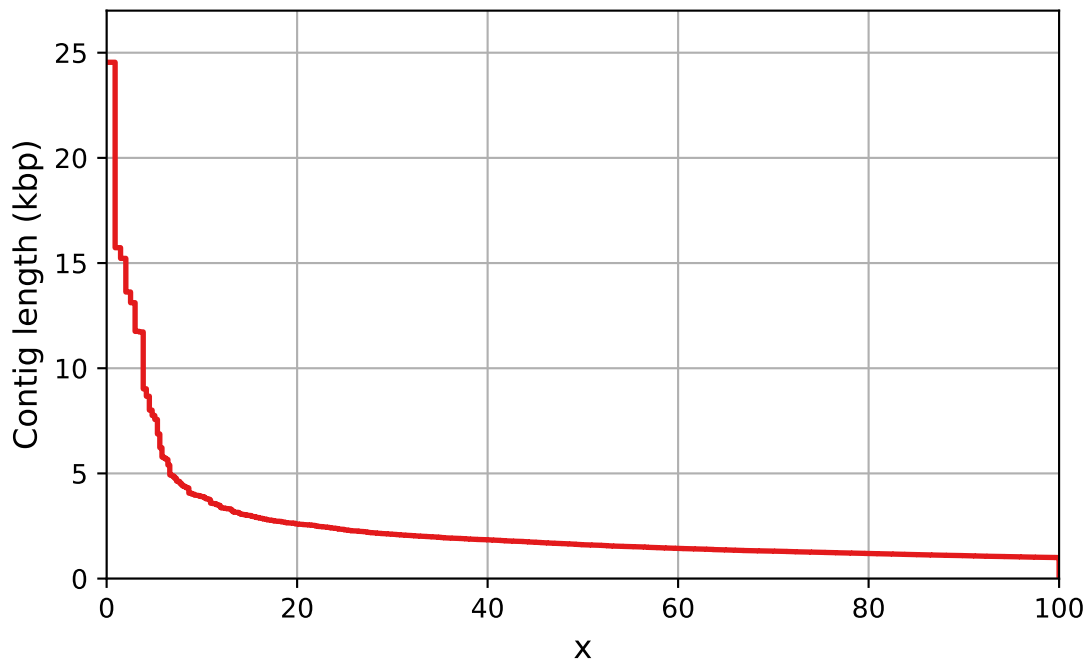


Report

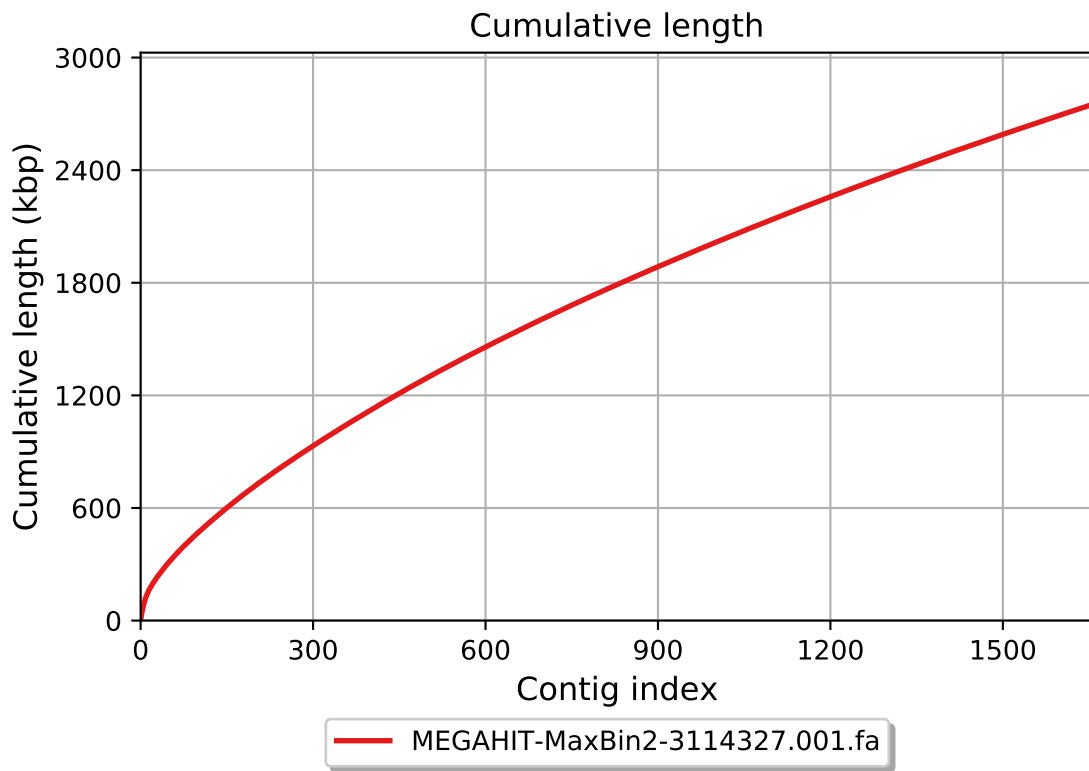
	MEGAHIT-MaxBin2-3114327.001.fa
# contigs (>= 0 bp)	1657
# contigs (>= 1000 bp)	1657
# contigs (>= 5000 bp)	18
# contigs (>= 10000 bp)	7
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2751112
Total length (>= 1000 bp)	2751112
Total length (>= 5000 bp)	182397
Total length (>= 10000 bp)	105716
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	1657
Largest contig	24545
Total length	2751112
GC (%)	42.43
N50	1609
N75	1249
L50	549
L75	1039
# 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).

