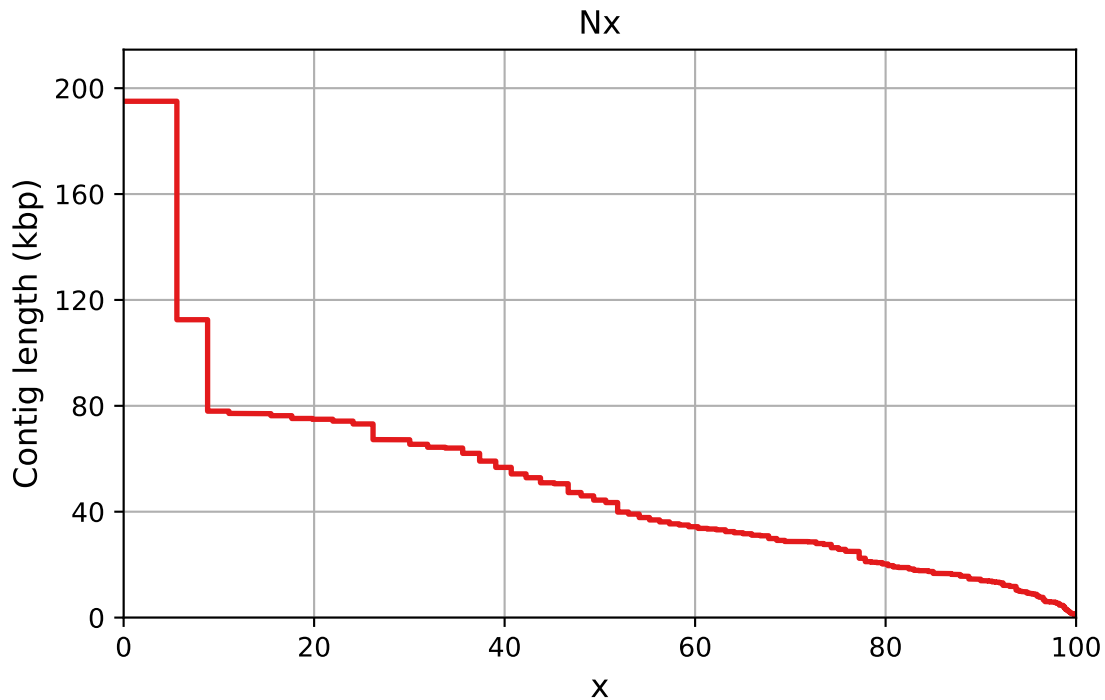


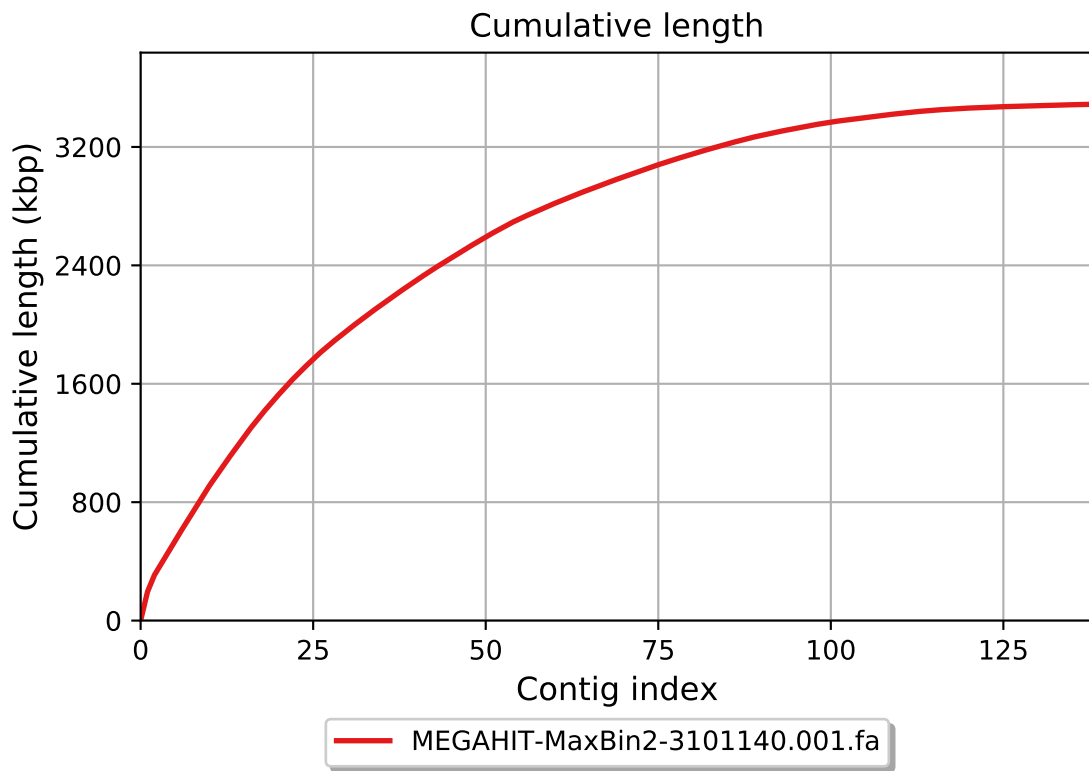
Report

	MEGAHIT-MaxBin2-3101140.001.fa
# contigs (>= 0 bp)	138
# contigs (>= 1000 bp)	138
# contigs (>= 5000 bp)	110
# contigs (>= 10000 bp)	90
# contigs (>= 25000 bp)	53
# contigs (>= 50000 bp)	22
Total length (>= 0 bp)	3488433
Total length (>= 1000 bp)	3488433
Total length (>= 5000 bp)	3426817
Total length (>= 10000 bp)	3278921
Total length (>= 25000 bp)	2668988
Total length (>= 50000 bp)	1628276
# contigs	138
Largest contig	195053
Total length	3488433
GC (%)	42.09
N50	44376
N75	26400
L50	25
L75	51
# N's per 100 kbp	0.00
# predicted rRNA genes	1 + 0 part

