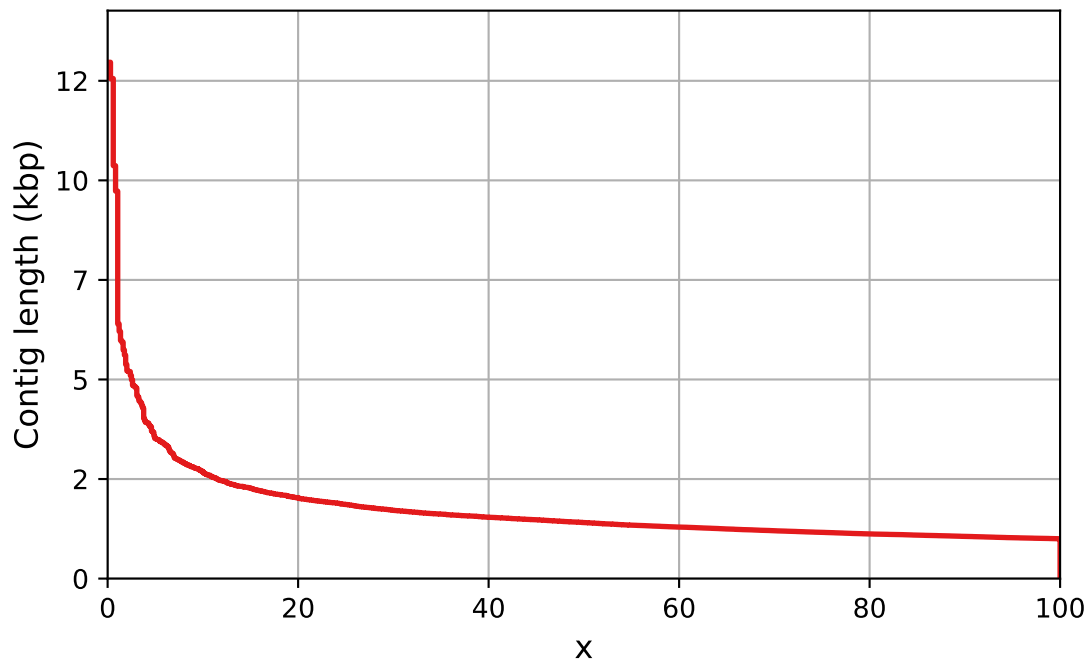


Report

	MEGAHIT-MaxBin2-3106384.007.fa
# contigs (>= 0 bp)	2941
# contigs (>= 1000 bp)	2941
# contigs (>= 5000 bp)	15
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	4304918
Total length (>= 1000 bp)	4304918
Total length (>= 5000 bp)	107583
Total length (>= 10000 bp)	35891
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	2941
Largest contig	12965
Total length	4304918
GC (%)	48.69
N50	1406
N75	1160
L50	1092
L75	1939
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 0 part

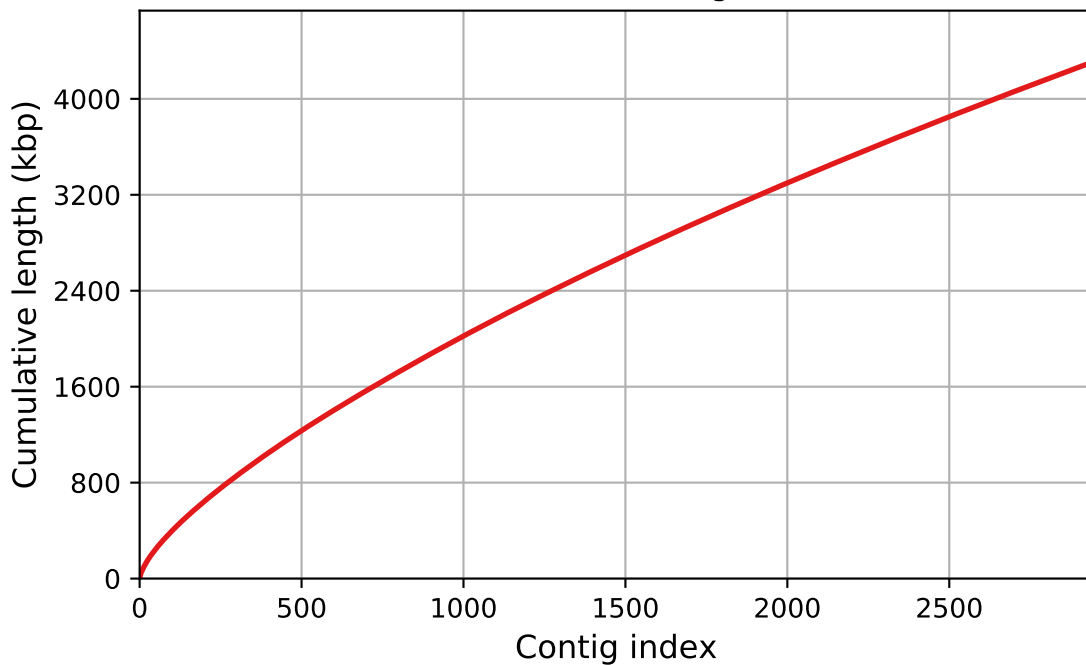
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



