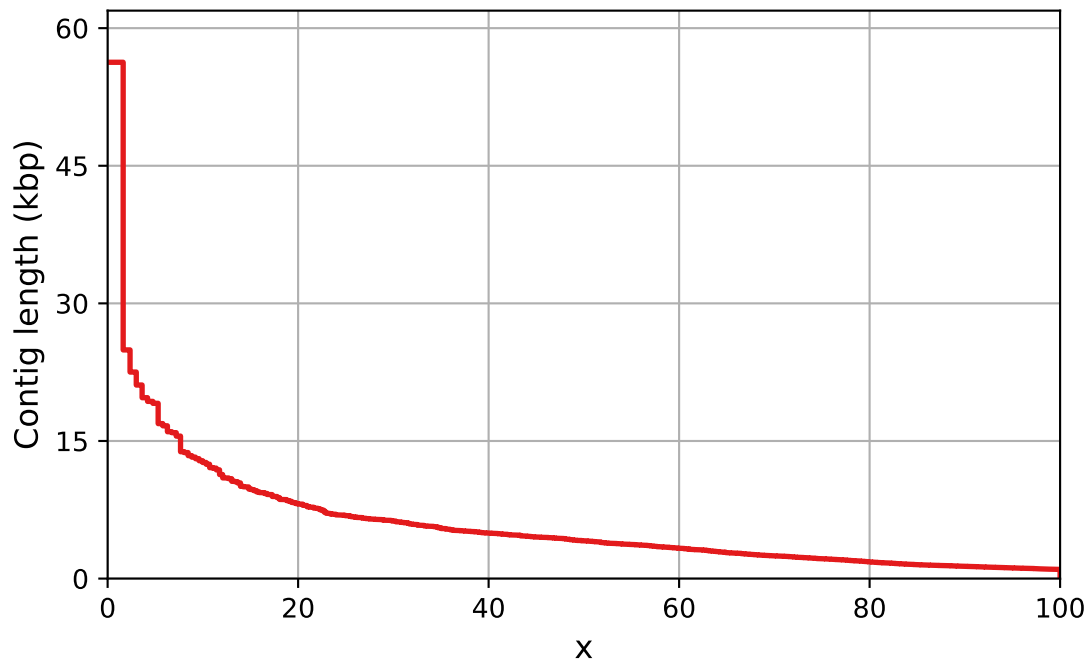


Report

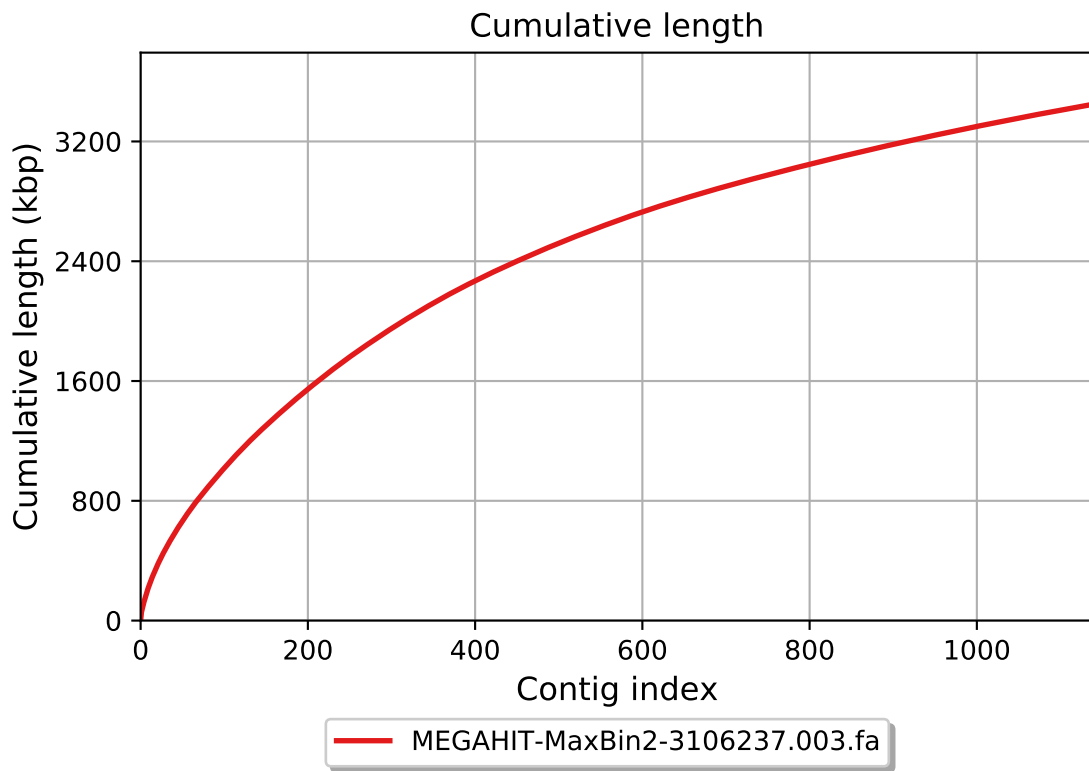
	MEGAHIT-MaxBin2-3106237.003.fa
# contigs (≥ 0 bp)	1139
# contigs (≥ 1000 bp)	1139
# contigs (≥ 5000 bp)	161
# contigs (≥ 10000 bp)	32
# contigs (≥ 25000 bp)	1
# contigs (≥ 50000 bp)	1
Total length (≥ 0 bp)	3448371
Total length (≥ 1000 bp)	3448371
Total length (≥ 5000 bp)	1359061
Total length (≥ 10000 bp)	501017
Total length (≥ 25000 bp)	56280
Total length (≥ 50000 bp)	56280
# contigs	1139
Largest contig	56280
Total length	3448371
GC (%)	42.34
N50	4120
N75	2167
L50	241
L75	530
# N's per 100 kbp	0.00
# predicted rRNA genes	1 + 1 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).