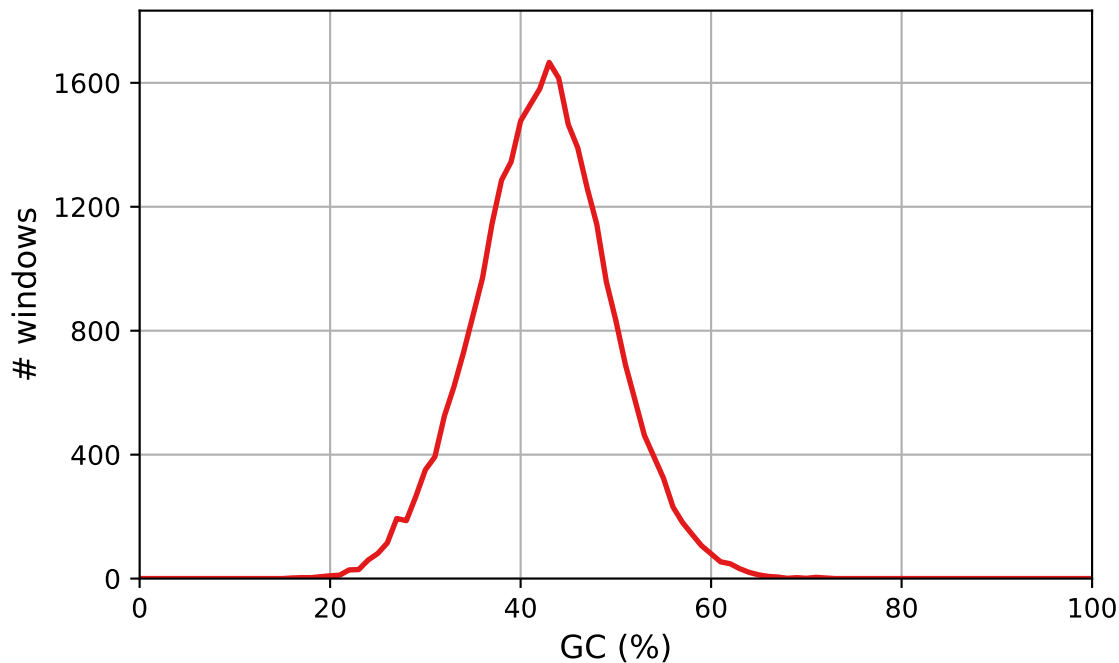
Nx



MEGAHIT-MaxBin2-3114327.001.fa

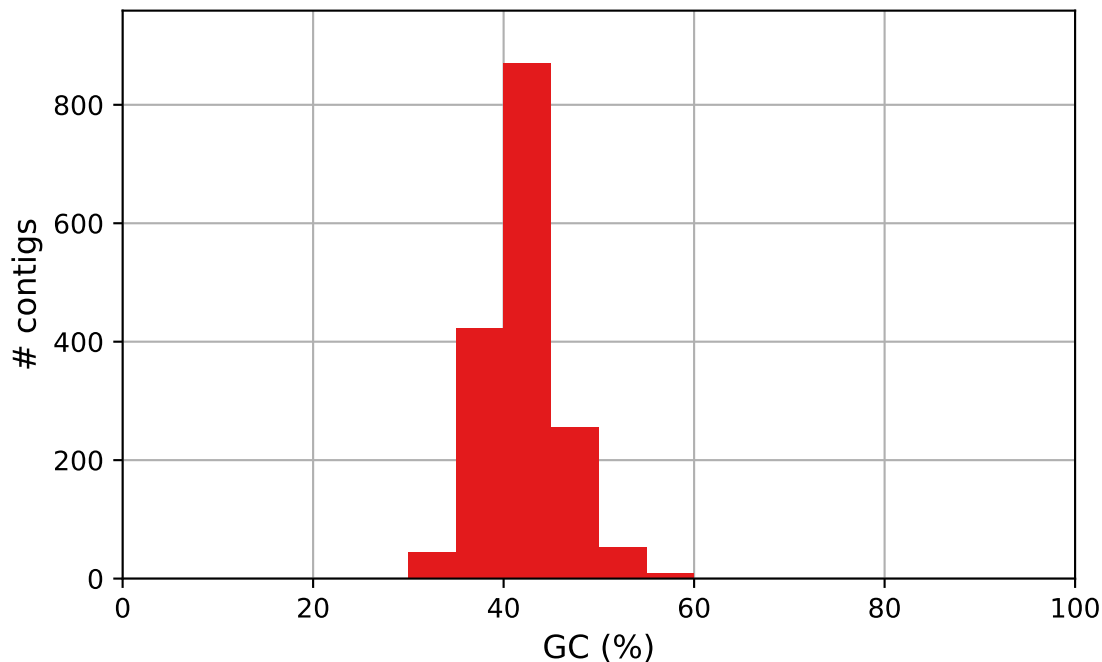


GC content



MEGAHIT-MaxBin2-3114327.001.fa

MEGAHIT-MaxBin2-3114327.001.fa GC content



MEGAHit-MaxBin2-3114327.001.fa