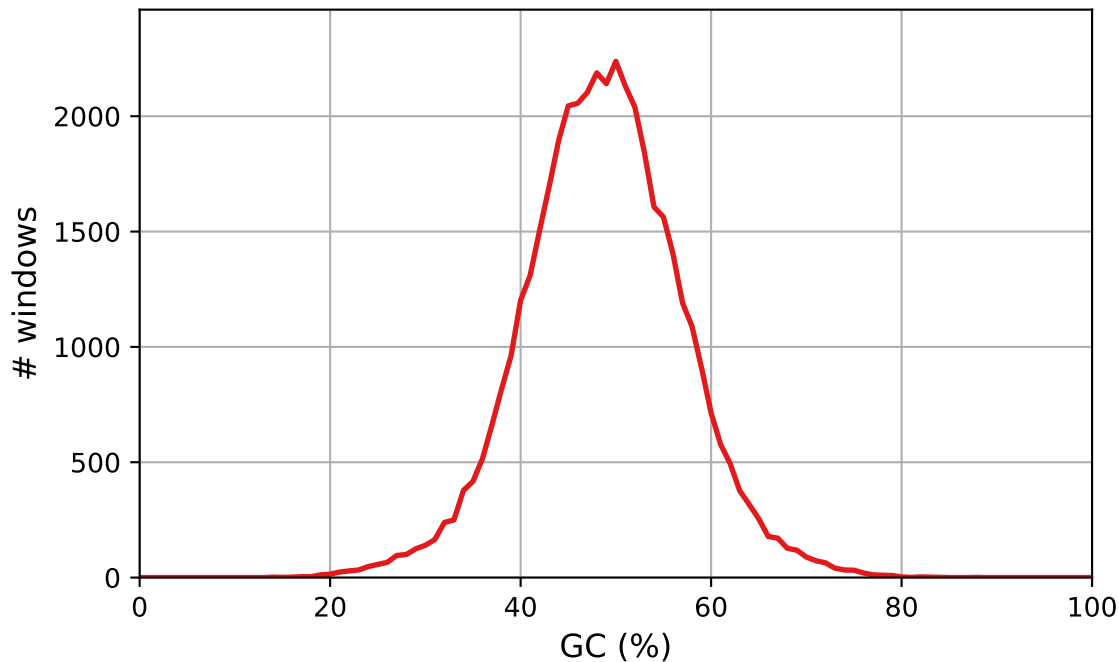
MEGAHIT-MaxBin2-3106384.007.fa

Cumulative length



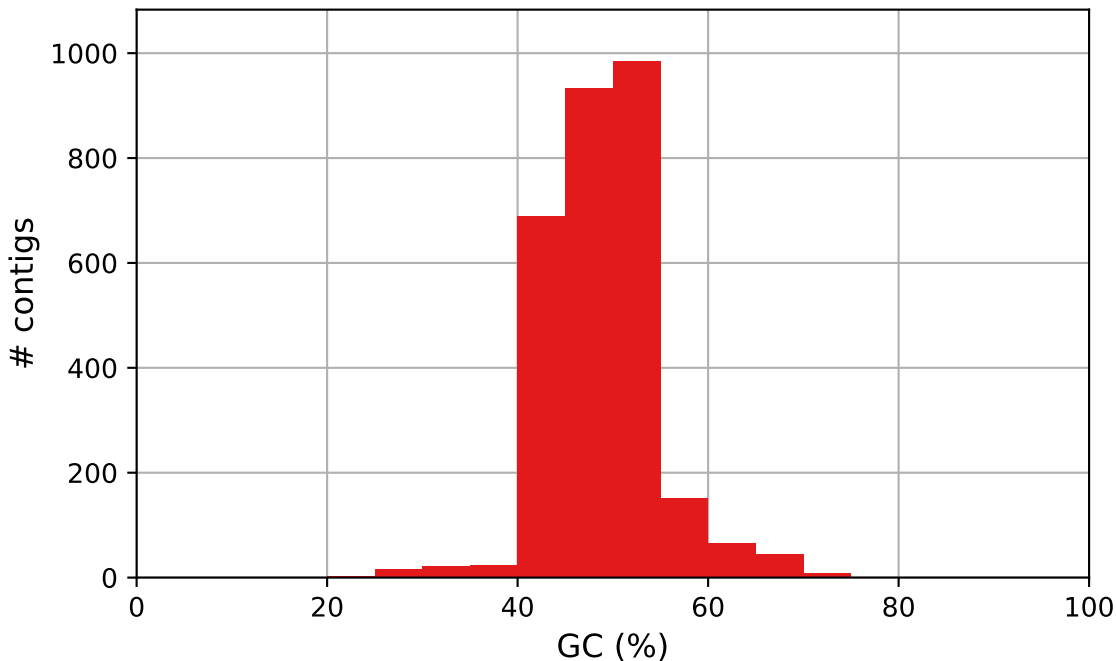
— MEGAHIT-MaxBin2-3106384.007.fa

GC content



MEGAHit-MaxBin2-3106384.007.fa

MEGAHIT-MaxBin2-3106384.007.fa GC content



MEGAHit-MaxBin2-3106384.007.fa