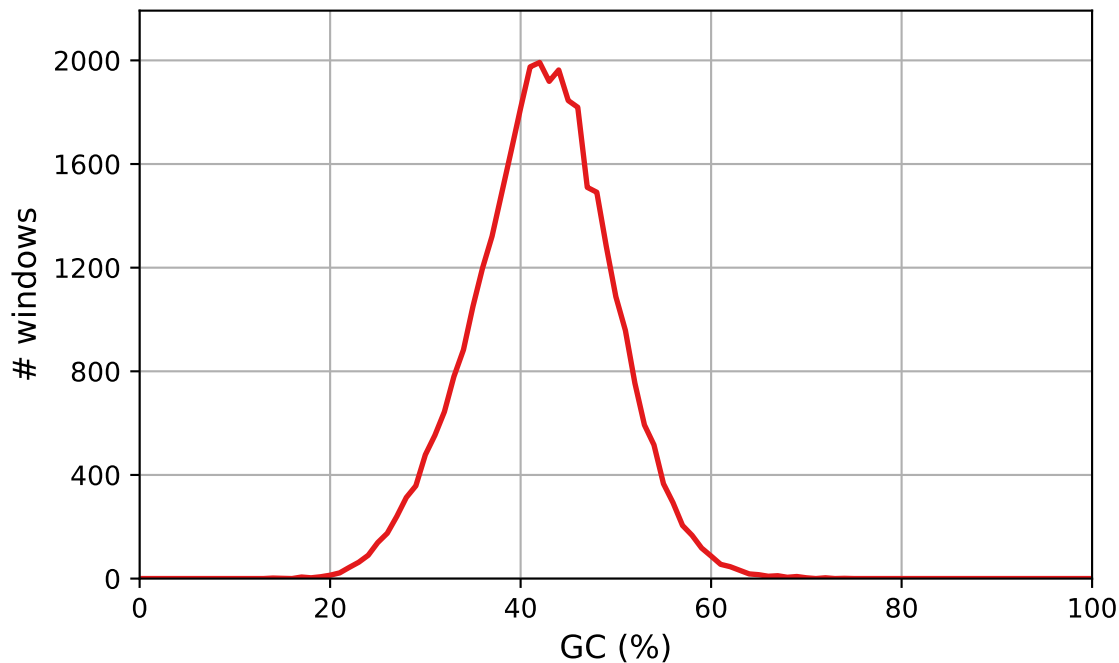
Nx



MEGAHIT-MaxBin2-3106237.003.fa

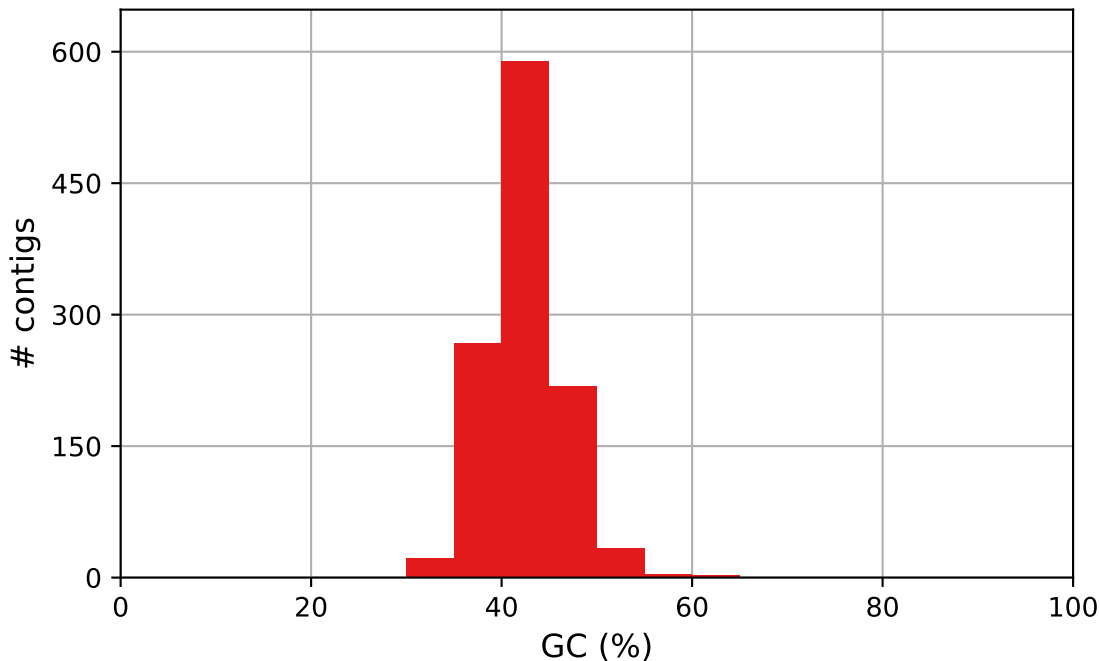


GC content



MEGAHIT-MaxBin2-3106237.003.fa

MEGAHIT-MaxBin2-3106237.003.fa GC content



MEGAHit-MaxBin2-3106237.003.fa