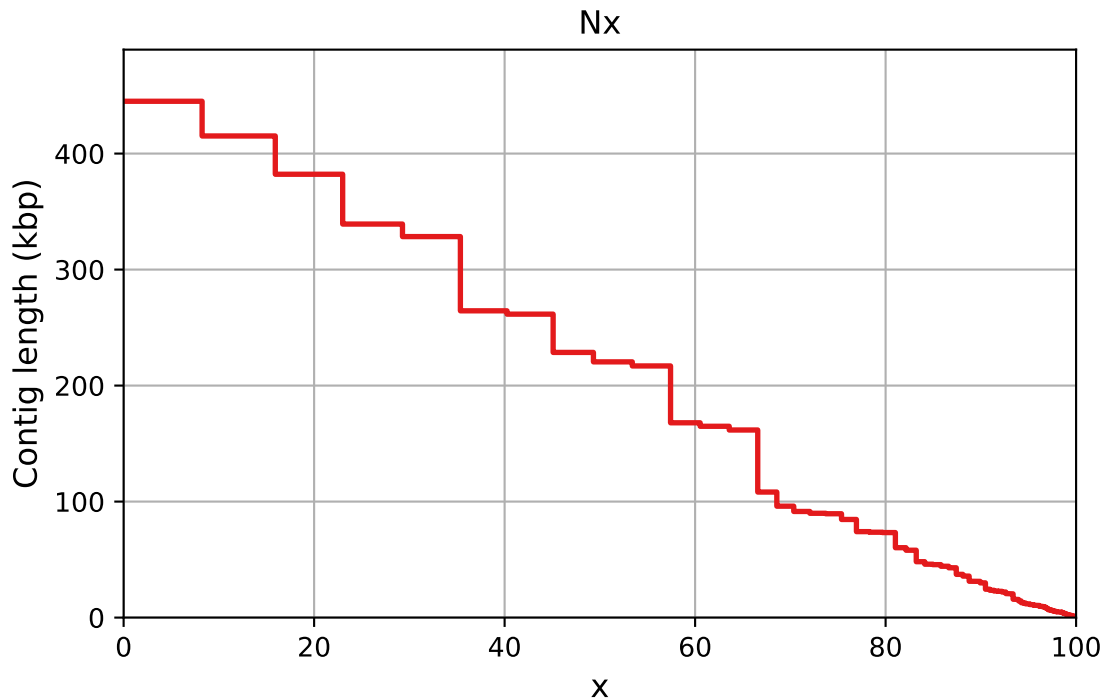


Report

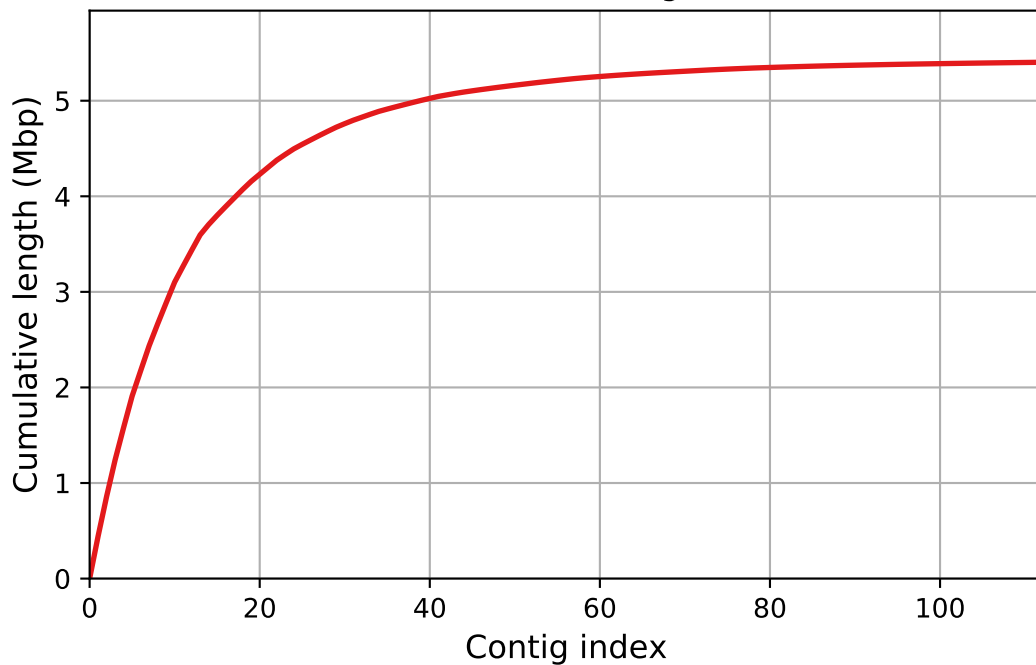
	MEGAHIT-MaxBin2-3101140.002.fa
# contigs (>= 0 bp)	112
# contigs (>= 1000 bp)	112
# contigs (>= 5000 bp)	67
# contigs (>= 10000 bp)	53
# contigs (>= 25000 bp)	34
# contigs (>= 50000 bp)	24
Total length (>= 0 bp)	5402312
Total length (>= 1000 bp)	5402312
Total length (>= 5000 bp)	5293945
Total length (>= 10000 bp)	5193696
Total length (>= 25000 bp)	4888201
Total length (>= 50000 bp)	4495668
# contigs	112
Largest contig	445087
Total length	5402312
GC (%)	46.12
N50	220451
N75	89517
L50	9
L75	18
# N's per 100 kbp	0.00
# predicted rRNA genes	3 + 2 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).



