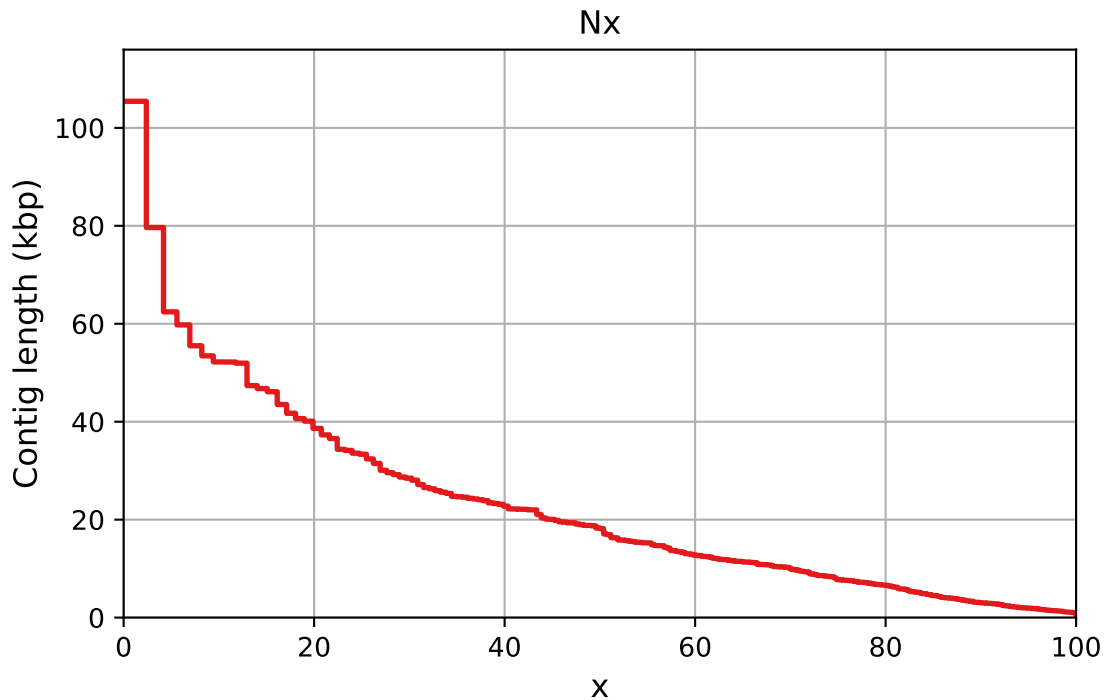


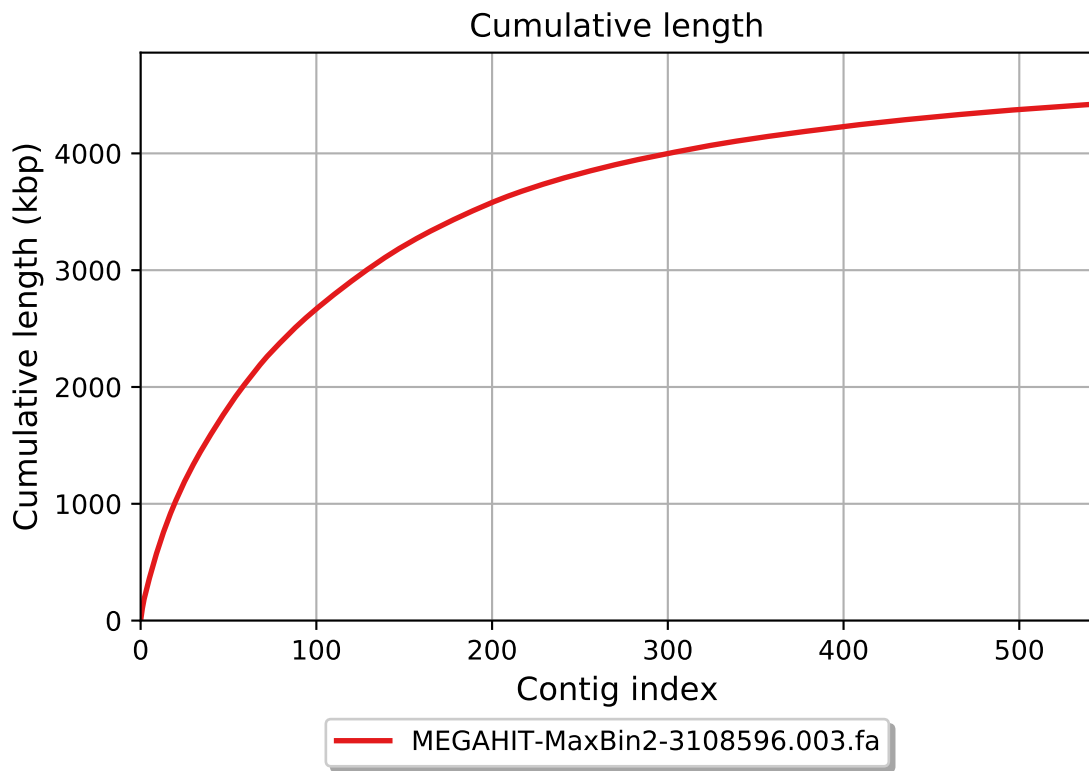
Report

	MEGAHIT-MaxBin2-3108596.003.fa
# contigs (>= 0 bp)	542
# contigs (>= 1000 bp)	542
# contigs (>= 5000 bp)	221
# contigs (>= 10000 bp)	138
# contigs (>= 25000 bp)	37
# contigs (>= 50000 bp)	9
Total length (>= 0 bp)	4420480
Total length (>= 1000 bp)	4420480
Total length (>= 5000 bp)	3697000
Total length (>= 10000 bp)	3098098
Total length (>= 25000 bp)	1521669
Total length (>= 50000 bp)	572594
# contigs	542
Largest contig	105433
Total length	4420480
GC (%)	47.70
N50	18171
N75	7736
L50	70
L75	163
# N's per 100 kbp	0.00
# predicted rRNA genes	4 + 5 part

