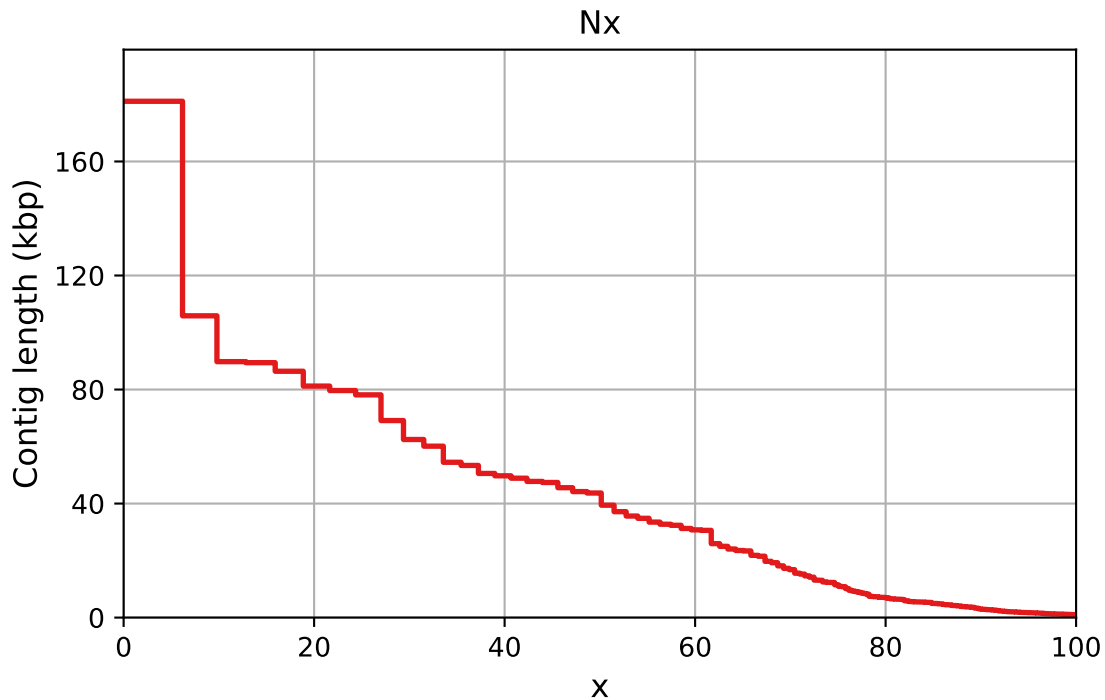


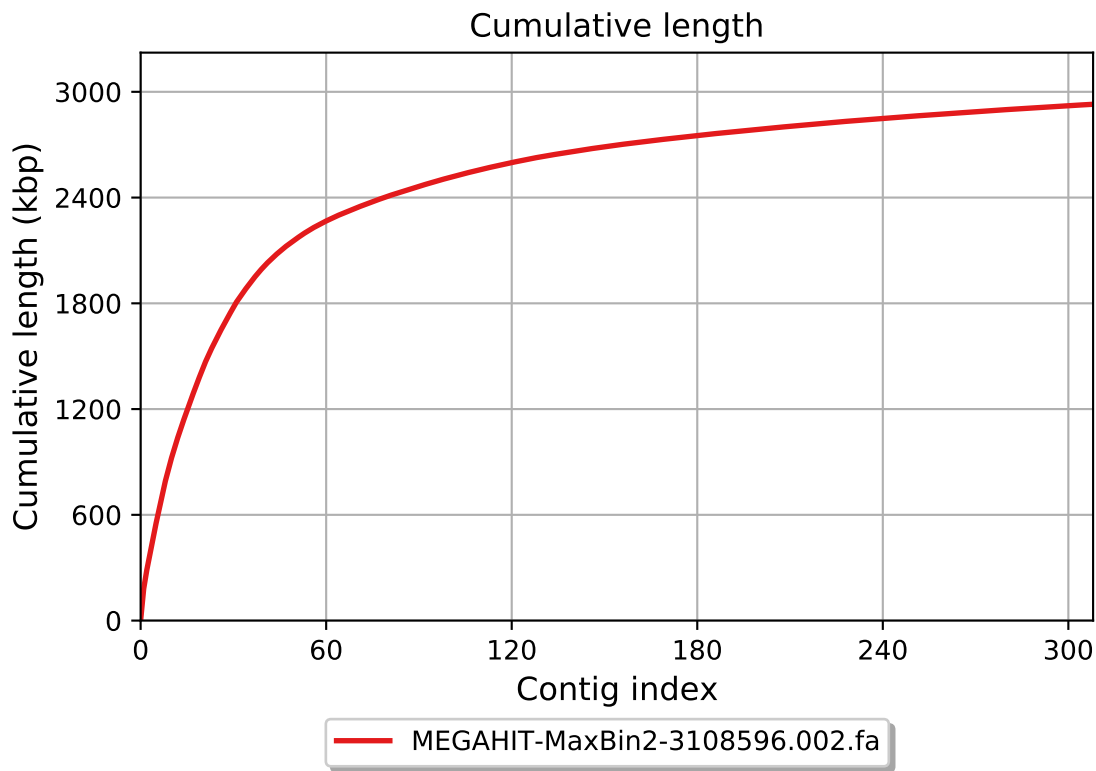
Report

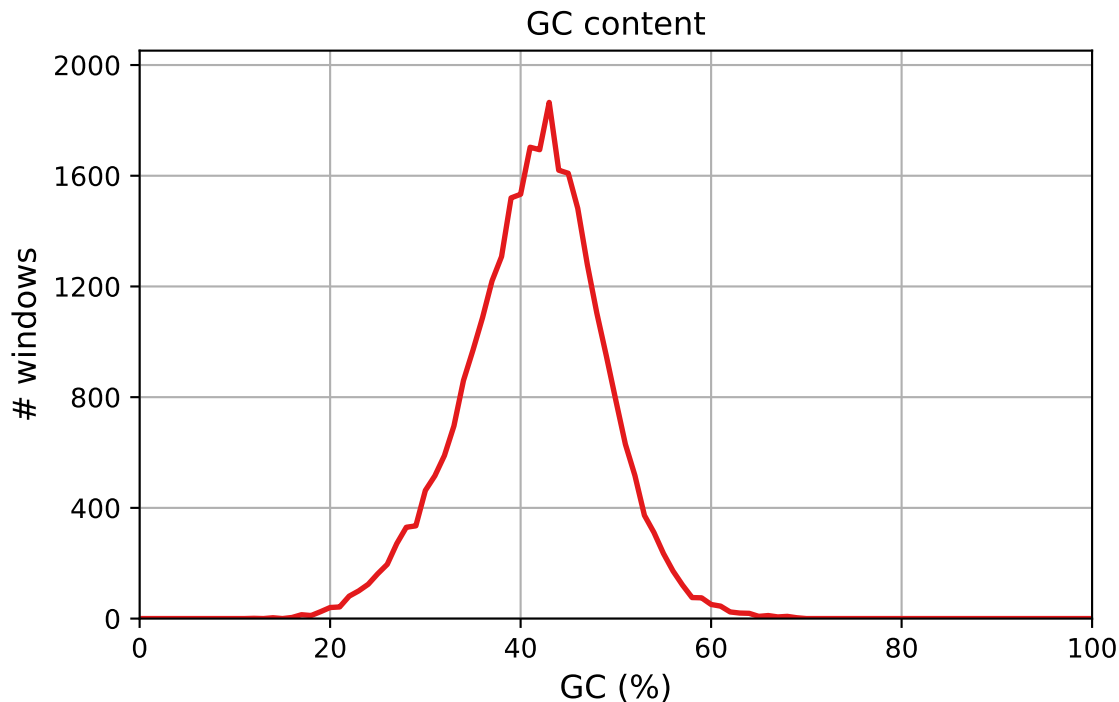
	MEGAHIT-MaxBin2-3108596.002.fa
# contigs (>= 0 bp)	308
# contigs (>= 1000 bp)	308
# contigs (>= 5000 bp)	94
# contigs (>= 10000 bp)	56
# contigs (>= 25000 bp)	32
# contigs (>= 50000 bp)	14
Total length (>= 0 bp)	2929362
Total length (>= 1000 bp)	2929362
Total length (>= 5000 bp)	2484942
