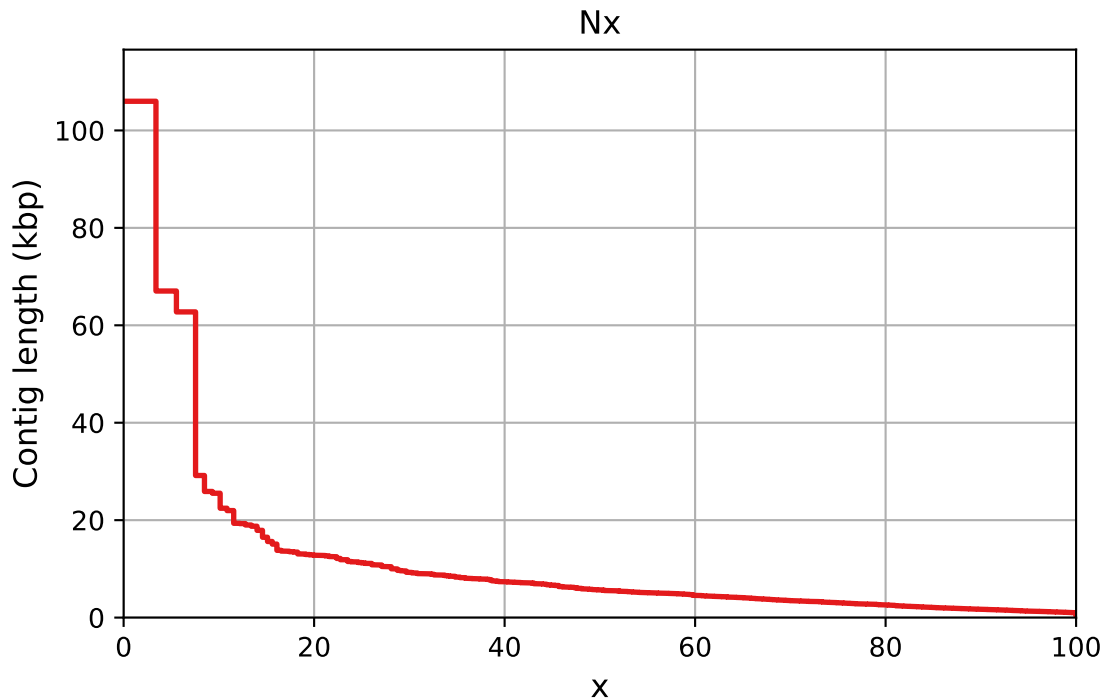


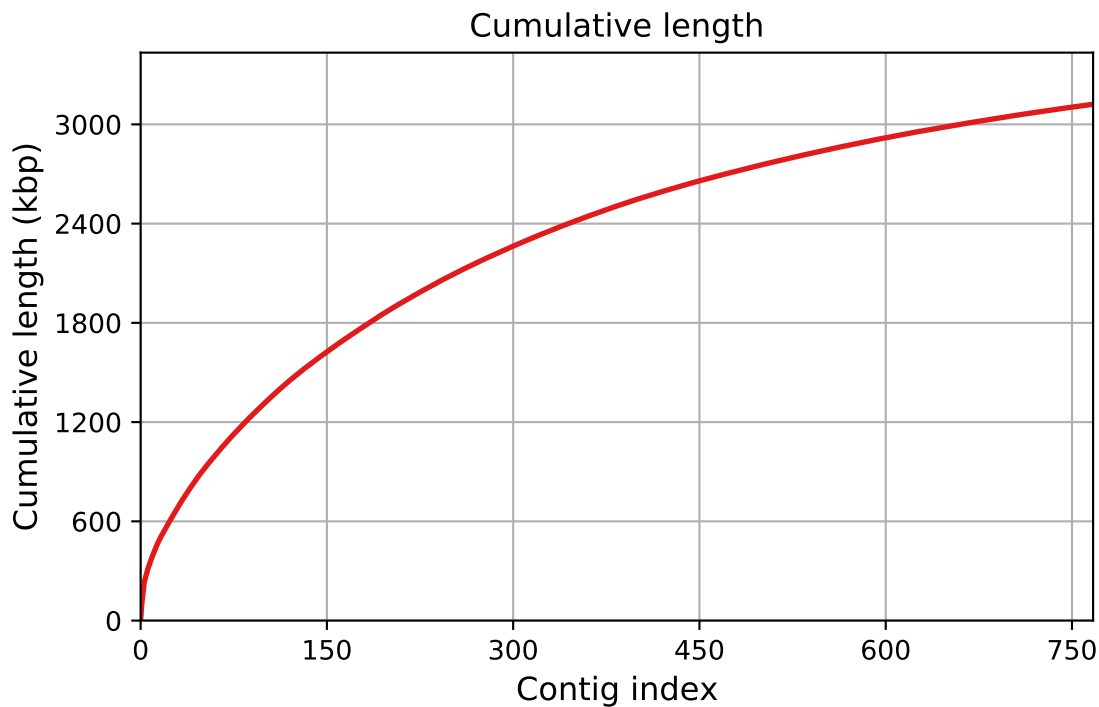
Report

	MEGAHIT-MaxBin2-3108596.004.fa
# contigs (>= 0 bp)	767
# contigs (>= 1000 bp)	767
# contigs (>= 5000 bp)	180
# contigs (>= 10000 bp)	48
# contigs (>= 25000 bp)	6
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	3121531
Total length (>= 1000 bp)	3121531
Total length (>= 5000 bp)	1780272
Total length (>= 10000 bp)	887206
Total length (>= 25000 bp)	316352
Total length (>= 50000 bp)	235771
# contigs	767
Largest contig	105987
Total length	3121531
GC (%)	40.52
N50	5693
N75	3045
L50	139
L75	325
# N's per 100 kbp	0.00
# predicted rRNA genes	1 + 0 part

