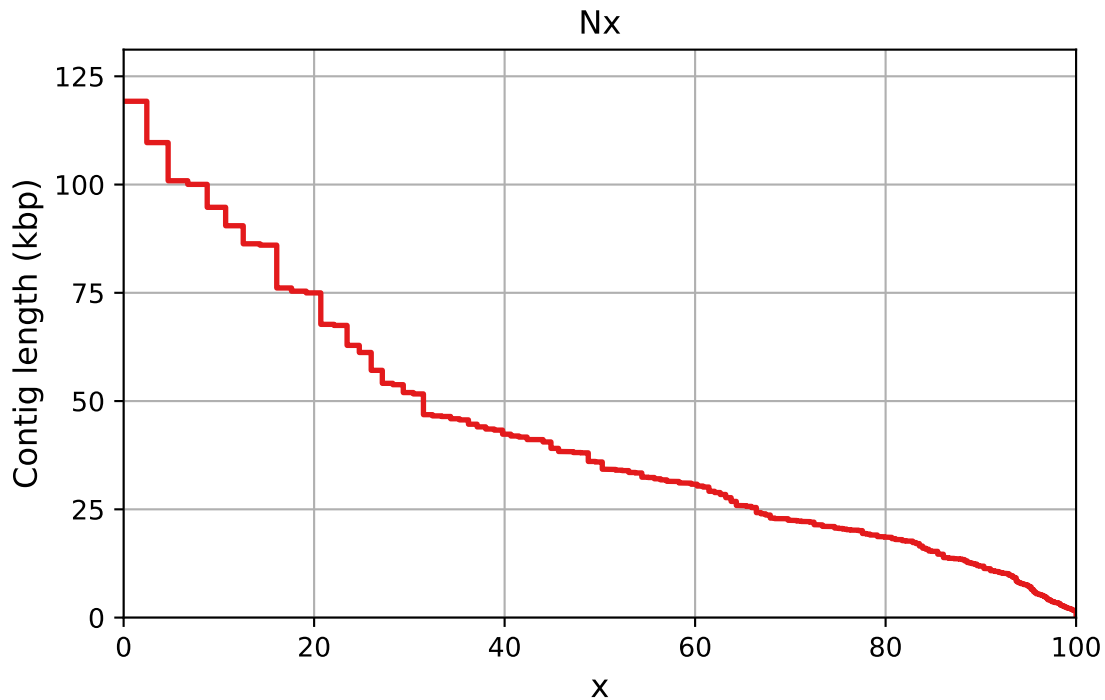
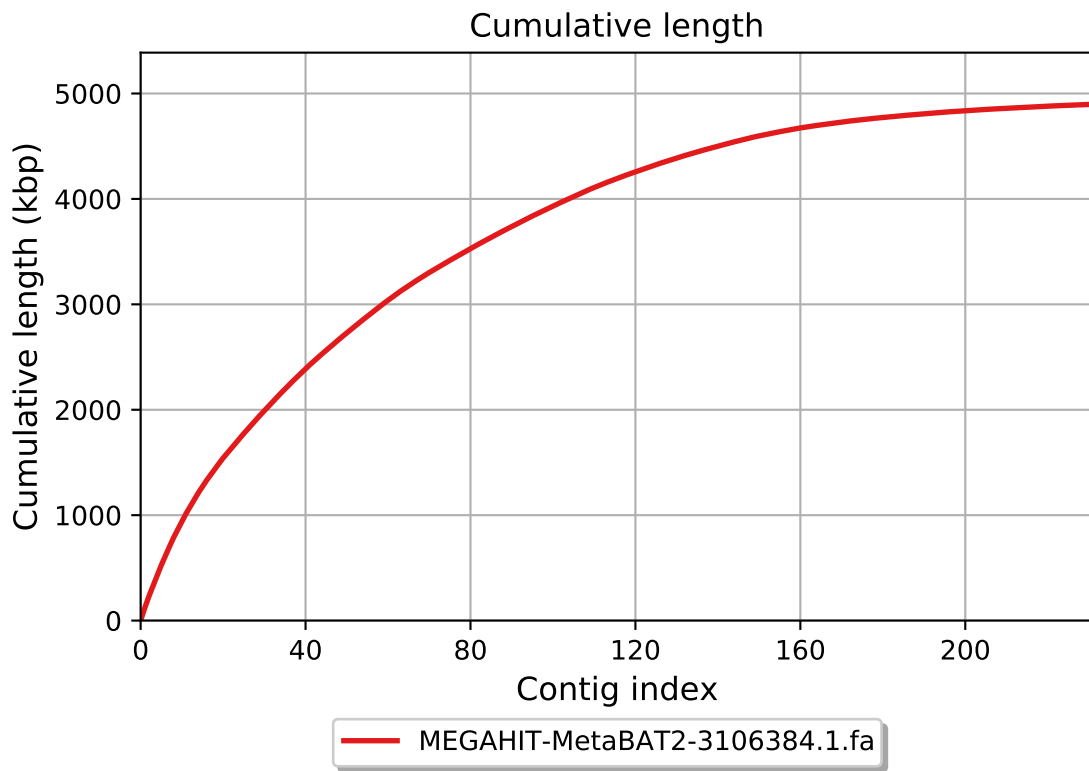


