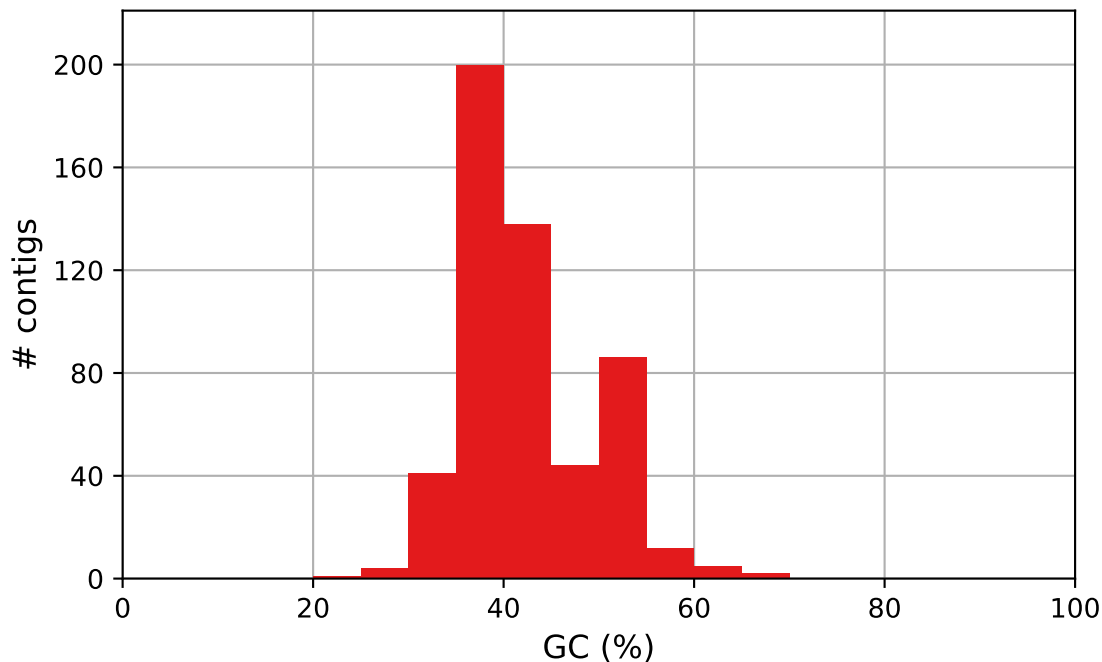
MEGAHIT-MetaBAT2-3106384.8.fa





— MEGAHit-MetaBAT2-3106384.8.fa

MEGAHIT-MetaBAT2-3106384.8.fa GC content



MEGAHit-MetaBAT2-3106384.8.fa