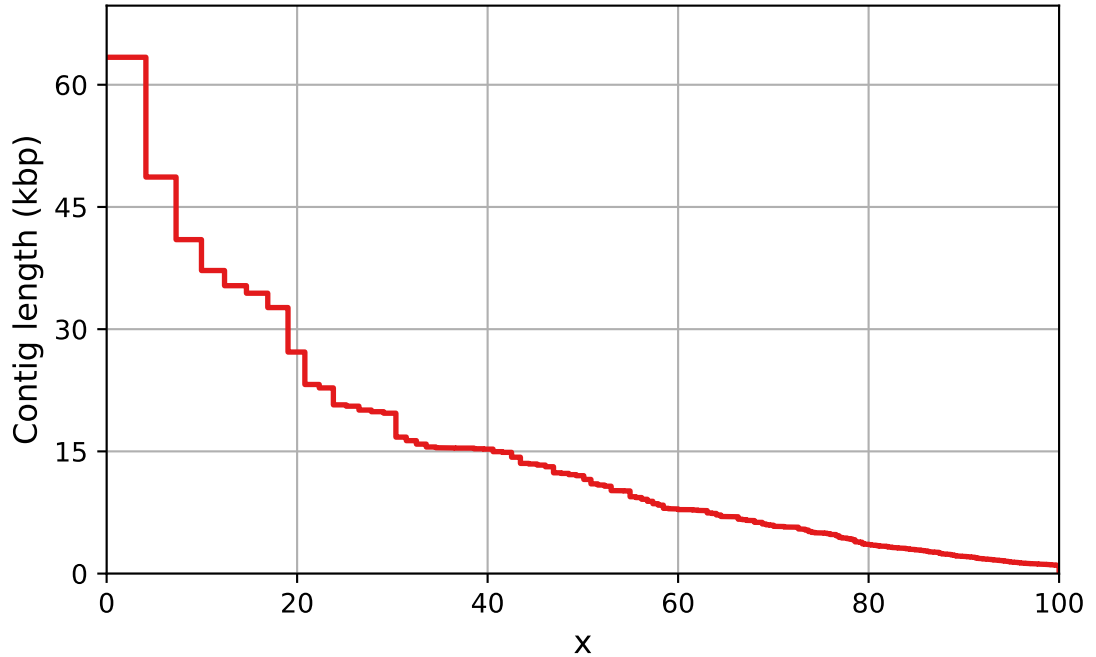


Report

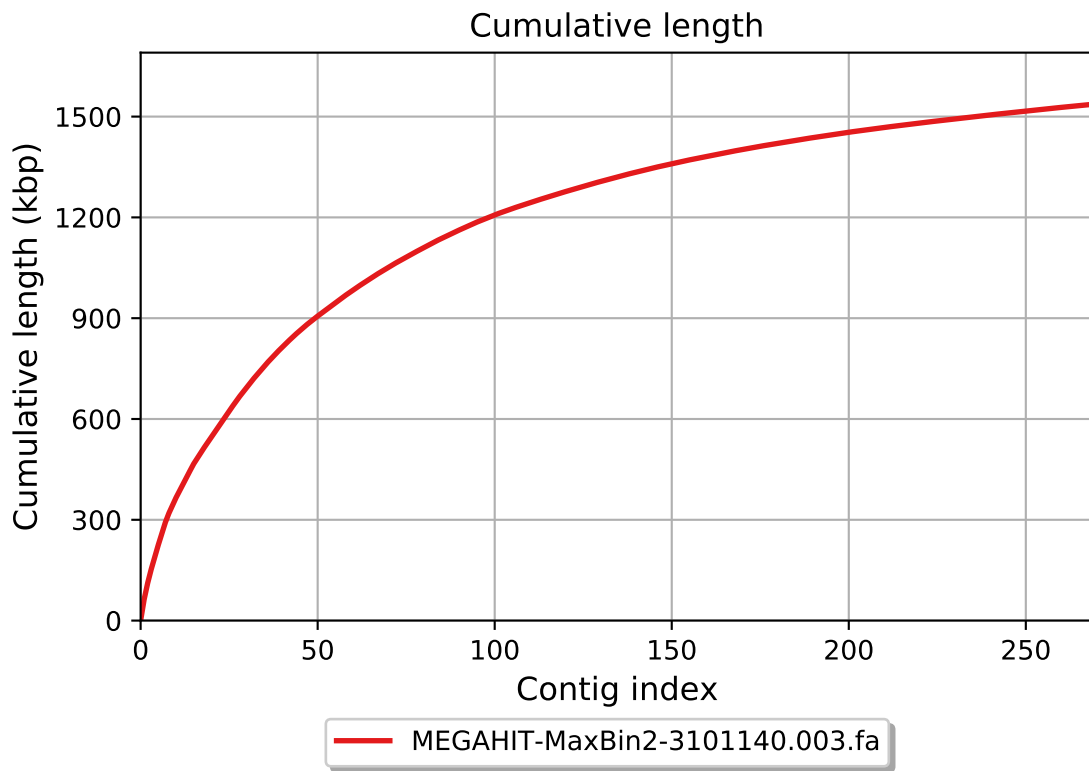
	MEGAHIT-MaxBin2-3101140.003.fa
# contigs (>= 0 bp)	269
# contigs (>= 1000 bp)	269
# contigs (>= 5000 bp)	87
# contigs (>= 10000 bp)	43
# contigs (>= 25000 bp)	8
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	1536682
Total length (>= 1000 bp)	1536682
Total length (>= 5000 bp)	1147539
Total length (>= 10000 bp)	844498
Total length (>= 25000 bp)	319828
Total length (>= 50000 bp)	63373
# contigs	269
Largest contig	63373
Total length	1536682
GC (%)	37.18
N50	12016
N75	4989
L50	36
L75	88
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 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).

