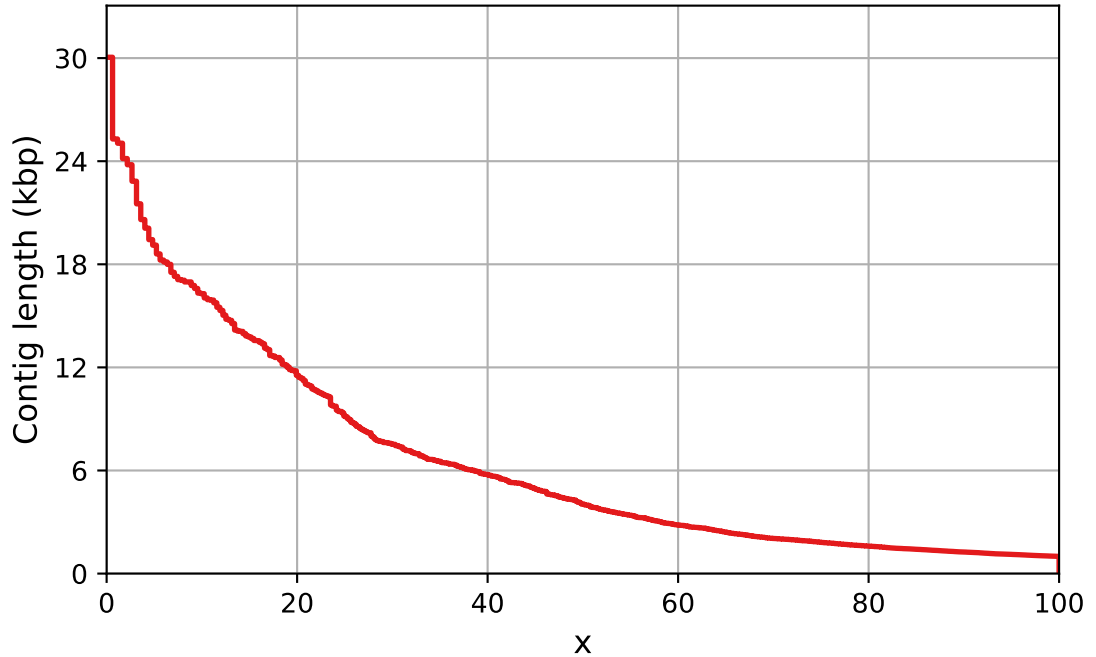


Report

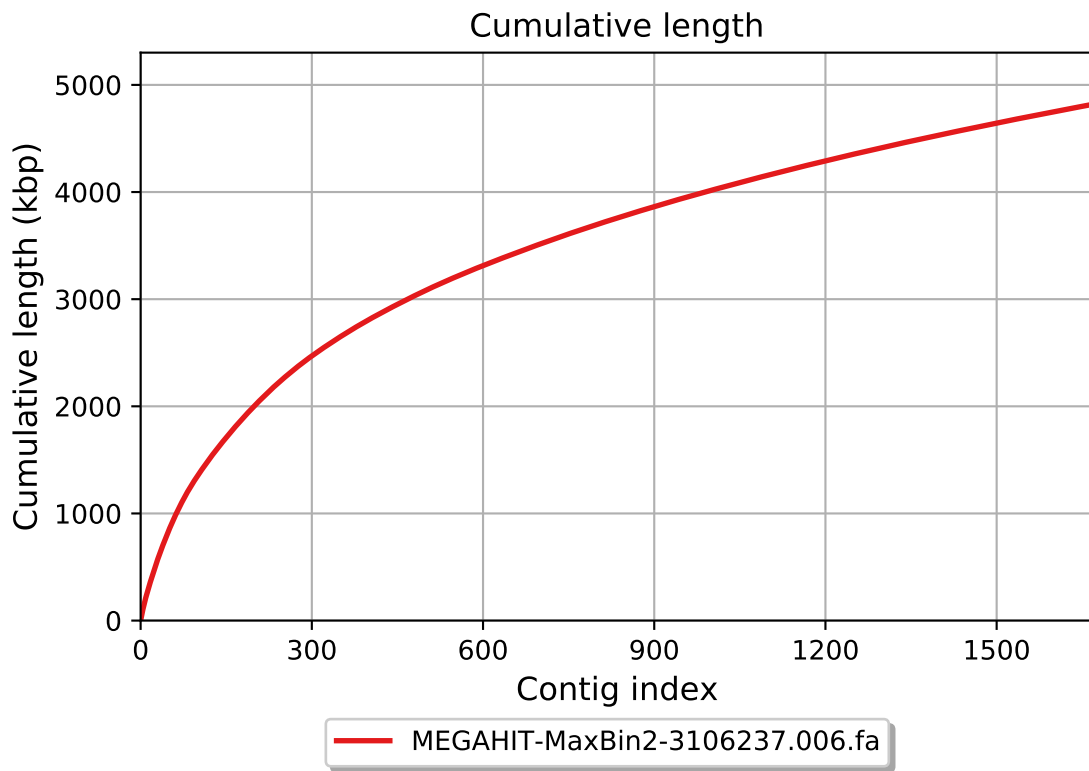
	MEGAHIT-MaxBin2-3106237.006.fa
# contigs (>= 0 bp)	1669
# contigs (>= 1000 bp)	1669
# contigs (>= 5000 bp)	229
# contigs (>= 10000 bp)	75
# contigs (>= 25000 bp)	3
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	4819227
Total length (>= 1000 bp)	4819227
Total length (>= 5000 bp)	2159383
Total length (>= 10000 bp)	1132836
Total length (>= 25000 bp)	80387
Total length (>= 50000 bp)	0
# contigs	1669
Largest contig	30048
Total length	4819227
GC (%)	47.90
N50	4039
N75	1814
L50	285
L75	754
# N's per 100 kbp	0.00
# predicted rRNA genes	6 + 6 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).

