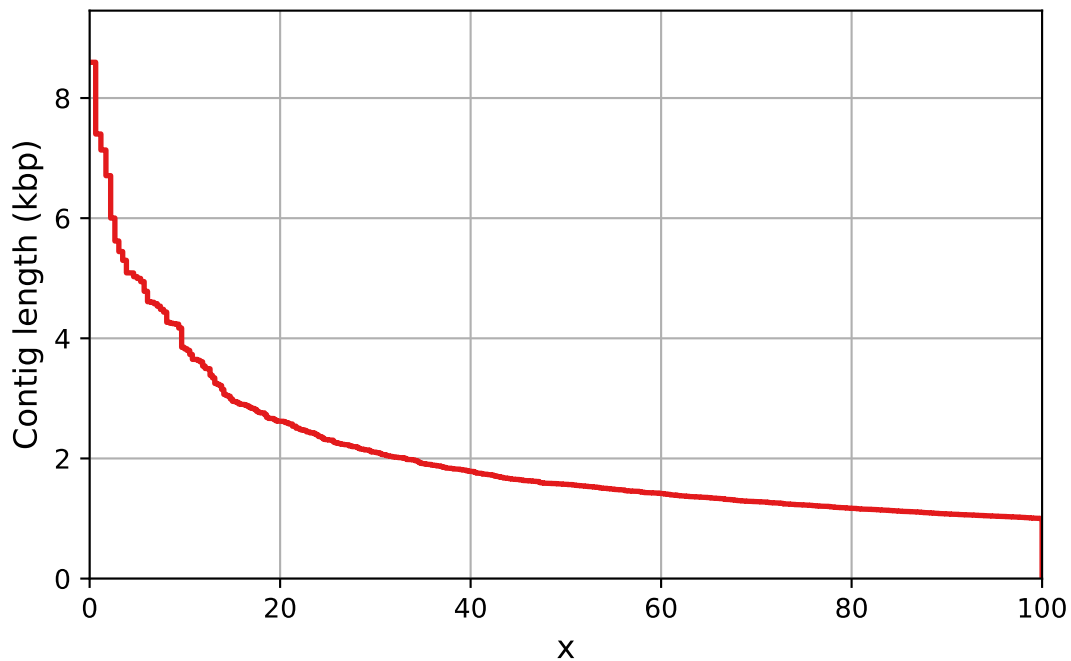


Report

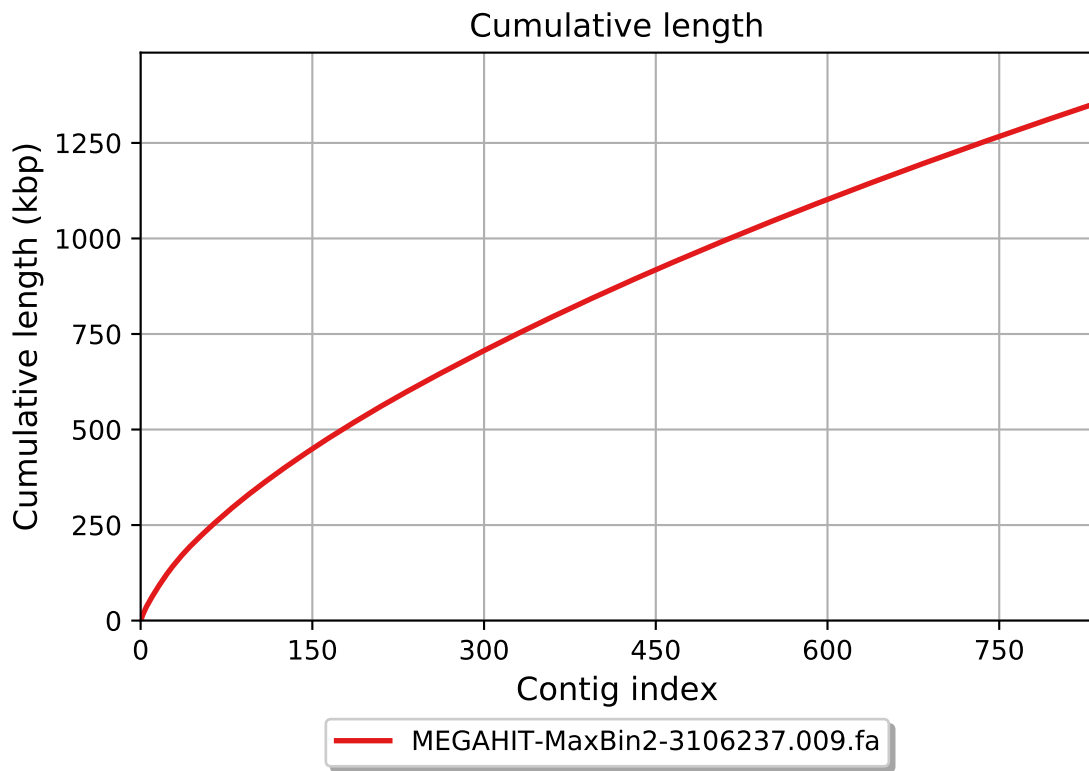
	MEGAHIT-MaxBin2-3106237.009.fa
# contigs (>= 0 bp)	832
# contigs (>= 1000 bp)	832
# contigs (>= 5000 bp)	11
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1351167
Total length (>= 1000 bp)	1351167
Total length (>= 5000 bp)	67408
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	832
Largest contig	8595
Total length	1351167
GC (%)	43.43
N50	1569
N75	1226
L50	281
L75	526
# 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).