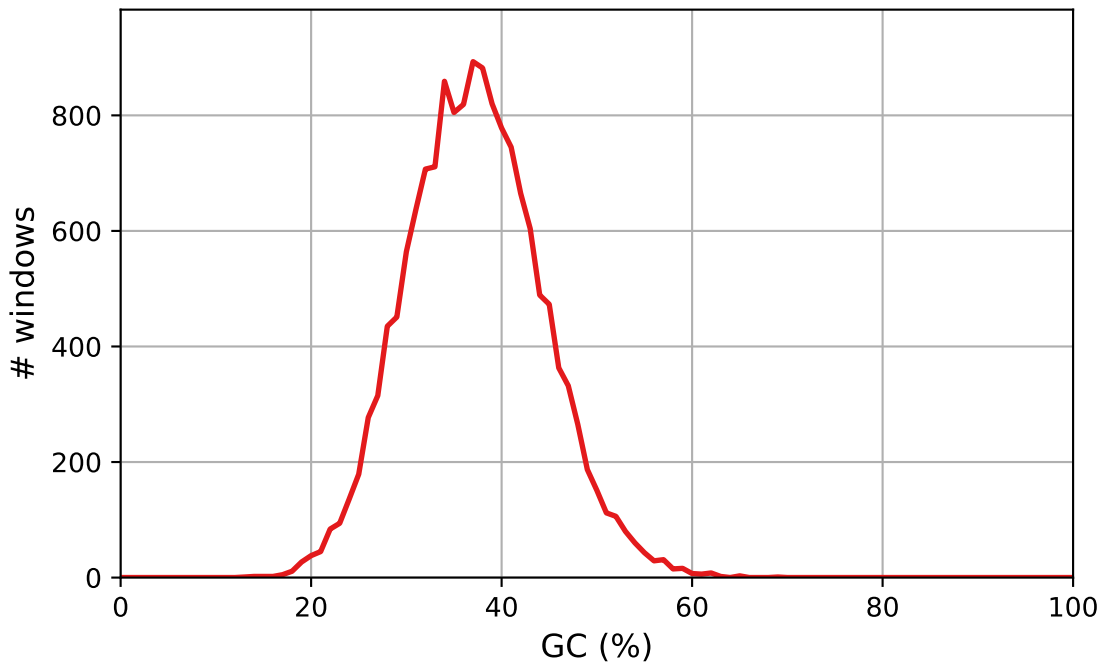
Nx



MEGAHIT-MaxBin2-3101140.003.fa

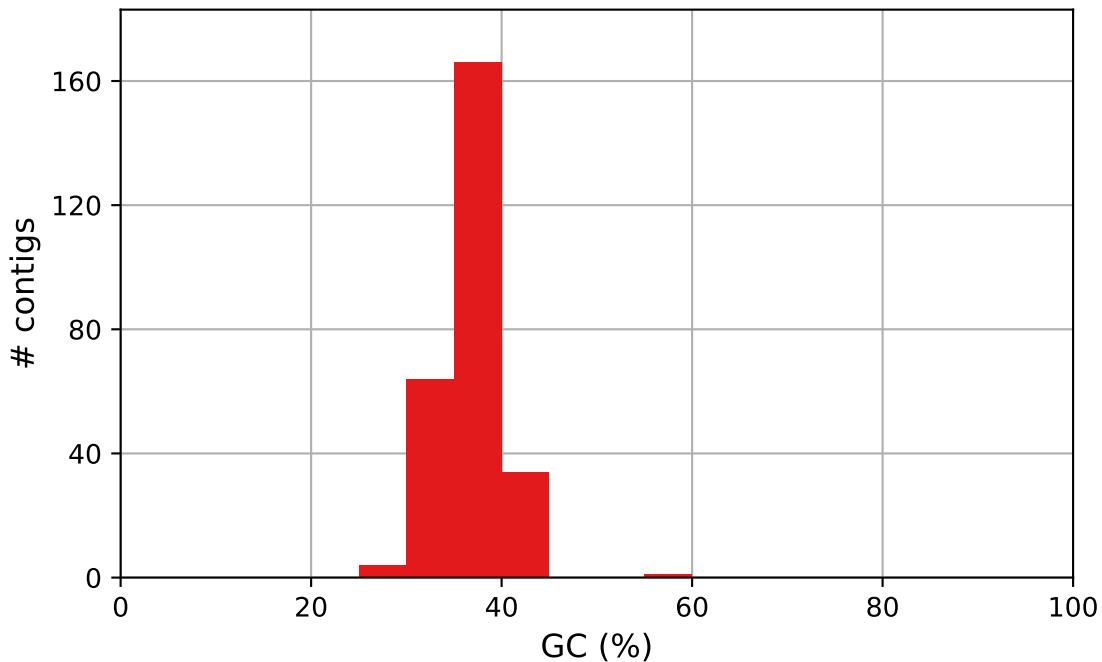


GC content



MEGAHIT-MaxBin2-3101140.003.fa

MEGAHIT-MaxBin2-3101140.003.fa GC content



MEGAHit-MaxBin2-3101140.003.fa