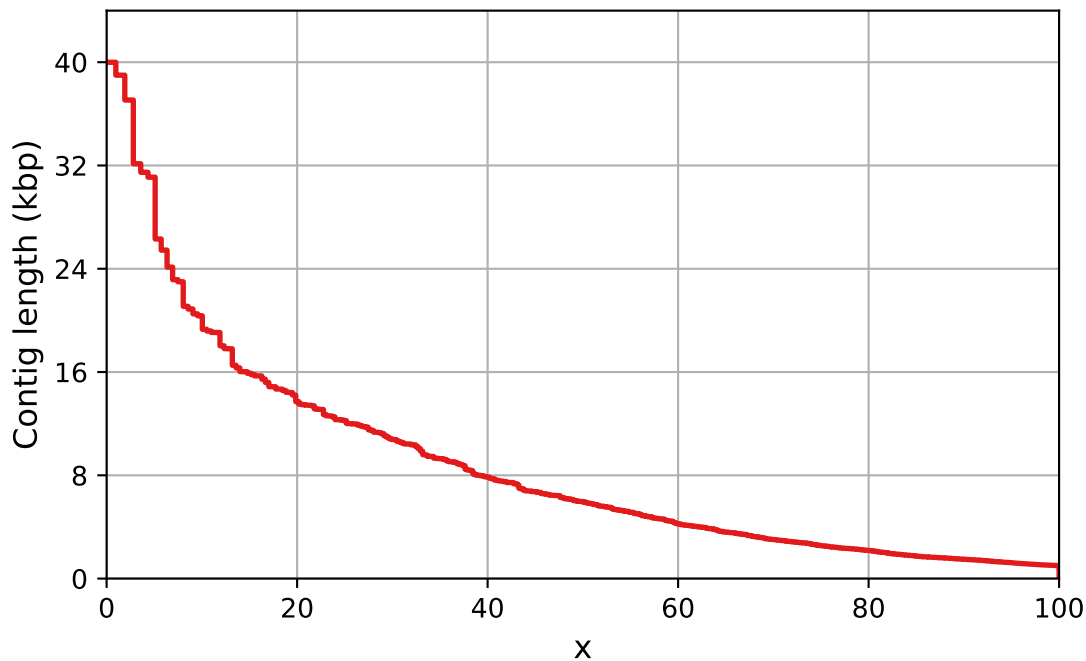


Report

	MEGAHIT-MaxBin2-3106384.005.fa
# contigs (>= 0 bp)	1113
# contigs (>= 1000 bp)	1113
# contigs (>= 5000 bp)	225
# contigs (>= 10000 bp)	86
# contigs (>= 25000 bp)	8
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	4136241
Total length (>= 1000 bp)	4136241
Total length (>= 5000 bp)	2312542
Total length (>= 10000 bp)	1362870
Total length (>= 25000 bp)	262506
Total length (>= 50000 bp)	0
# contigs	1113
Largest contig	39994
Total length	4136241
GC (%)	39.06
N50	5945
N75	2544
L50	181
L75	451
# N's per 100 kbp	0.00
# predicted rRNA genes	3 + 2 part

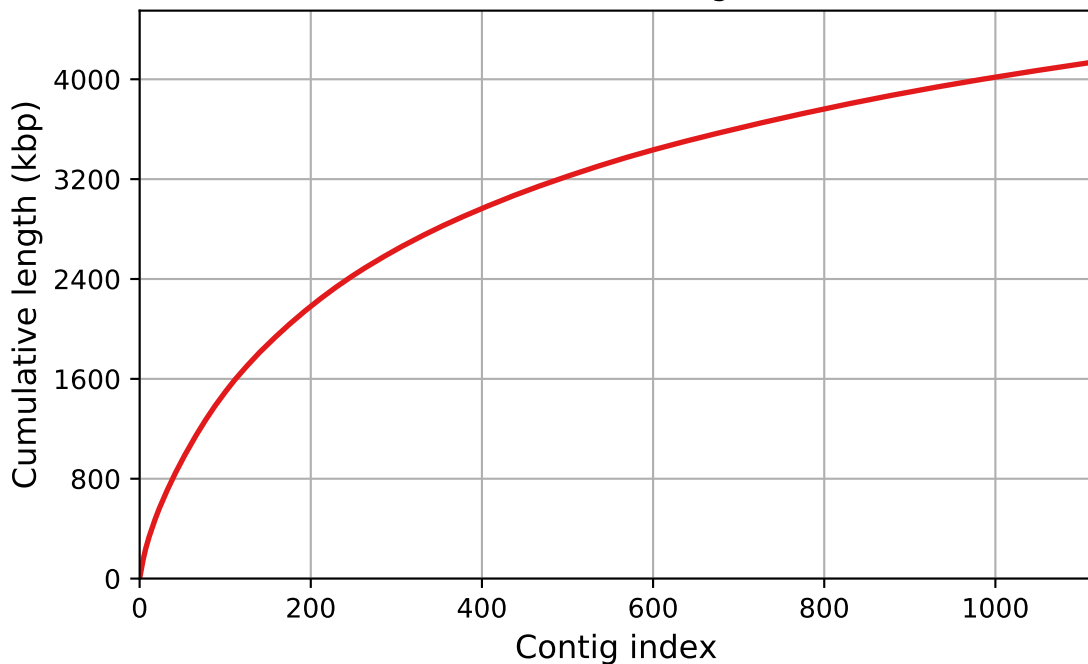
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

