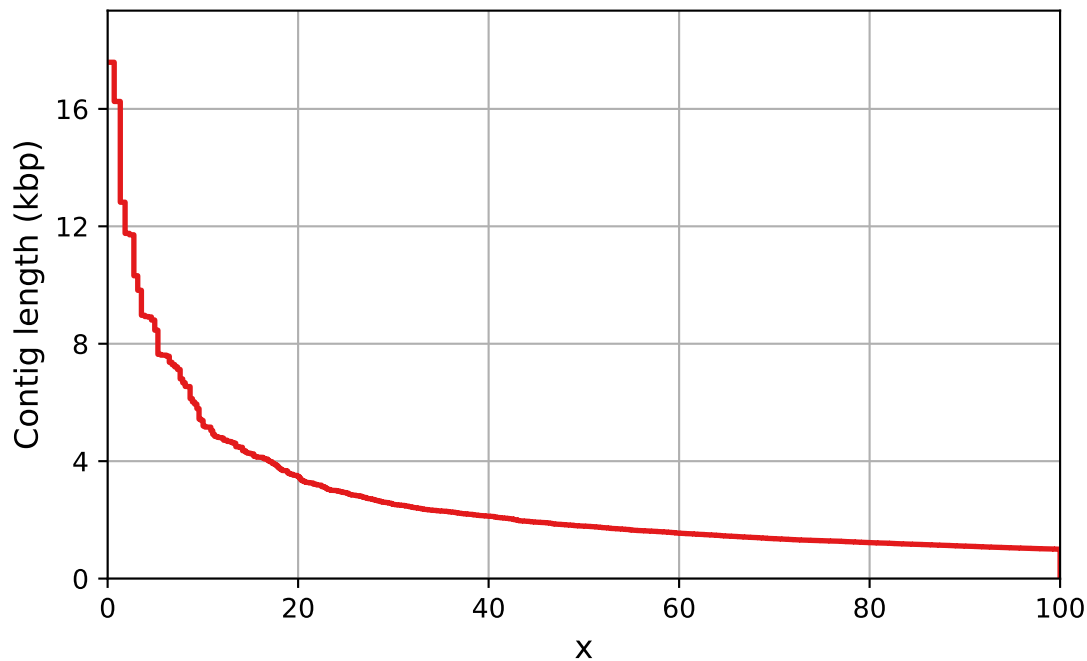


Report

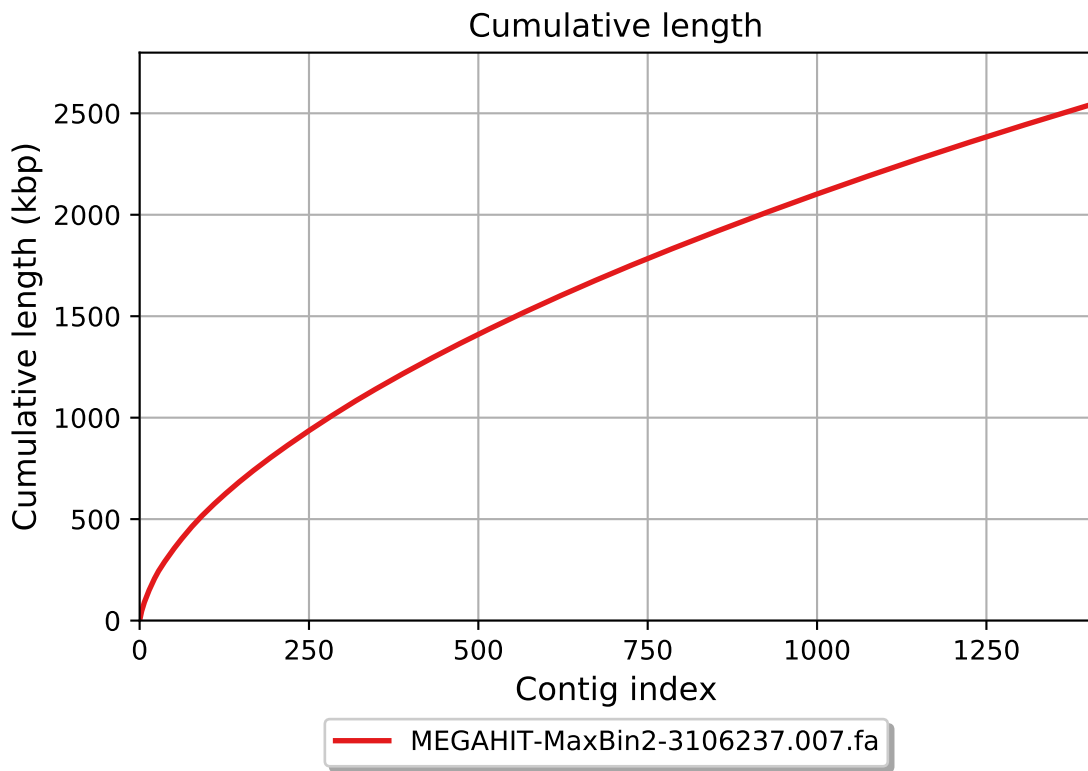
	MEGAHIT-MaxBin2-3106237.007.fa
# contigs (>= 0 bp)	1406
# contigs (>= 1000 bp)	1406
# contigs (>= 5000 bp)	35
# contigs (>= 10000 bp)	6
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2544490
Total length (>= 1000 bp)	2544490
Total length (>= 5000 bp)	280753
Total length (>= 10000 bp)	80448
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	1406
Largest contig	17588
Total length	2544490
GC (%)	37.48
N50	1786
N75	1292
L50	420
L75	845
# 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).