Report

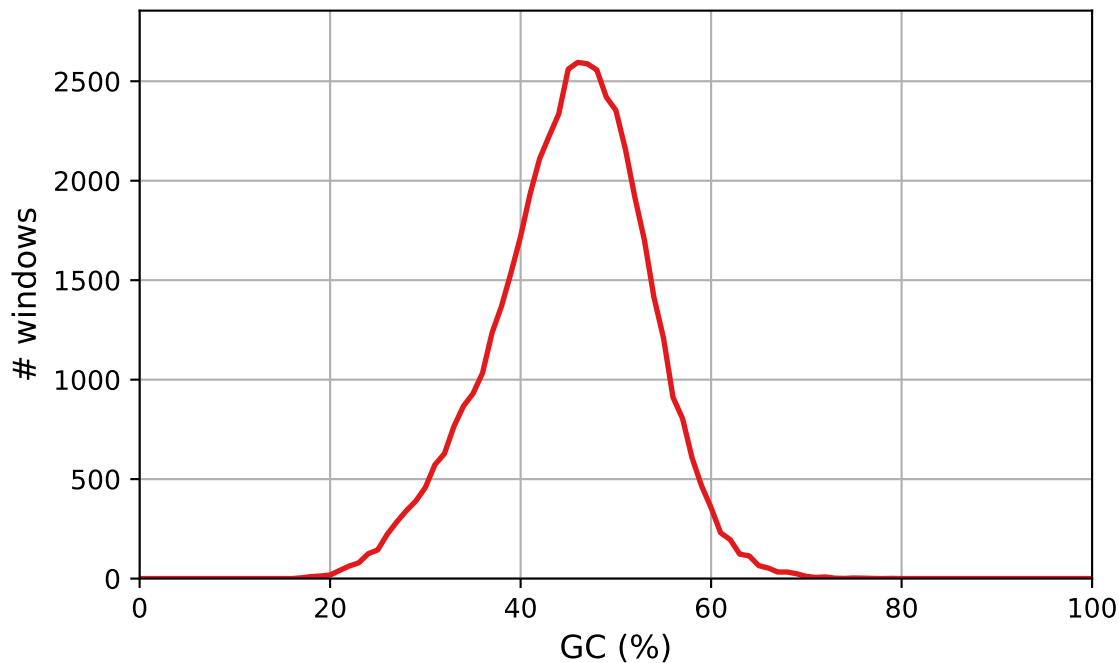
	MEGAHIT-MetaBAT2-3106384.1.fa
# contigs (>= 0 bp)	231
# contigs (>= 1000 bp)	231
# contigs (>= 5000 bp)	170
# contigs (>= 10000 bp)	145
# contigs (>= 25000 bp)	68
# contigs (>= 50000 bp)	20
Total length (>= 0 bp)	4897843
Total length (>= 1000 bp)	4897843
Total length (>= 5000 bp)	4728989
Total length (>= 10000 bp)	4551844
Total length (>= 25000 bp)	3253496
Total length (>= 50000 bp)	1541742
# contigs	231
Largest contig	119232
Total length	4897843
GC (%)	45.15
N50	35936
N75	20667
L50	42
L75	87
# N's per 100 kbp	0.00
# predicted rRNA genes	1 + 6 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).



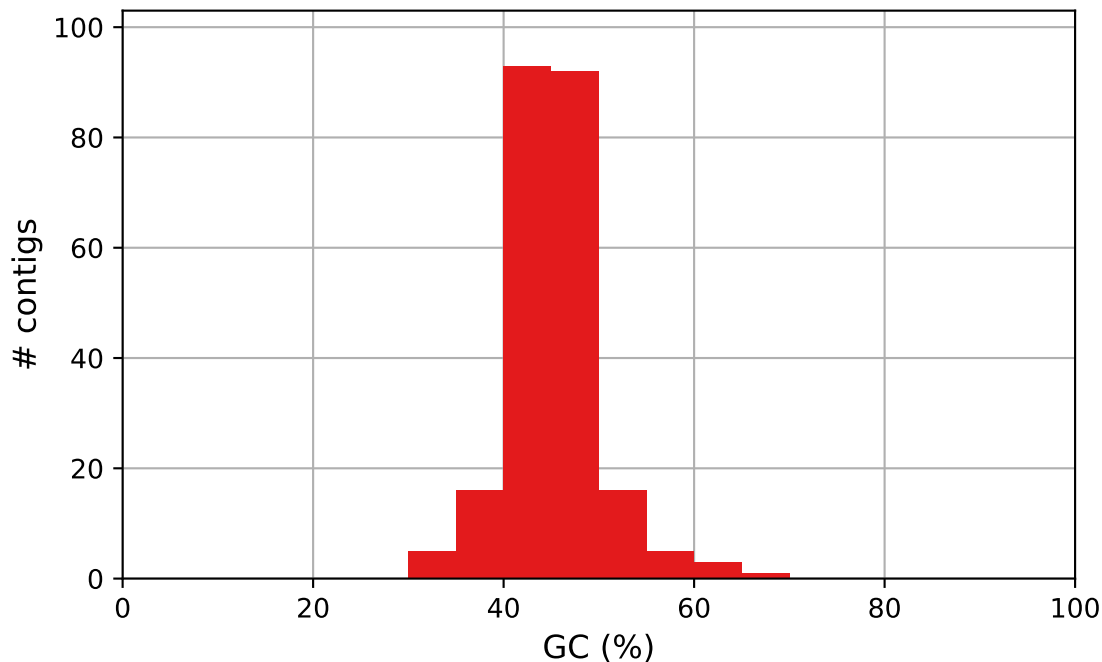


GC content



— MEGAHIT-MetaBAT2-3106384.1.fa

MEGAHIT-MetaBAT2-3106384.1.fa GC content



MEGAhit-MetaBAT2-3106384.1.fa