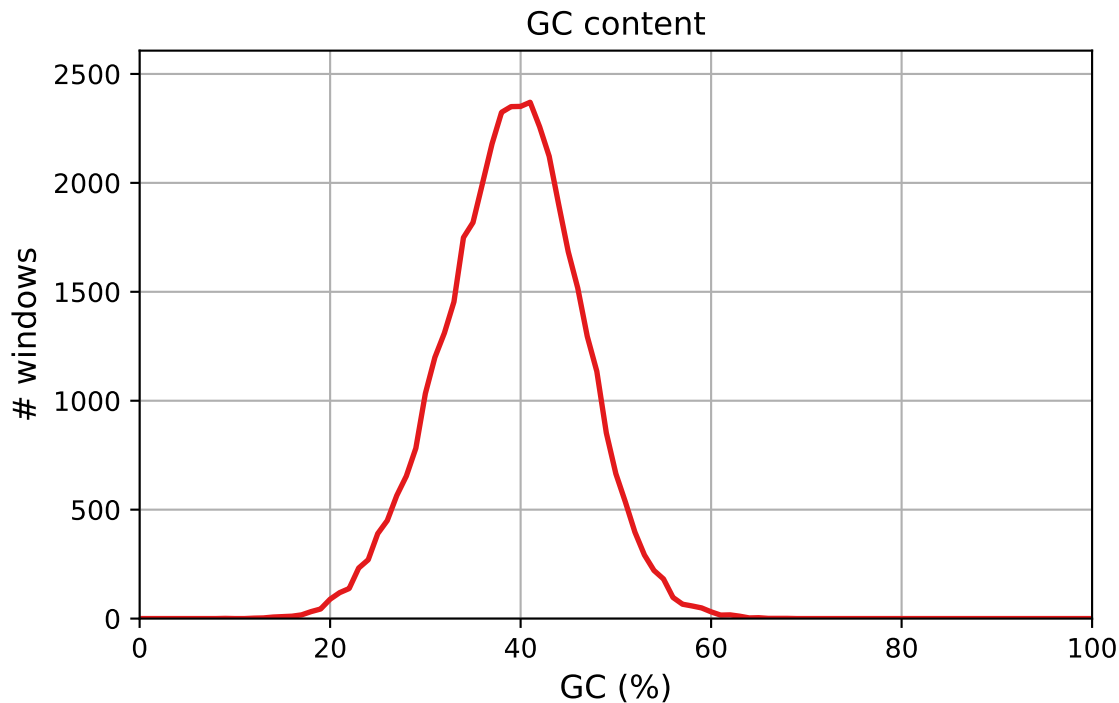


MEGAHIT-MaxBin2-3106384.005.fa

Cumulative length

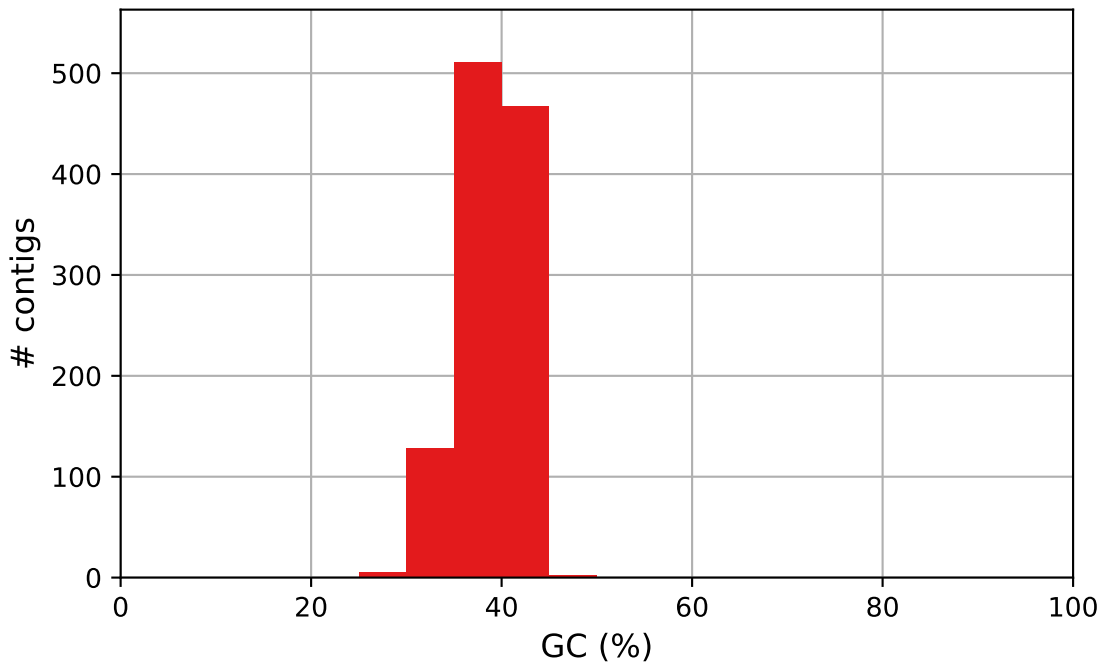


MEGAHIT-MaxBin2-3106384.005.fa



MEGAHit-MaxBin2-3106384.005.fa

MEGAHIT-MaxBin2-3106384.005.fa GC content



MEGAHit-MaxBin2-3106384.005.fa