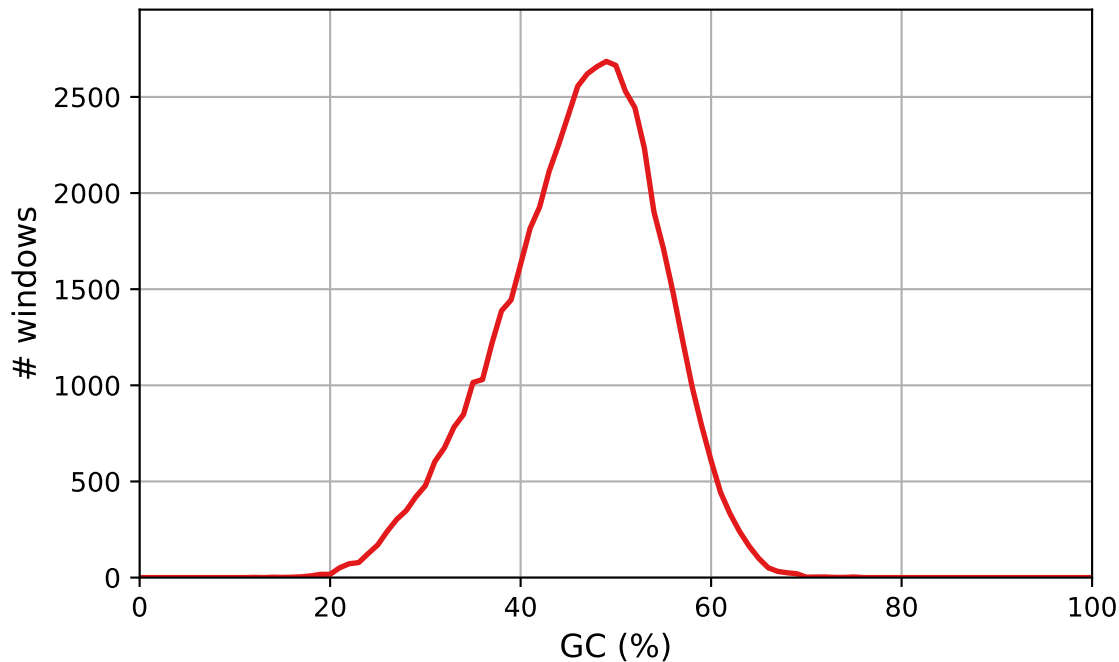
MEGAHIT-MaxBin2-3101140.002.fa

Cumulative length



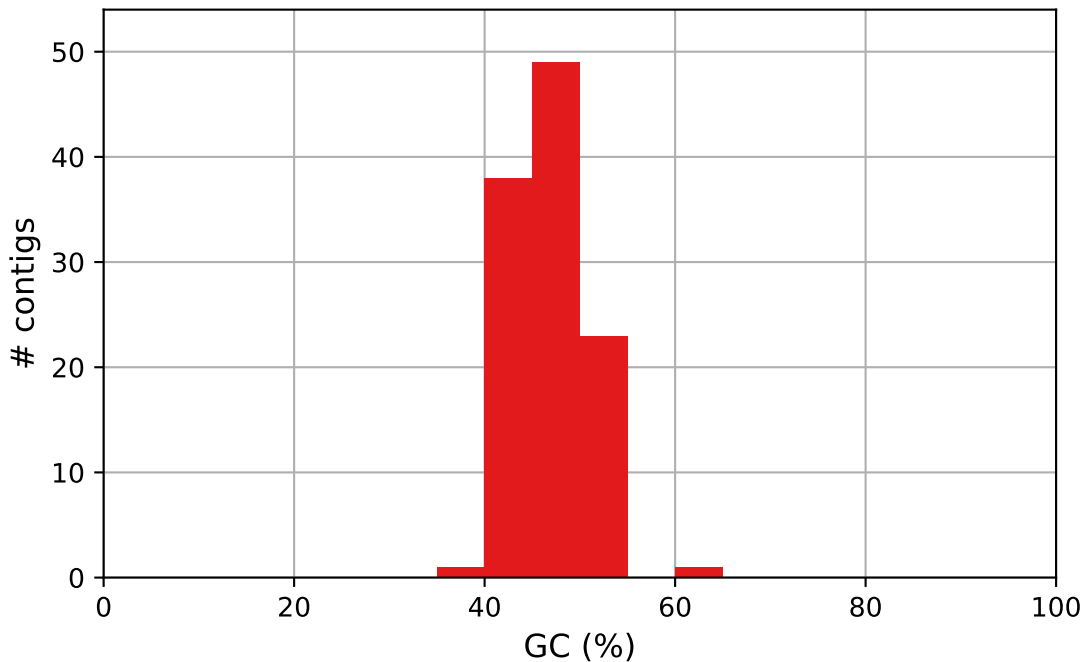
MEGAHIT-MaxBin2-3101140.002.fa

GC content



MEGAHIT-MaxBin2-3101140.002.fa

MEGAHIT-MaxBin2-3101140.002.fa GC content



MEGAHit-MaxBin2-3101140.002.fa