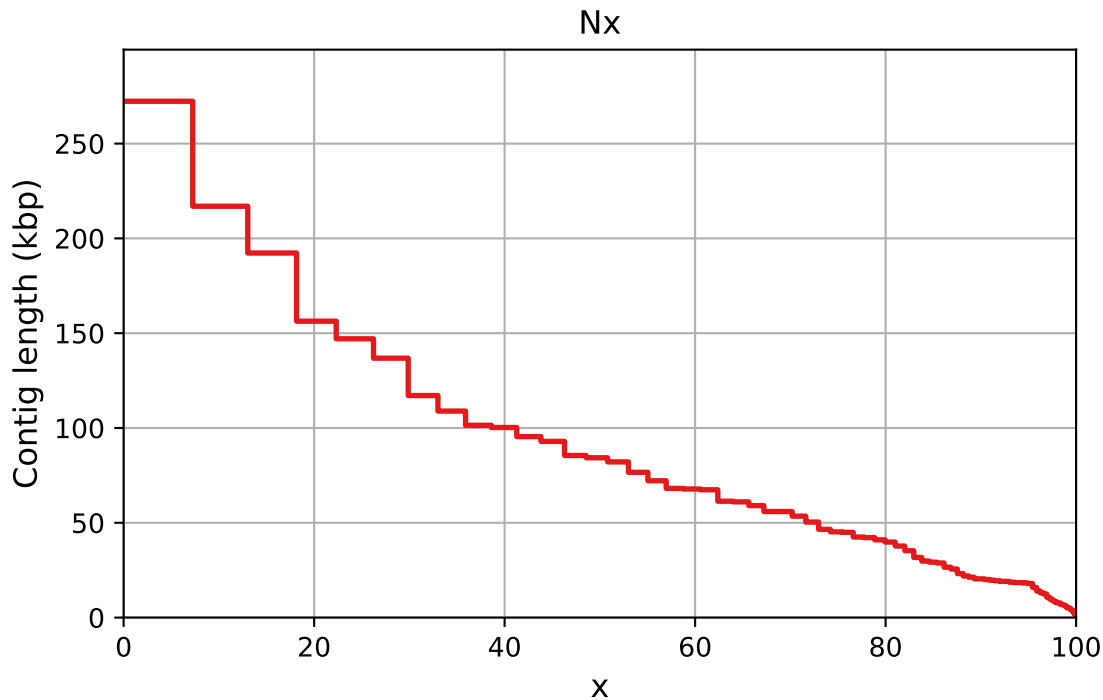


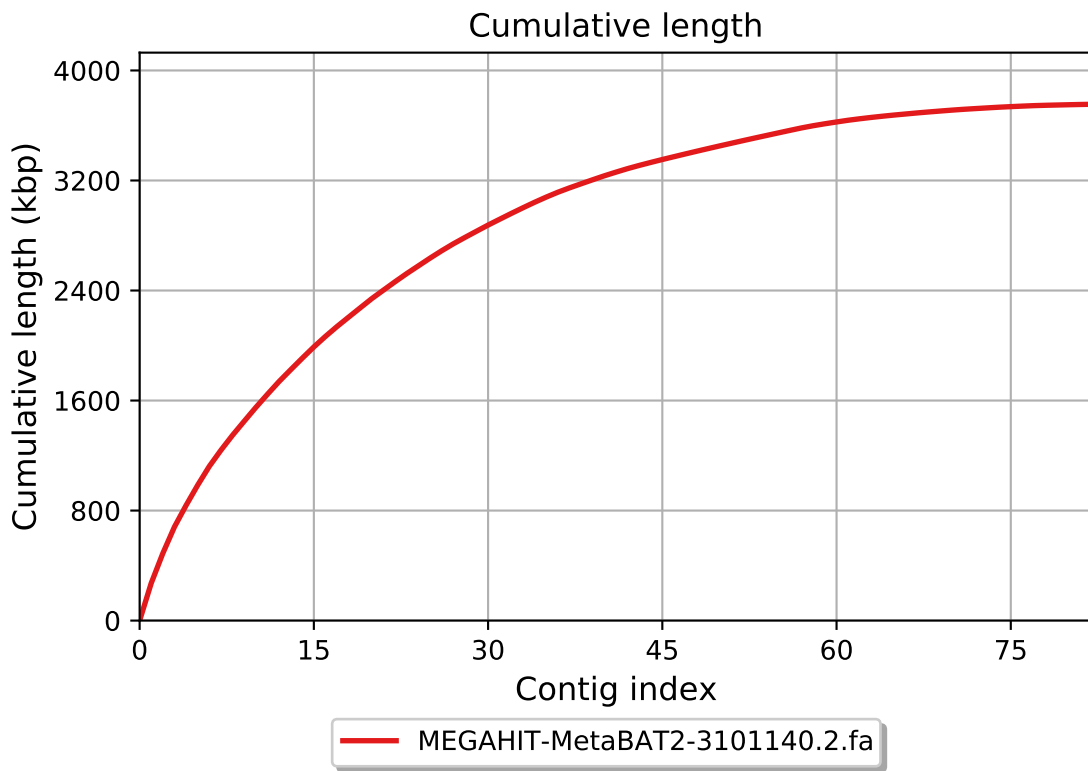
Report

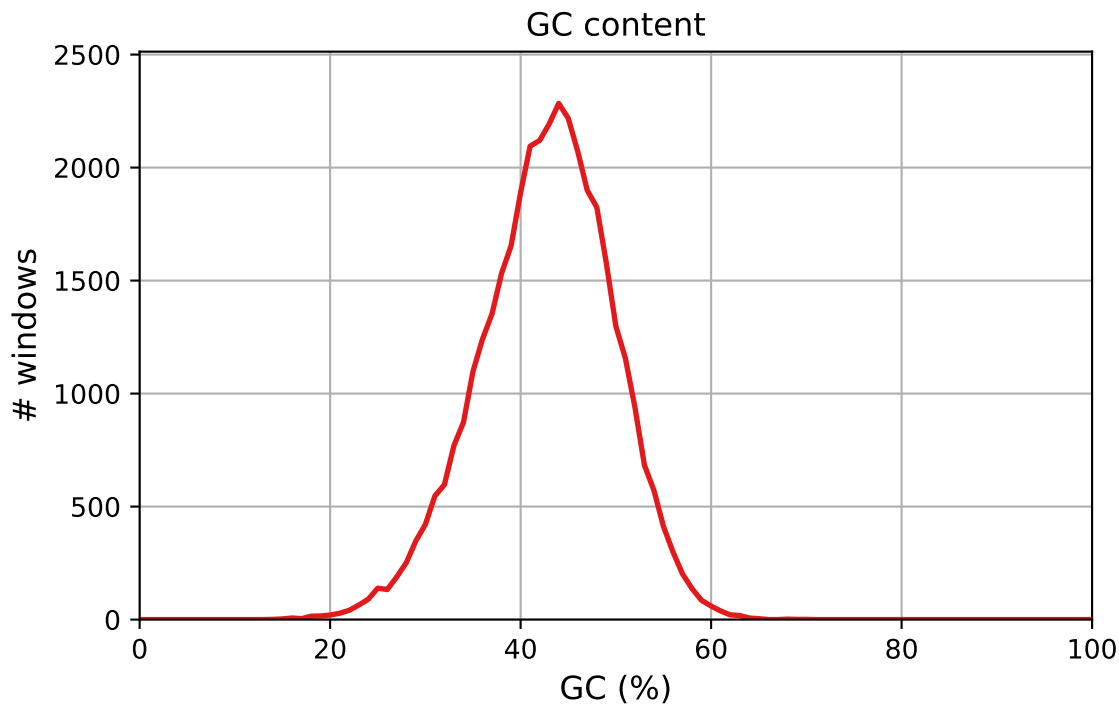
	MEGAHIT-MetaBAT2-3101140.2.fa
# contigs (>= 0 bp)	82
# contigs (>= 1000 bp)	82
# contigs (>= 5000 bp)	72
# contigs (>= 10000 bp)	62
# contigs (>= 25000 bp)	42
# contigs (>= 50000 bp)	27
Total length (>= 0 bp)	3754182
Total length (>= 1000 bp)	3754182
Total length (>= 5000 bp)	3723043