Total length (>= 10000 bp)	2230259
Total length (>= 25000 bp)	1833199
Total length (>= 50000 bp)	1141731
# contigs	308
Largest contig	181120
Total length	2929362
GC (%)	41.40
N50	43700
N75	11510
L50	21
L75	53
# 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).



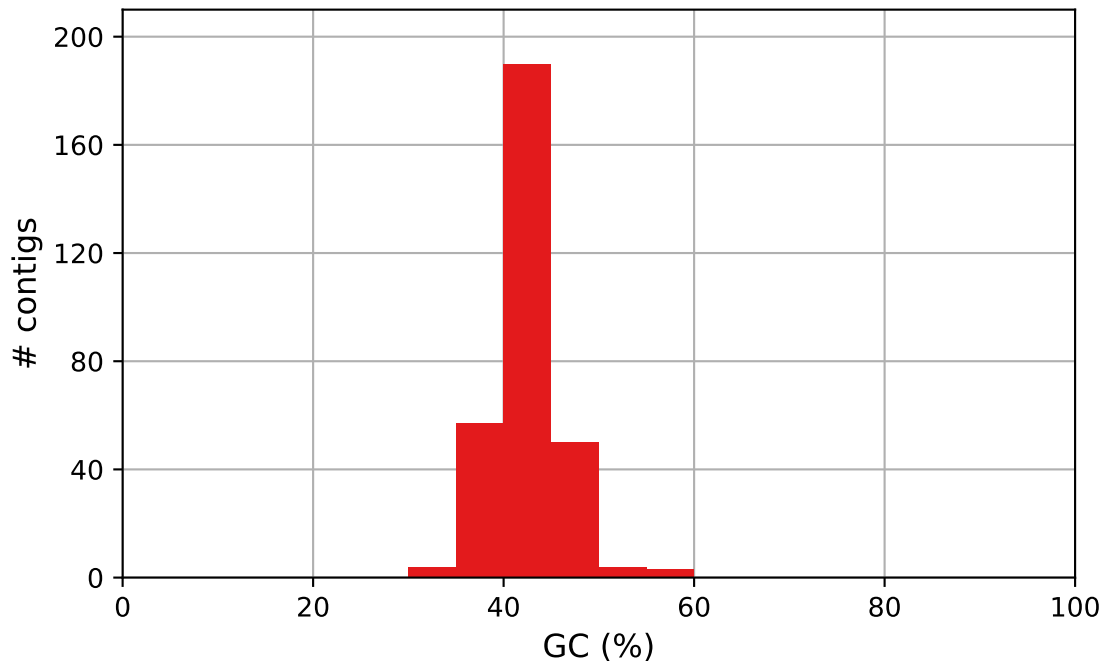
MEGAHIT-MaxBin2-3108596.002.fa





MEGAHit-MaxBin2-3108596.002.fa

MEGAHIT-MaxBin2-3108596.002.fa GC content



MEGAHit-MaxBin2-3108596.002.fa