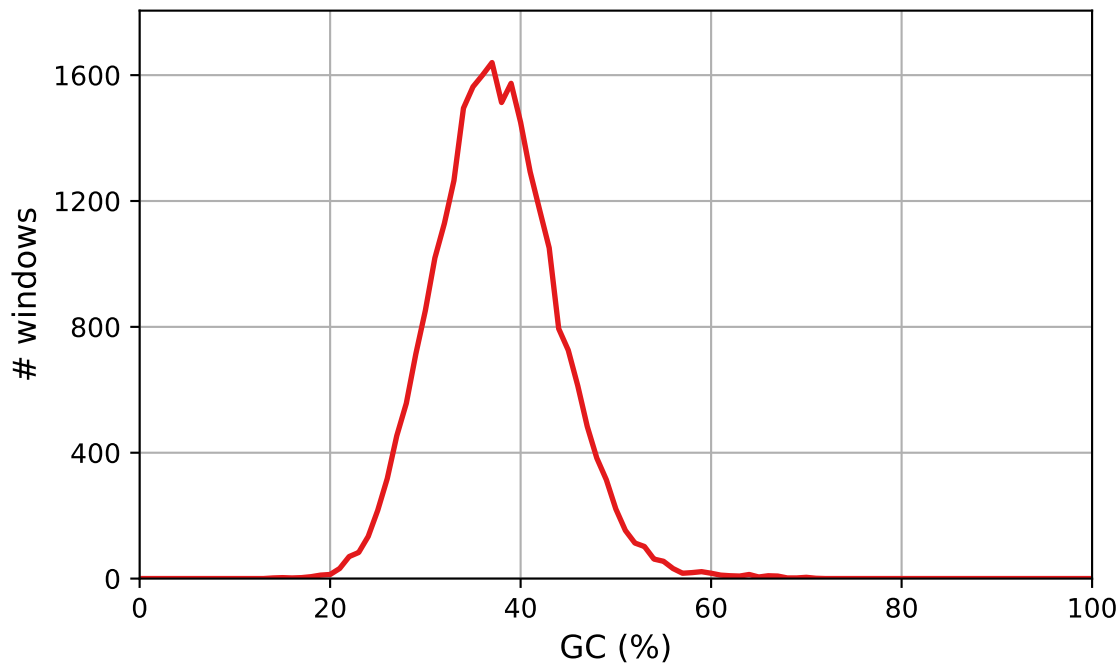
Nx



MEGAHIT-MaxBin2-3106237.007.fa

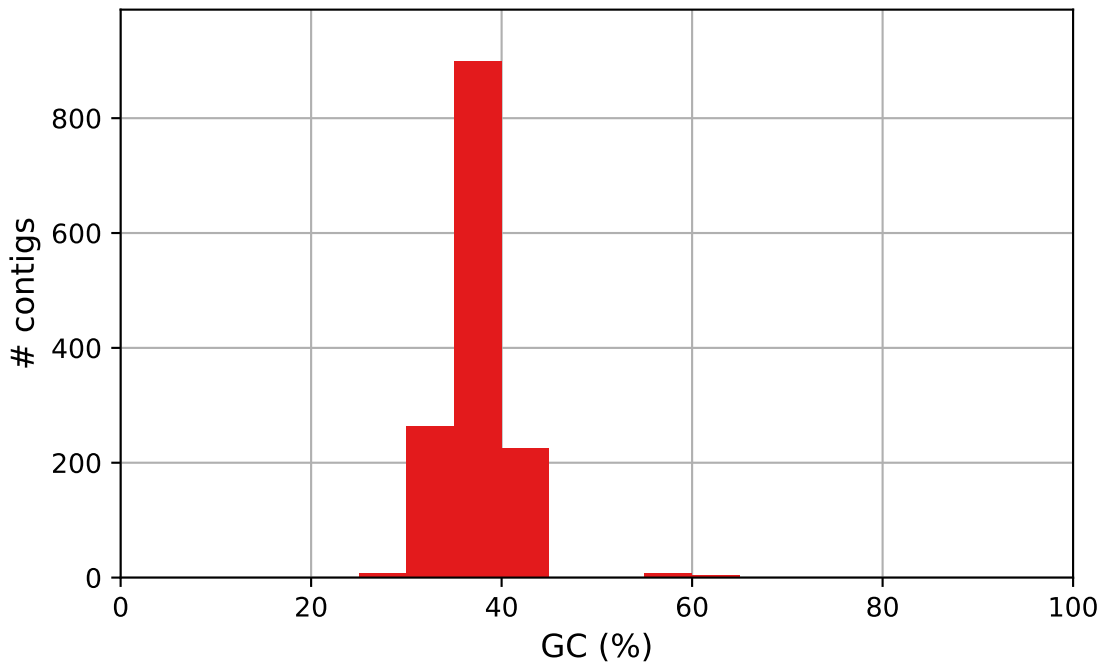


GC content



MEGAHIT-MaxBin2-3106237.007.fa

MEGAHIT-MaxBin2-3106237.007.fa GC content



MEGAHIT-MaxBin2-3106237.007.fa