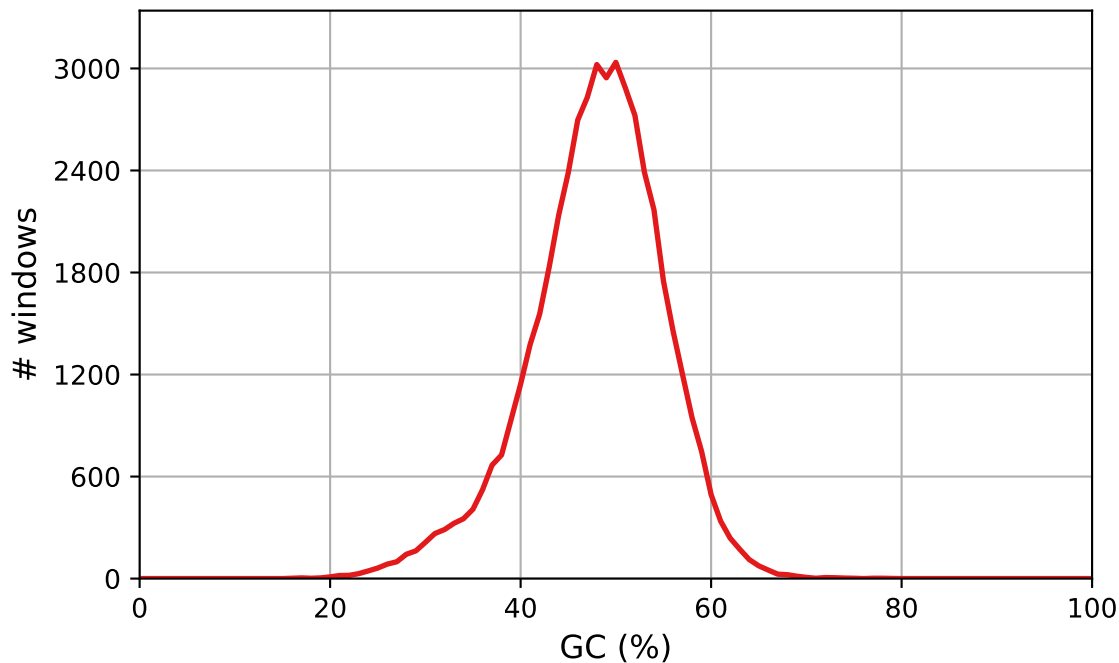
Nx



MEGAHIT-MaxBin2-3106237.006.fa

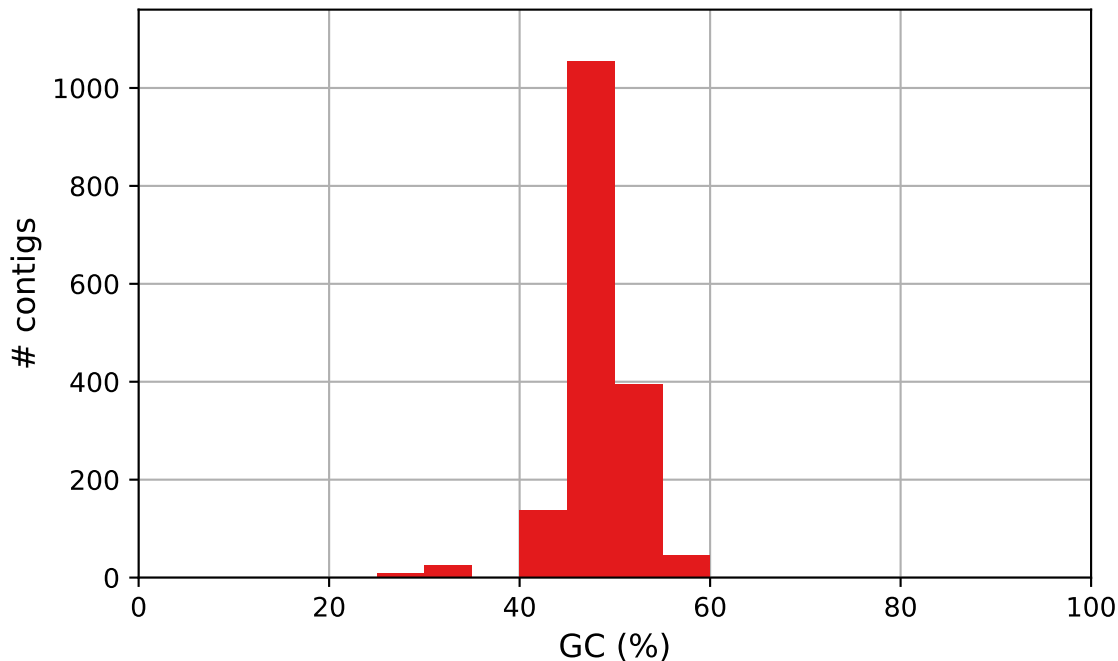


GC content



MEGAHit-MaxBin2-3106237.006.fa

MEGAHIT-MaxBin2-3106237.006.fa GC content



MEGAHit-MaxBin2-3106237.006.fa