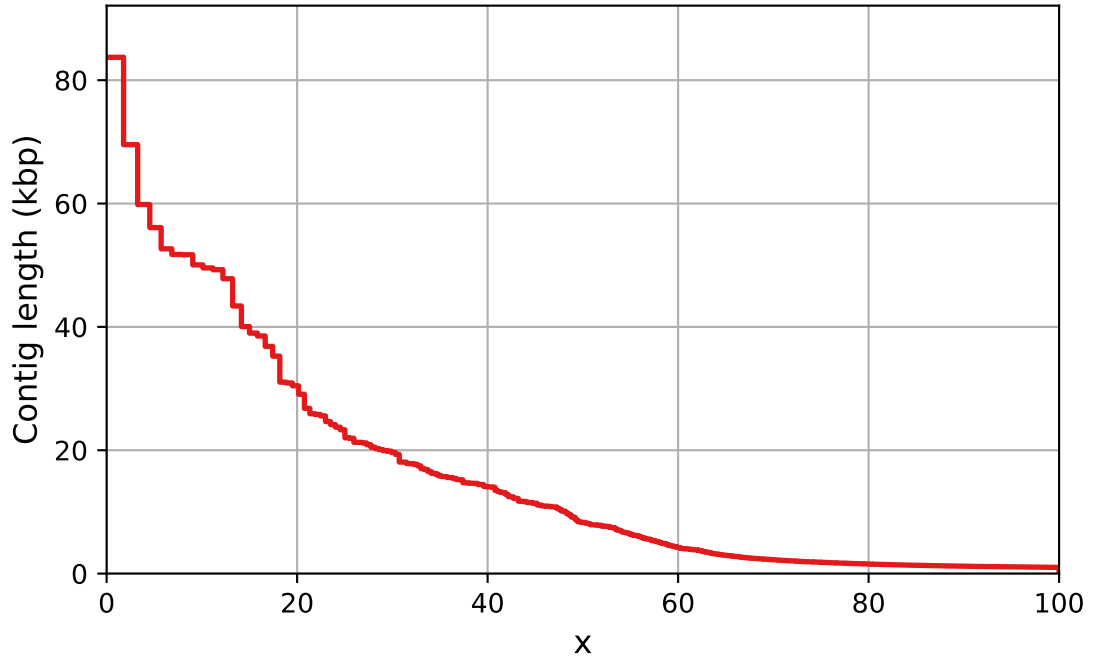


Report

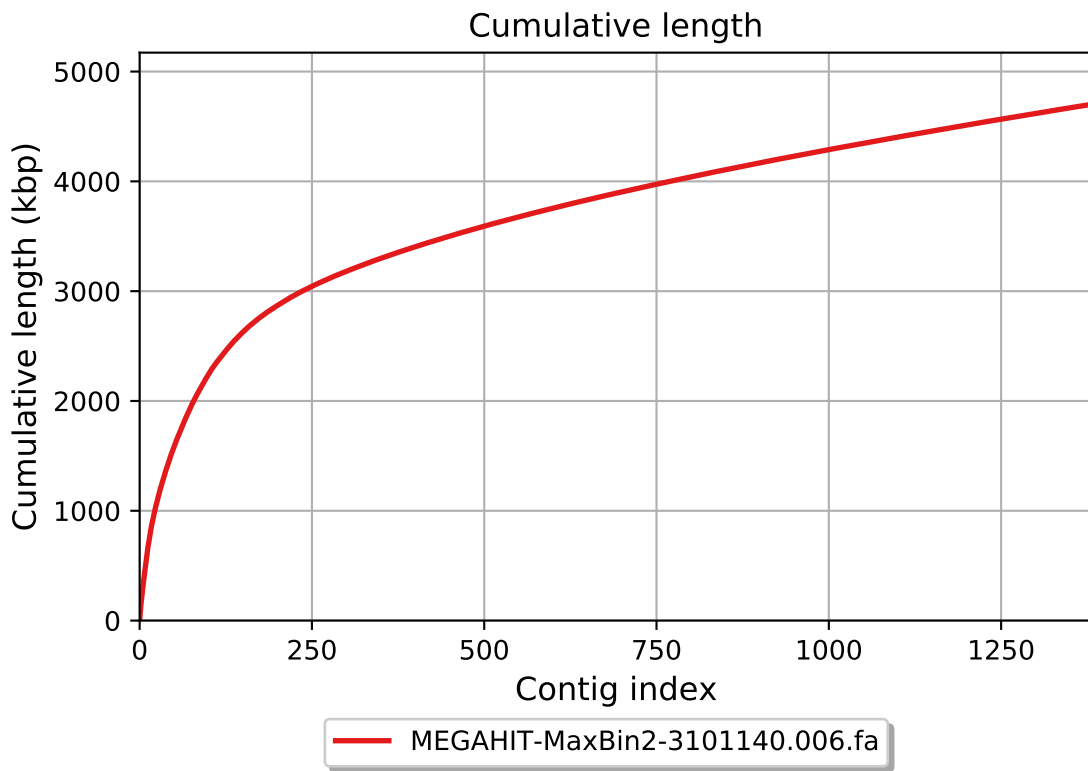
	MEGAHIT-MaxBin2-3101140.006.fa
# contigs (>= 0 bp)	1382
# contigs (>= 1000 bp)	1382
# contigs (>= 5000 bp)	170
# contigs (>= 10000 bp)	102
# contigs (>= 25000 bp)	25
# contigs (>= 50000 bp)	8
Total length (>= 0 bp)	4702080
Total length (>= 1000 bp)	4702080
Total length (>= 5000 bp)	2738484
Total length (>= 10000 bp)	2264691
Total length (>= 25000 bp)	1080577
Total length (>= 50000 bp)	475348
# contigs	1382
Largest contig	83717
Total length	4702080
GC (%)	42.66
N50	8284
N75	1832
L50	112
L75	464
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 2 part