Total length (>= 10000 bp)	3648788
Total length (>= 25000 bp)	3286051
Total length (>= 50000 bp)	2739103
# contigs	82
Largest contig	272320
Total length	3754182
GC (%)	42.69
N50	84297
N75	45215
L50	14
L75	29
# 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).



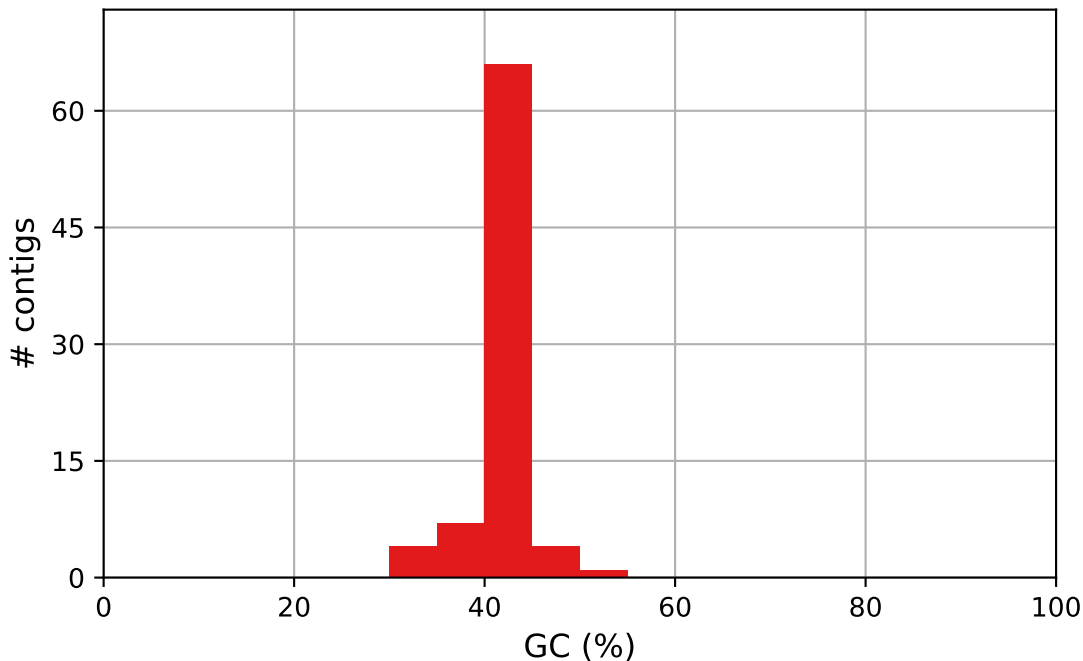
— MEGAHIT-MetaBAT2-3101140.2.fa





— MEGAHit-MetaBAT2-3101140.2.fa

MEGAHIT-MetaBAT2-3101140.2.fa GC content



MEGAHIT-MetaBAT2-3101140.2.fa