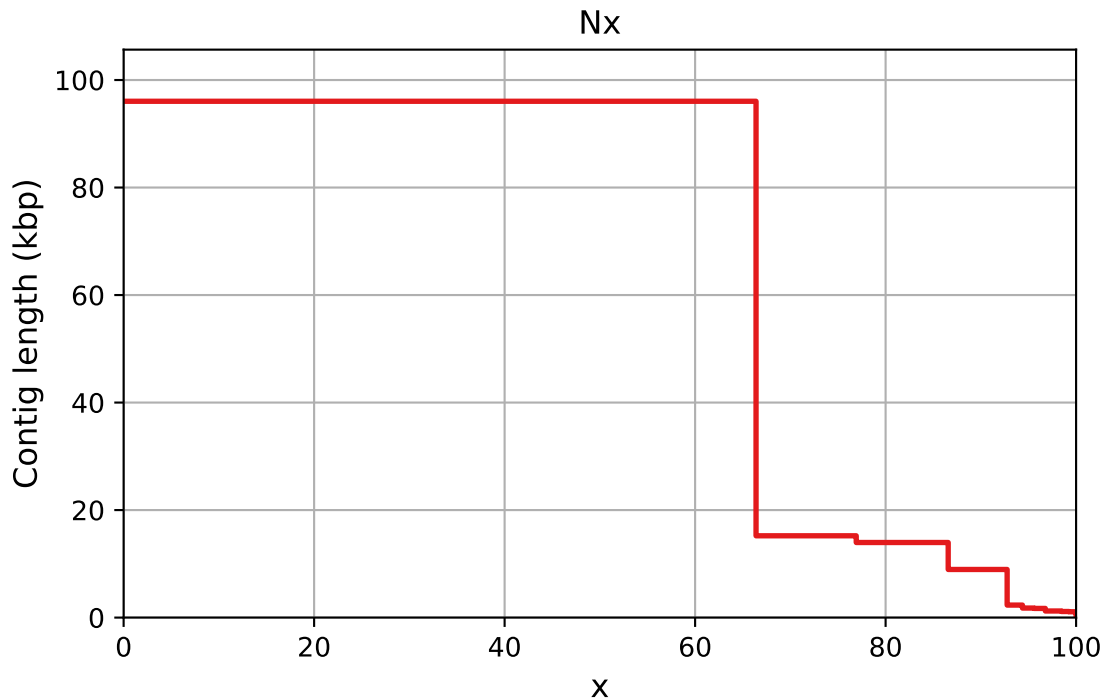


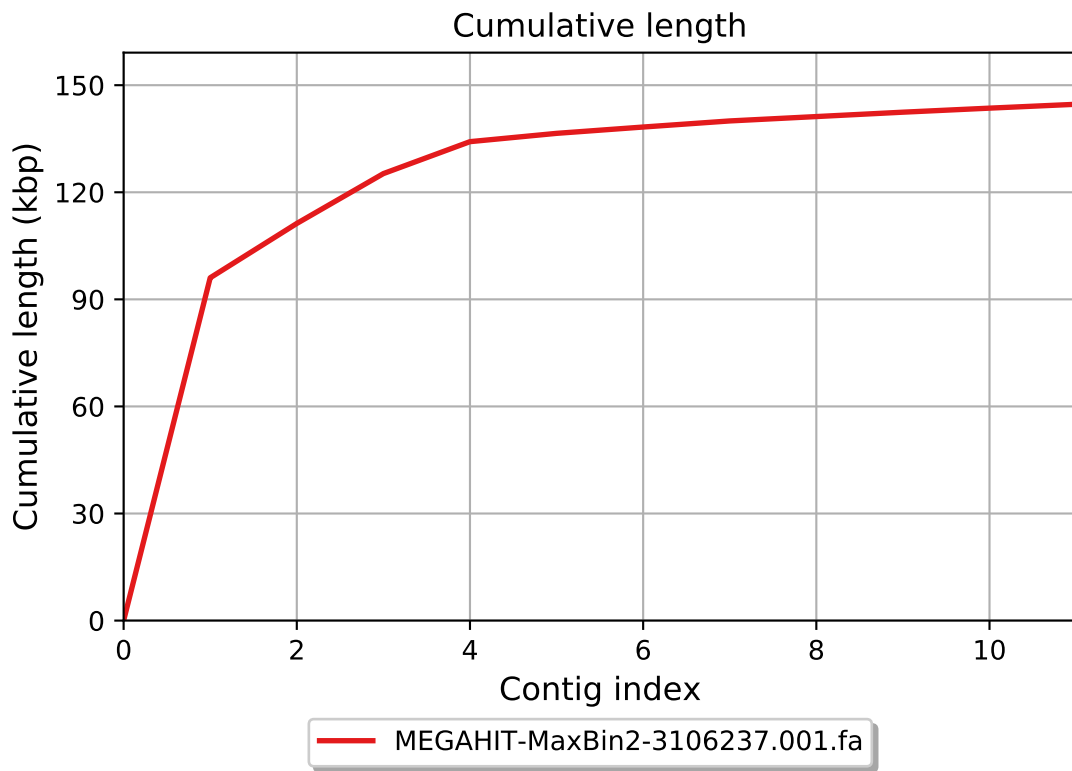
Report

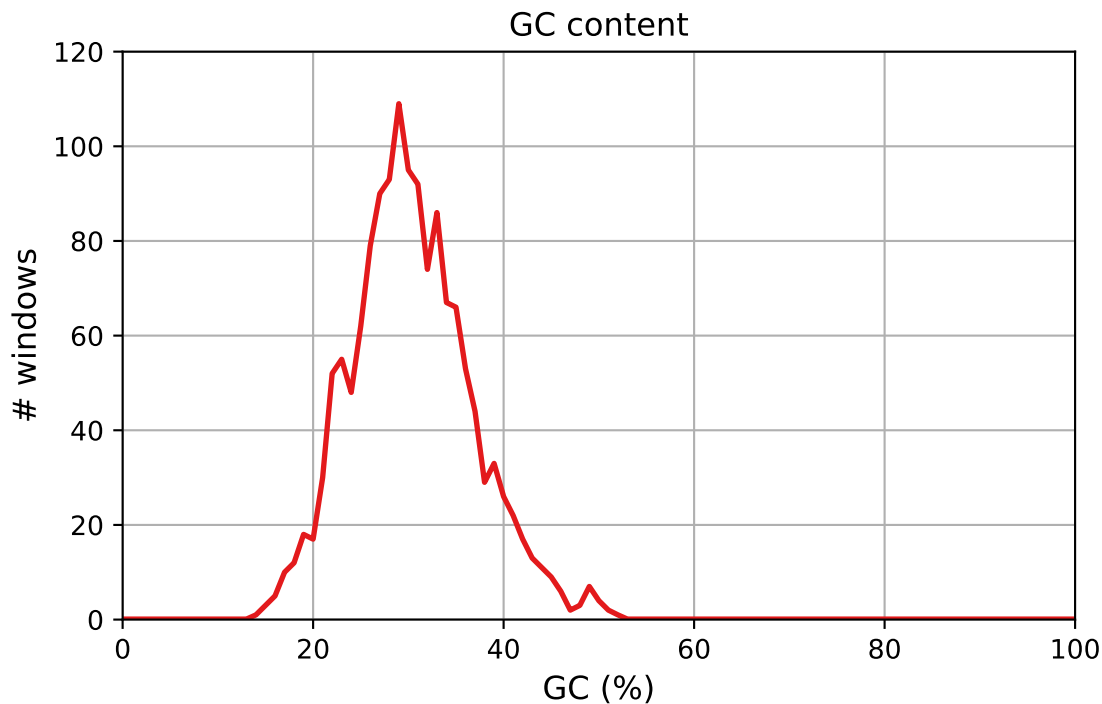
	MEGAHIT-MaxBin2-3106237.001.fa
# contigs (>= 0 bp)	11
# contigs (>= 1000 bp)	11
# contigs (>= 5000 bp)	4
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	144655
Total length (>= 1000 bp)	144655
Total length (>= 5000 bp)	134182
Total length (>= 10000 bp)	125231