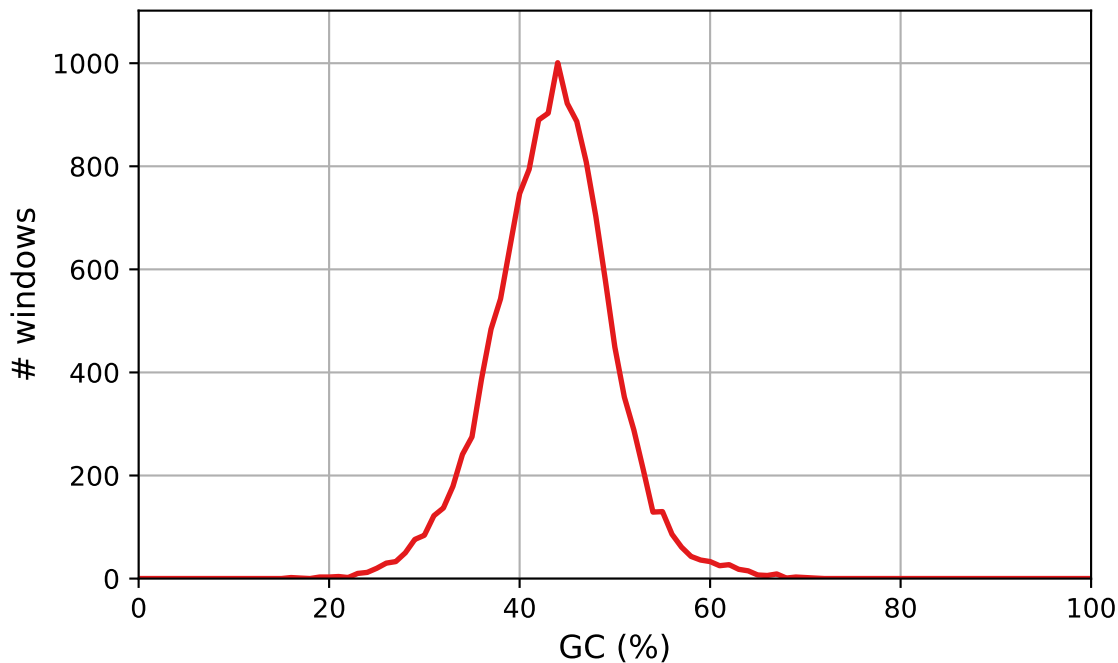
Nx



MEGAHIT-MaxBin2-3106237.009.fa

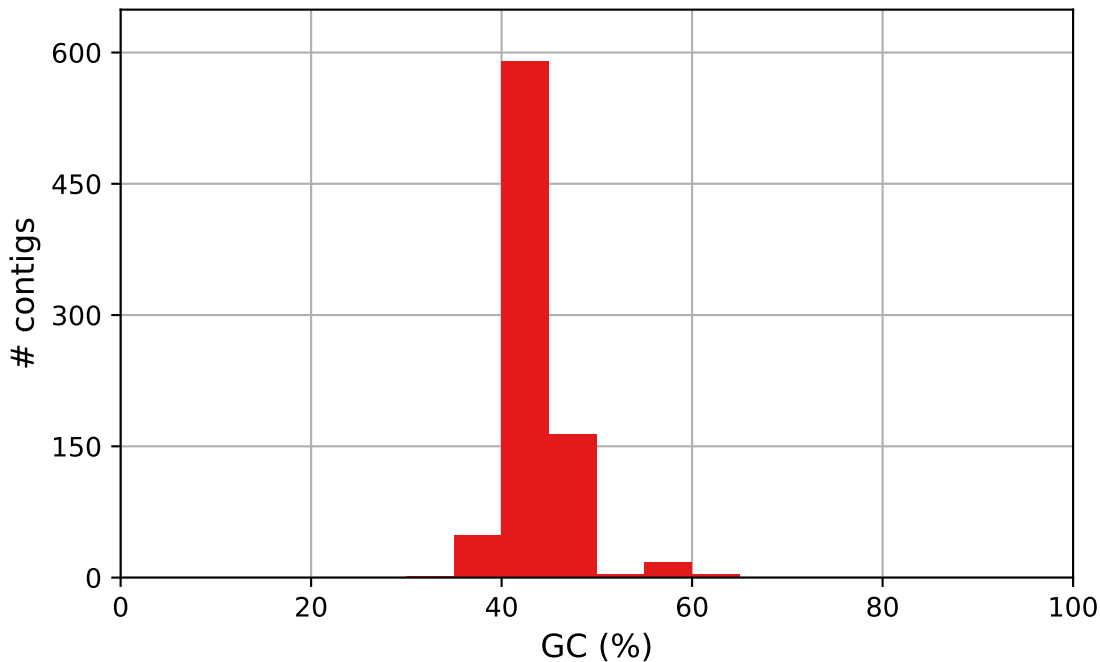


GC content



MEGAHIT-MaxBin2-3106237.009.fa

MEGAHIT-MaxBin2-3106237.009.fa GC content



MEGAHit-MaxBin2-3106237.009.fa