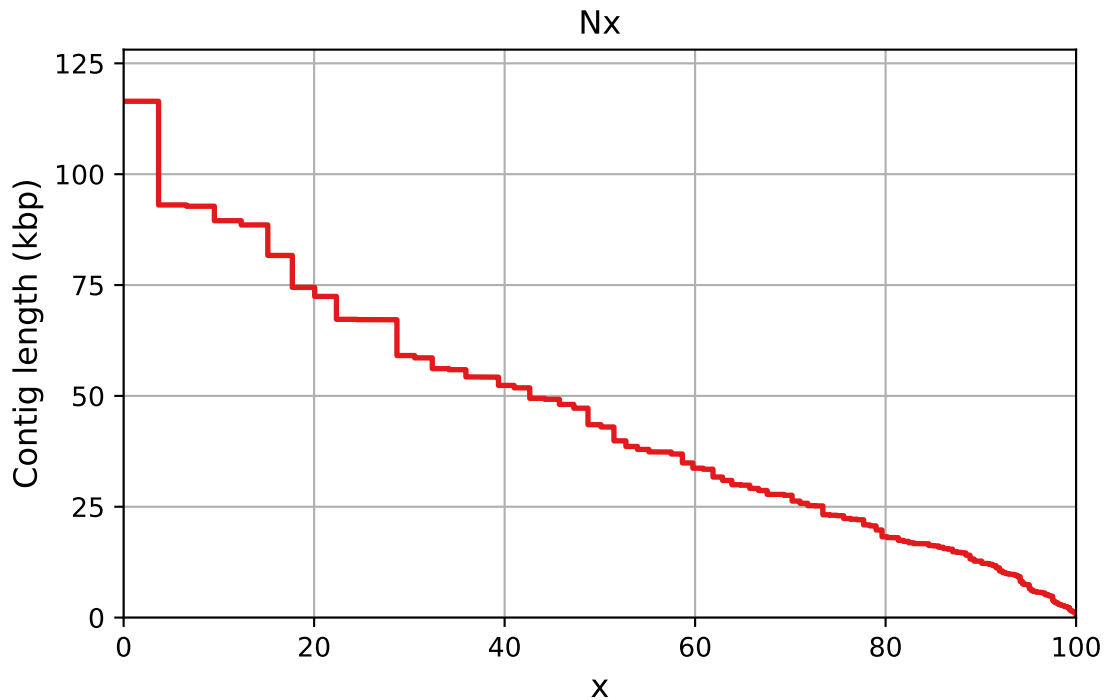


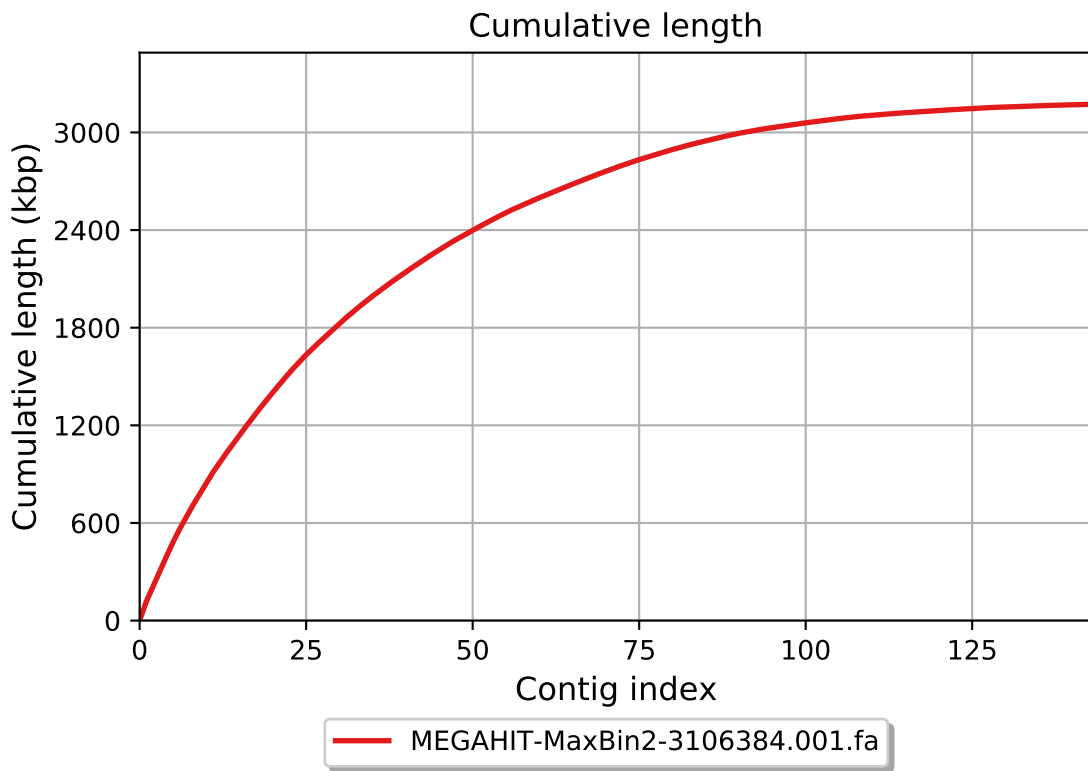
# Report

	MEGAHIT-MaxBin2-3106384.001.fa
# contigs (>= 0 bp)	143
# contigs (>= 1000 bp)	143
# contigs (>= 5000 bp)	104
# contigs (>= 10000 bp)	84
# contigs (>= 25000 bp)	47
# contigs (>= 50000 bp)	19
Total length (>= 0 bp)	3173138
Total length (>= 1000 bp)	3173138
Total length (>= 5000 bp)	3080331
Total length (>= 10000 bp)	2938581
Total length (>= 25000 bp)	2329413
Total length (>= 50000 bp)	1353002
# contigs	143
Largest contig	116447
Total length	3173138
GC (%)	42.22
N50	43503
N75	22989
L50	24
L75	50
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 2 part