Total length (>= 25000 bp)	96047
Total length (>= 50000 bp)	96047
# contigs	11
Largest contig	96047
Total length	144655
GC (%)	30.38
N50	96047
N75	15216
L50	1
L75	2
# 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).



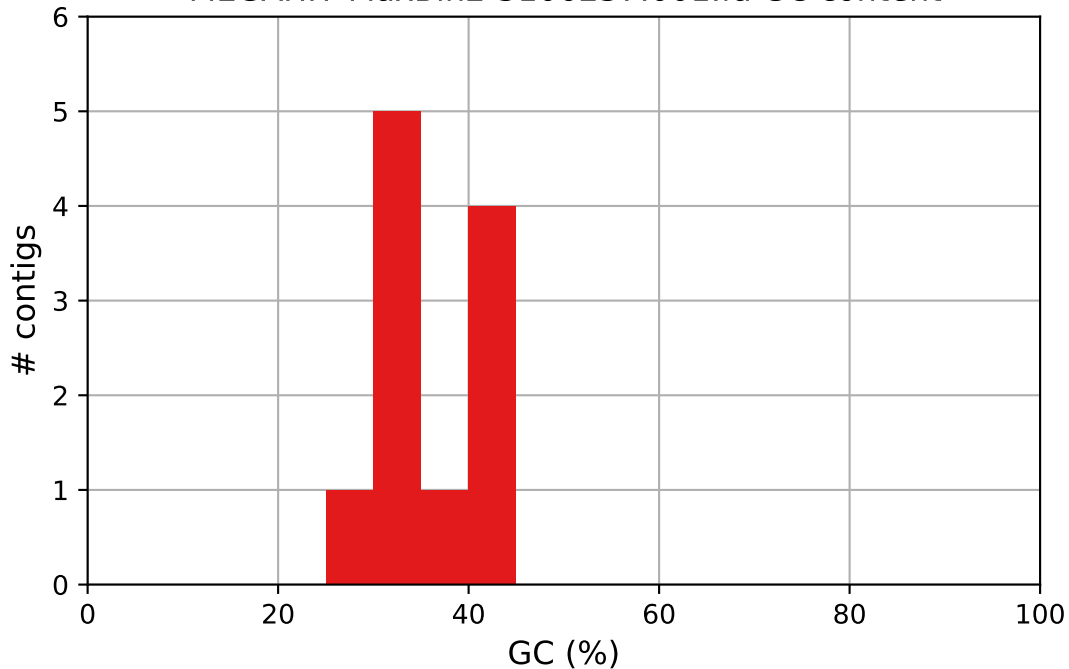
MEGAHIT-MaxBin2-3106237.001.fa





MEGAHit-MaxBin2-3106237.001.fa

MEGAHIT-MaxBin2-3106237.001.fa GC content



MEGAHit-MaxBin2-3106237.001.fa