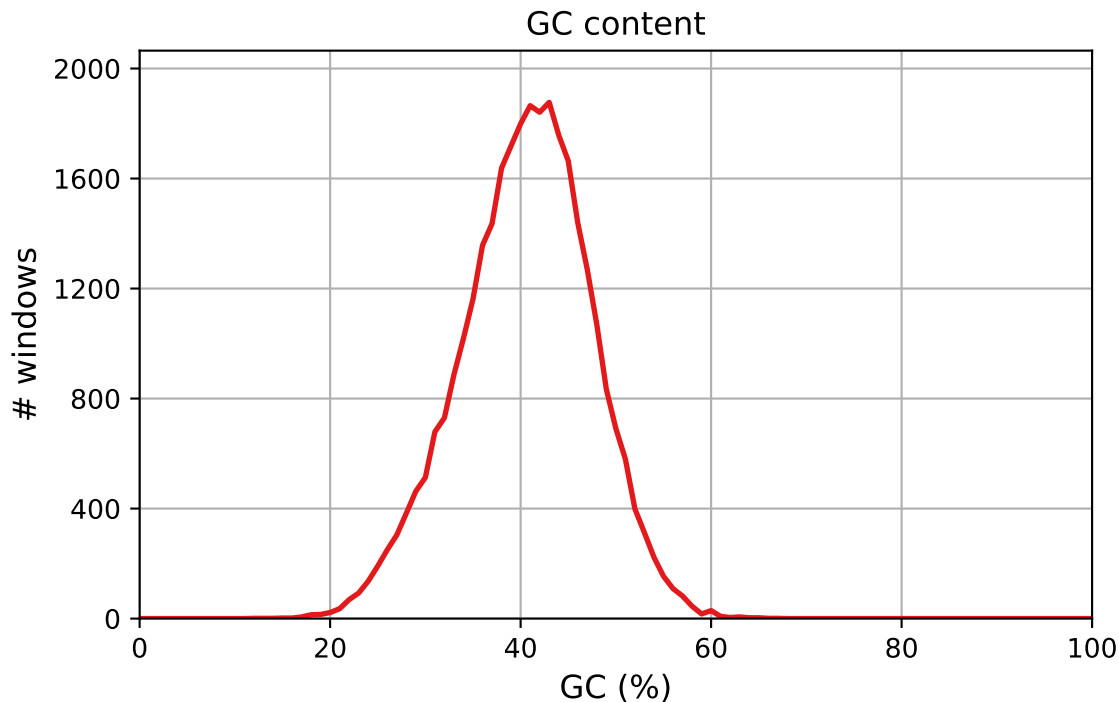
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



MEGAHIT-MaxBin2-3108596.004.fa

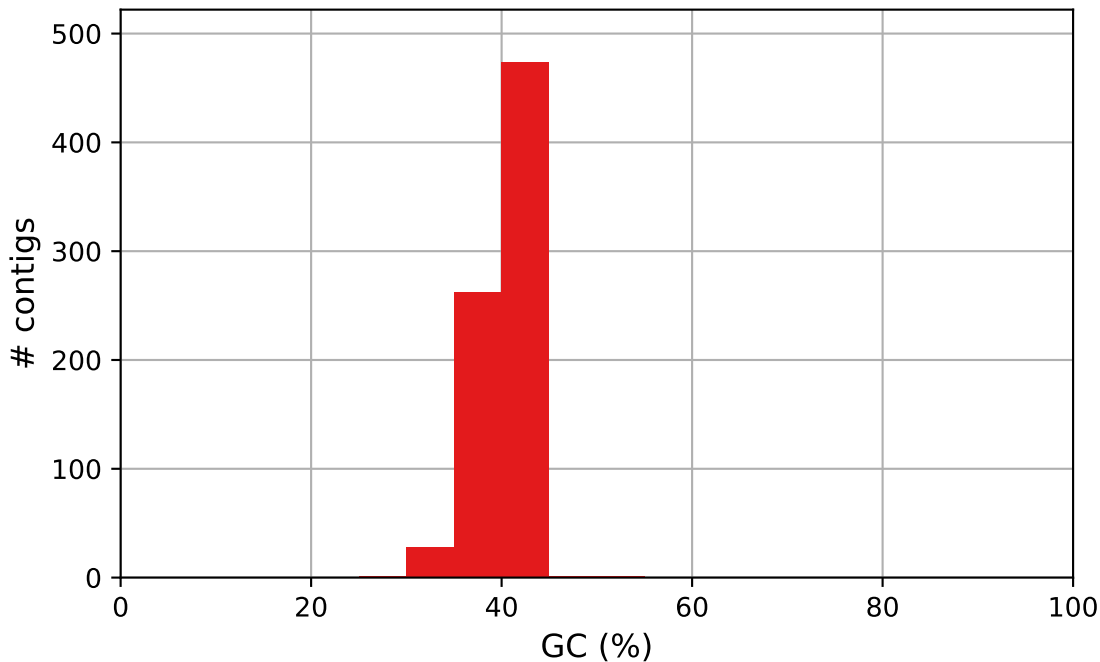


MEGAHIT-MaxBin2-3108596.004.fa



MEGAHit-MaxBin2-3108596.004.fa

MEGAHIT-MaxBin2-3108596.004.fa GC content



MEGAHit-MaxBin2-3108596.004.fa