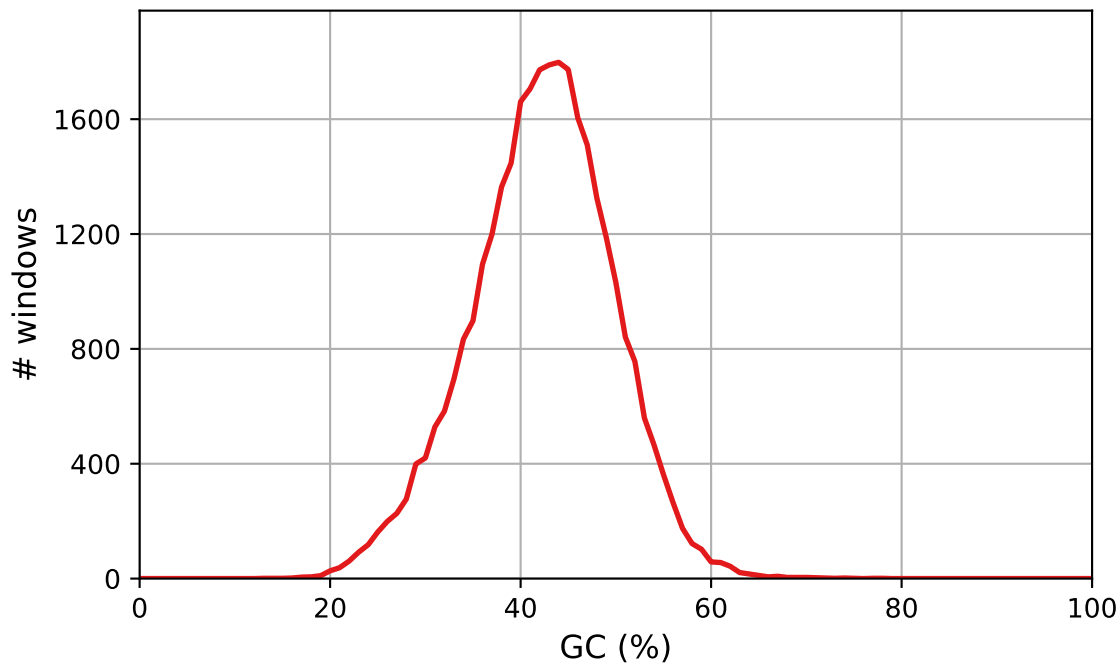
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



MEGAHIT-MaxBin2-3106384.001.fa

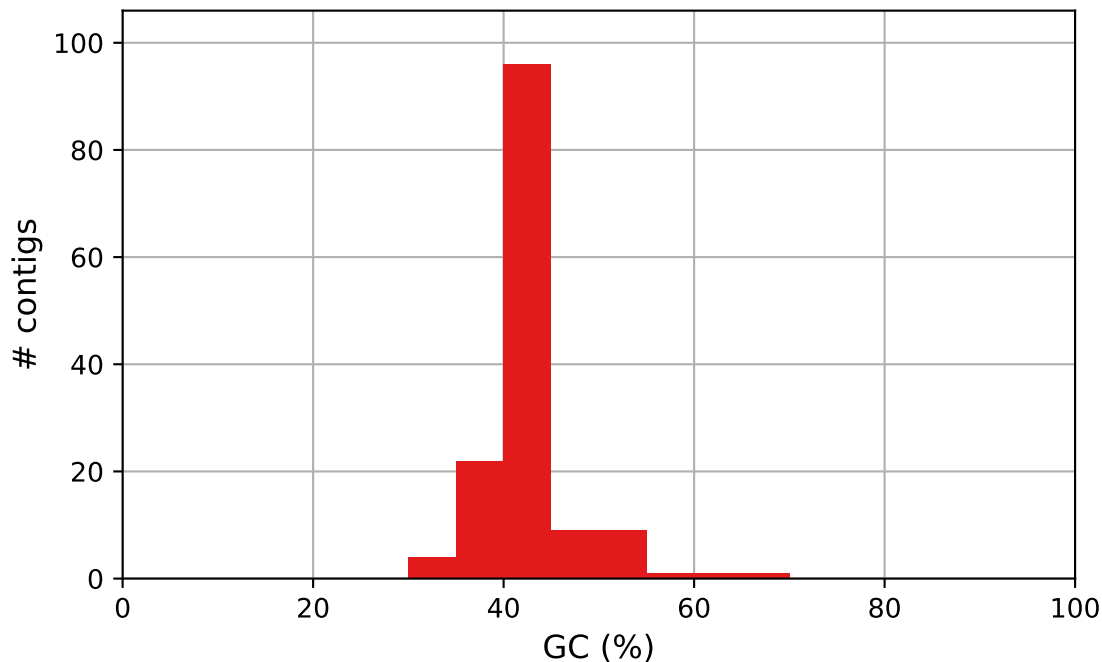


# GC content



MEGAHIT-MaxBin2-3106384.001.fa

MEGAHIT-MaxBin2-3106384.001.fa GC content



MEGAHit-MaxBin2-3106384.001.fa