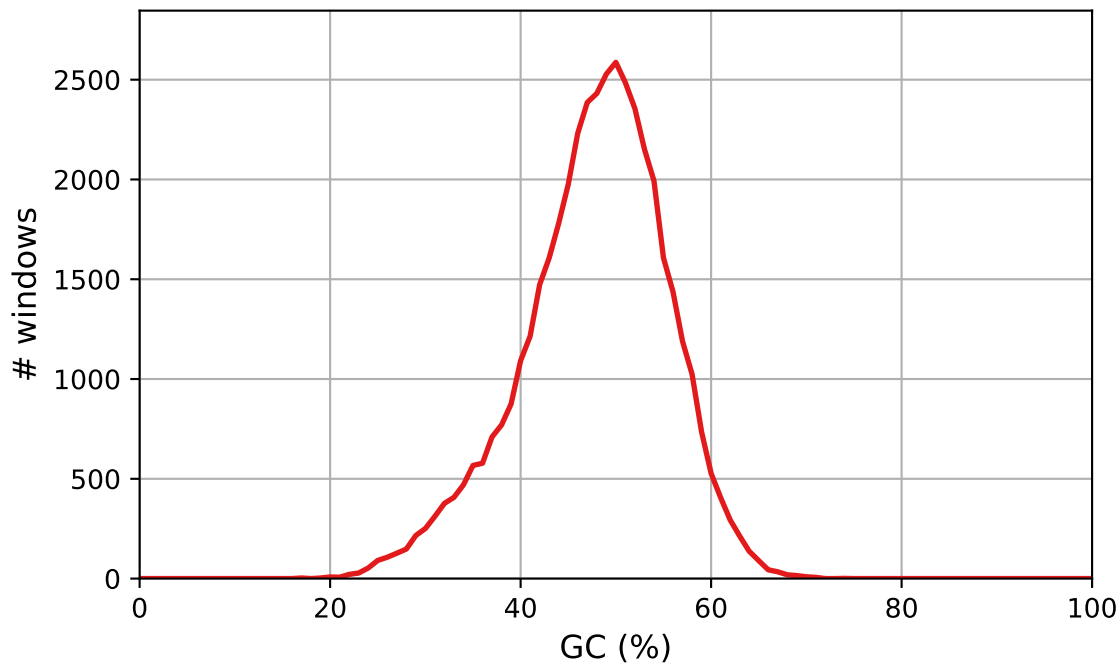
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



MEGAHIT-MaxBin2-3108596.003.fa

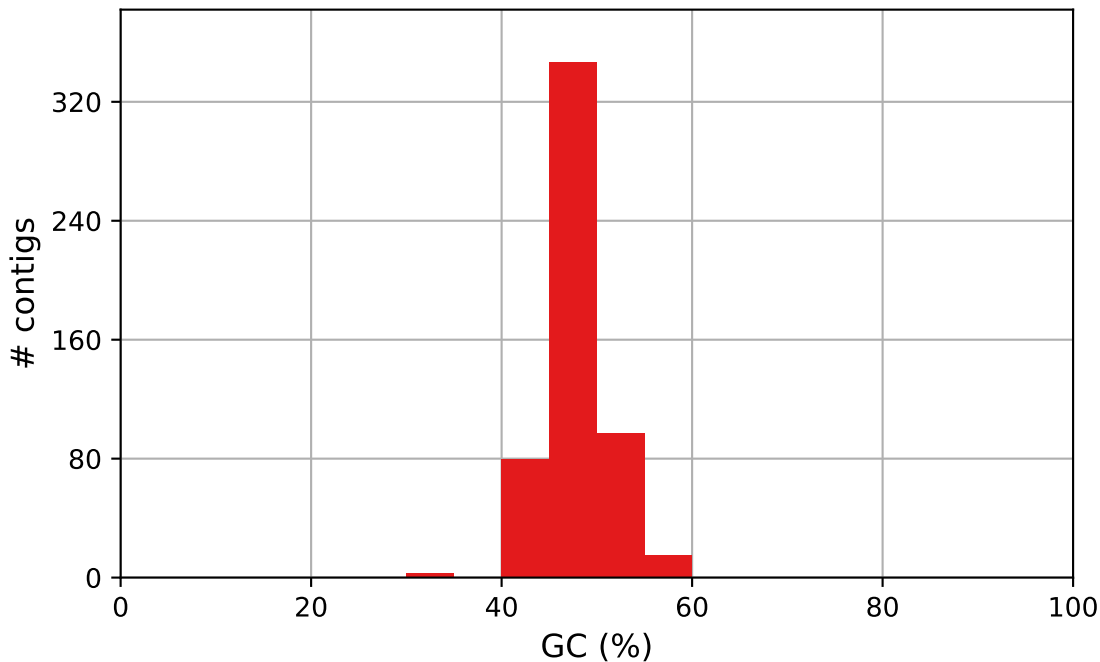


GC content



MEGAHit-MaxBin2-3108596.003.fa

MEGAHIT-MaxBin2-3108596.003.fa GC content



MEGAHIT-MaxBin2-3108596.003.fa