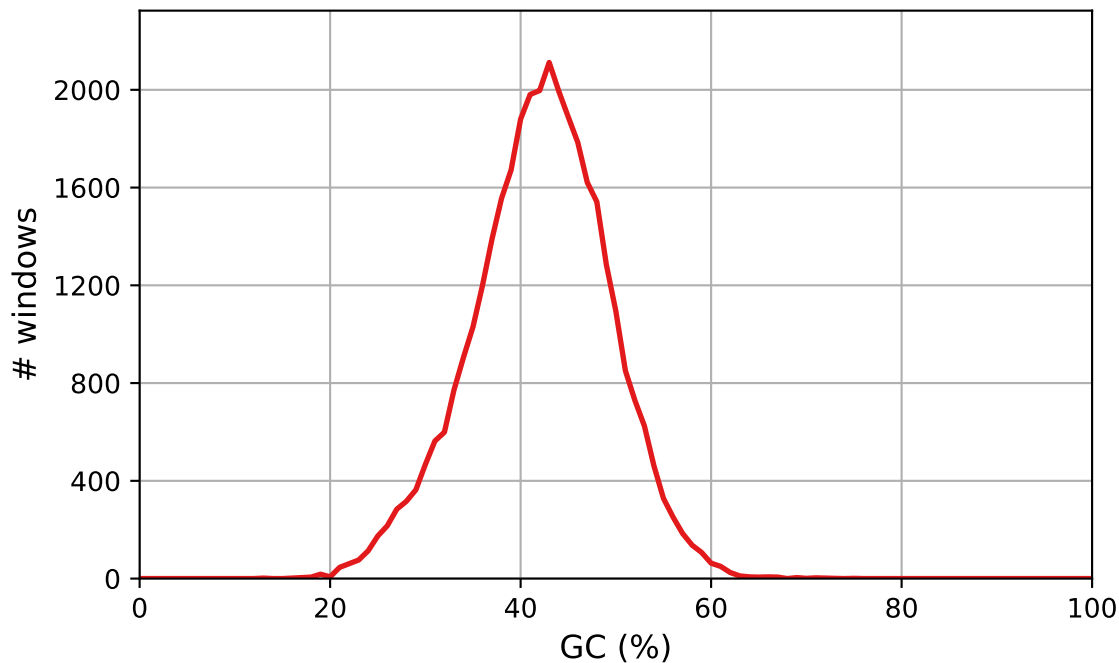
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



MEGAHIT-MaxBin2-3101140.001.fa

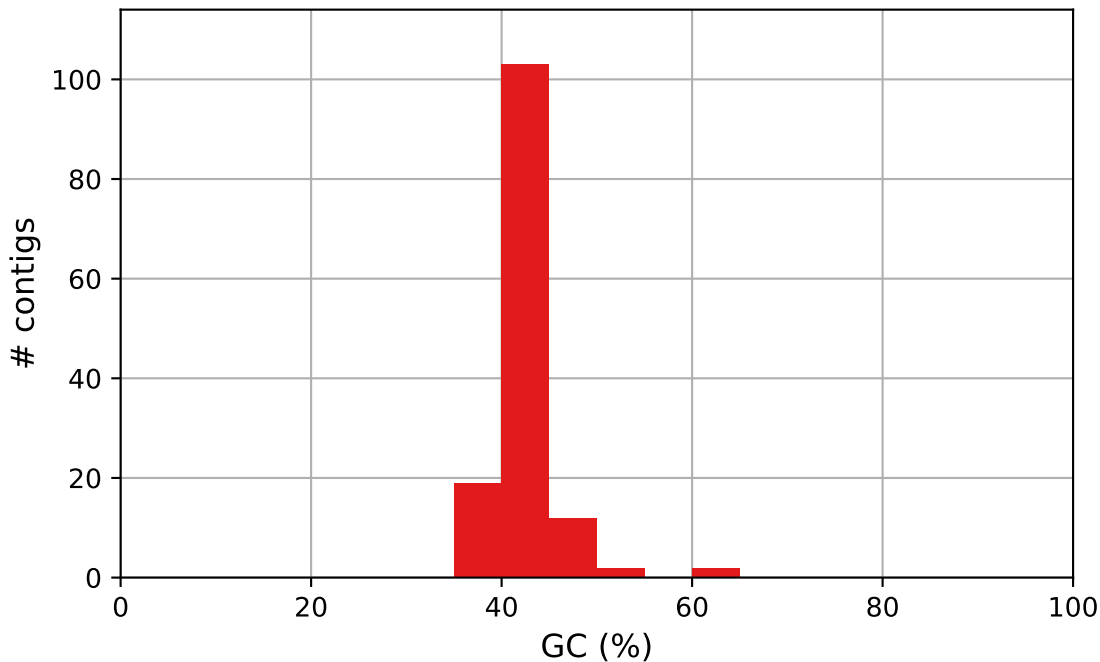


GC content



MEGAHIT-MaxBin2-3101140.001.fa

MEGAHIT-MaxBin2-3101140.001.fa GC content



MEGAHit-MaxBin2-3101140.001.fa