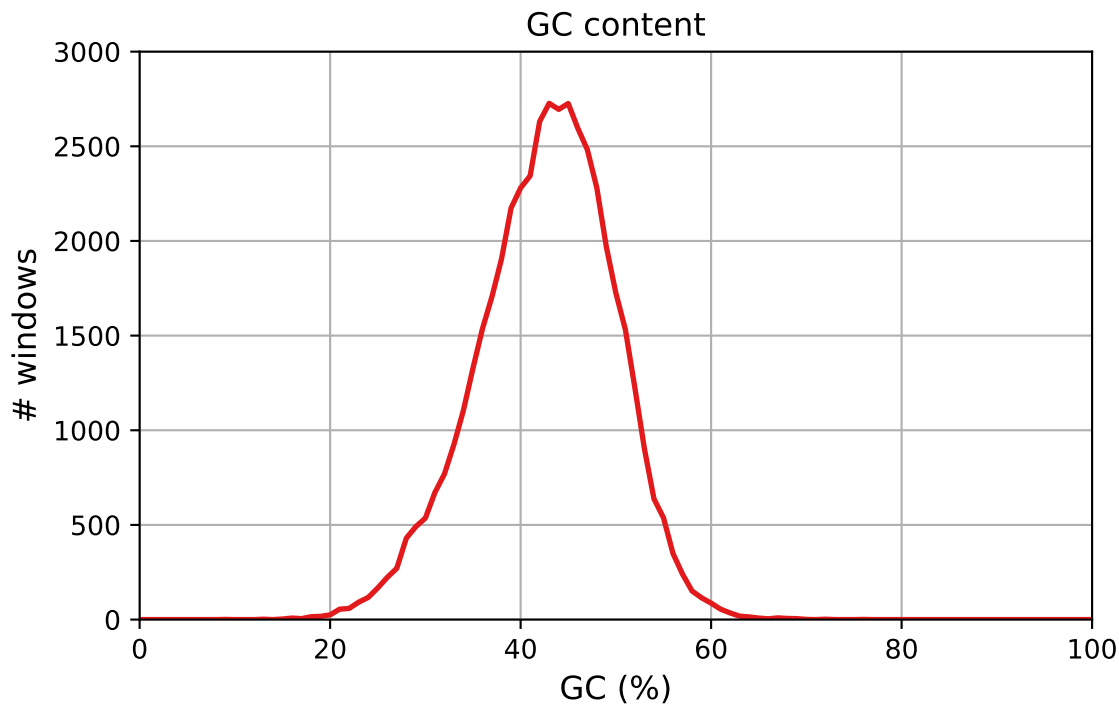
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



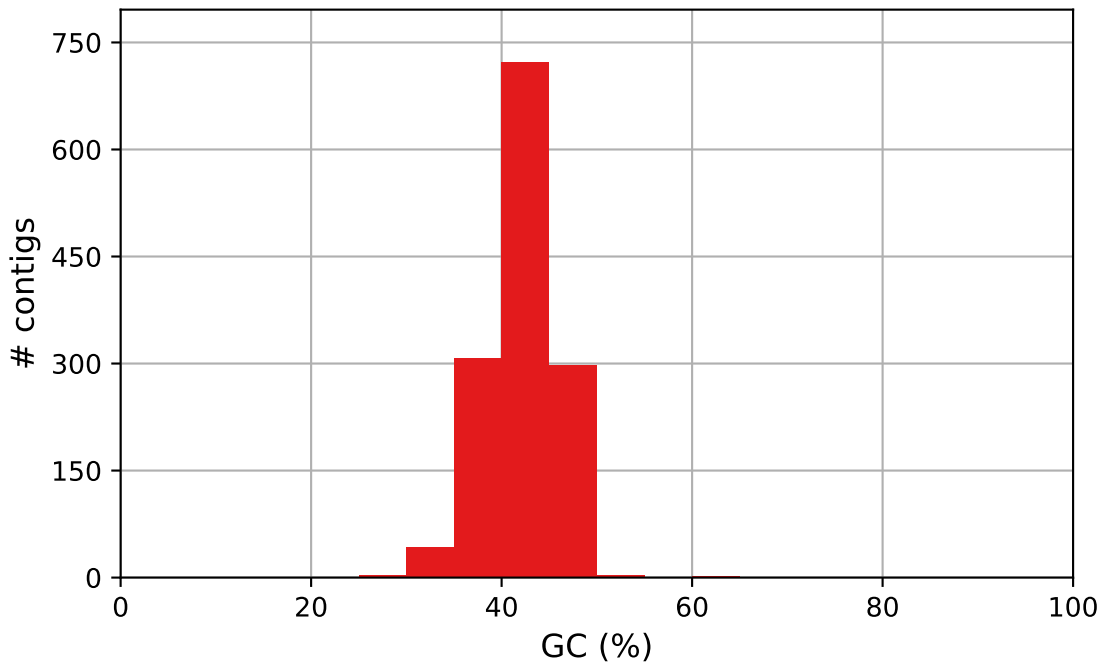
MEGAHIT-MaxBin2-3101140.006.fa





MEGAHIT-MaxBin2-3101140.006.fa

MEGAHIT-MaxBin2-3101140.006.fa GC content



MEGAHit-MaxBin2-3101140.006.fa