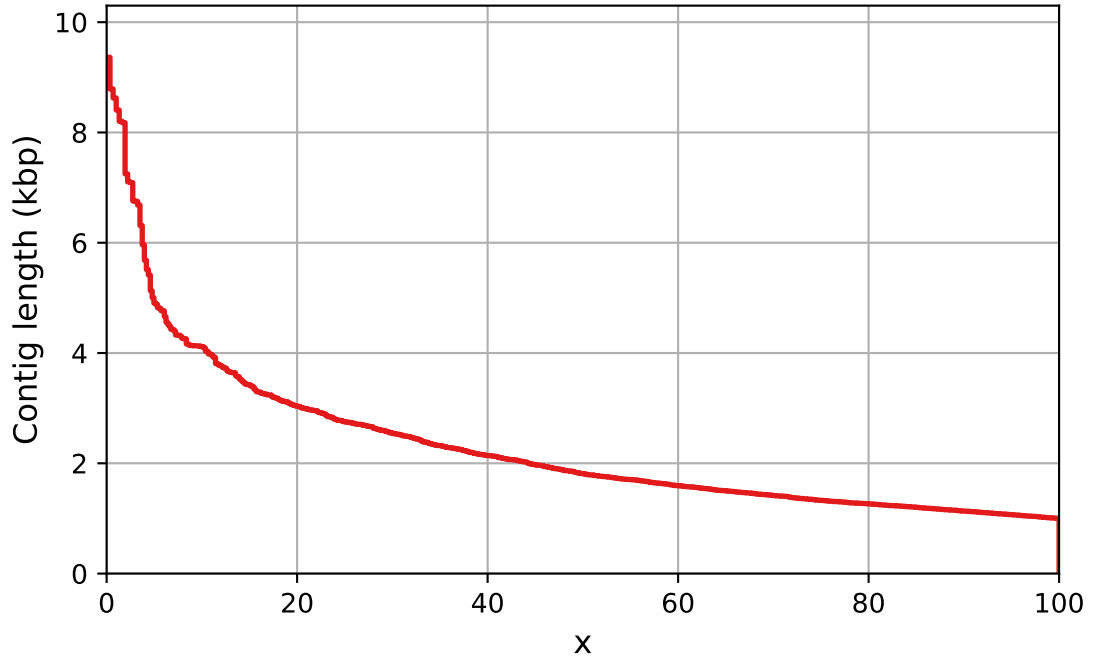


Report

	MEGAHIT-MaxBin2-3101140.007.fa
# contigs (>= 0 bp)	1477
# contigs (>= 1000 bp)	1477
# contigs (>= 5000 bp)	19
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2661750
Total length (>= 1000 bp)	2661750
Total length (>= 5000 bp)	132227
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	1477
Largest contig	9363
Total length	2661750
GC (%)	50.43
N50	1814
N75	1328
L50	472
L75	903
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 0 part

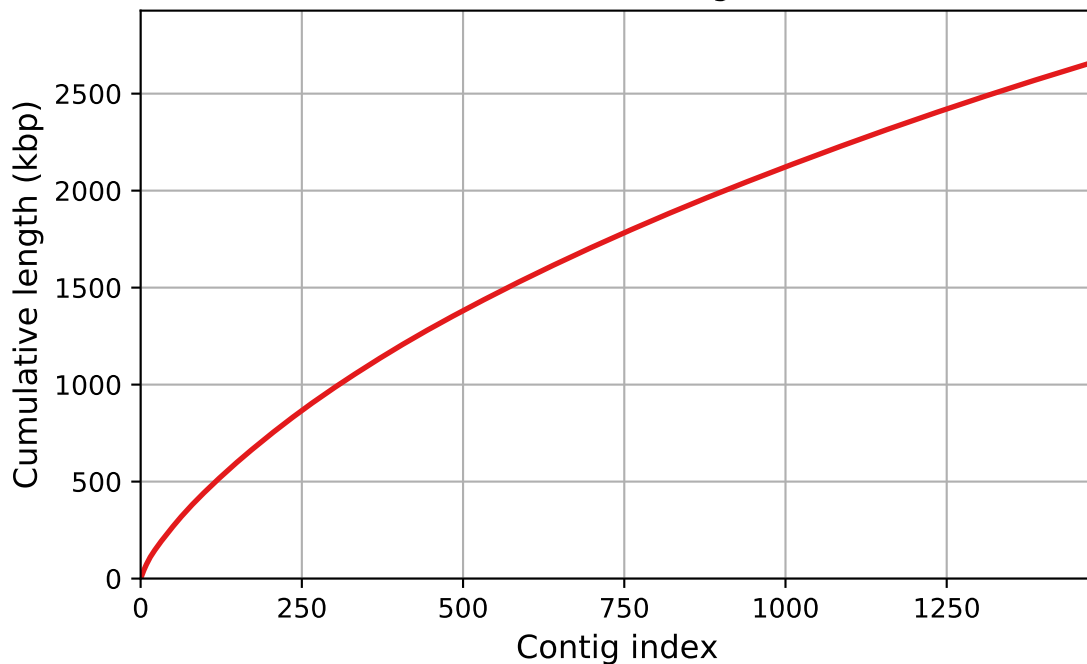
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



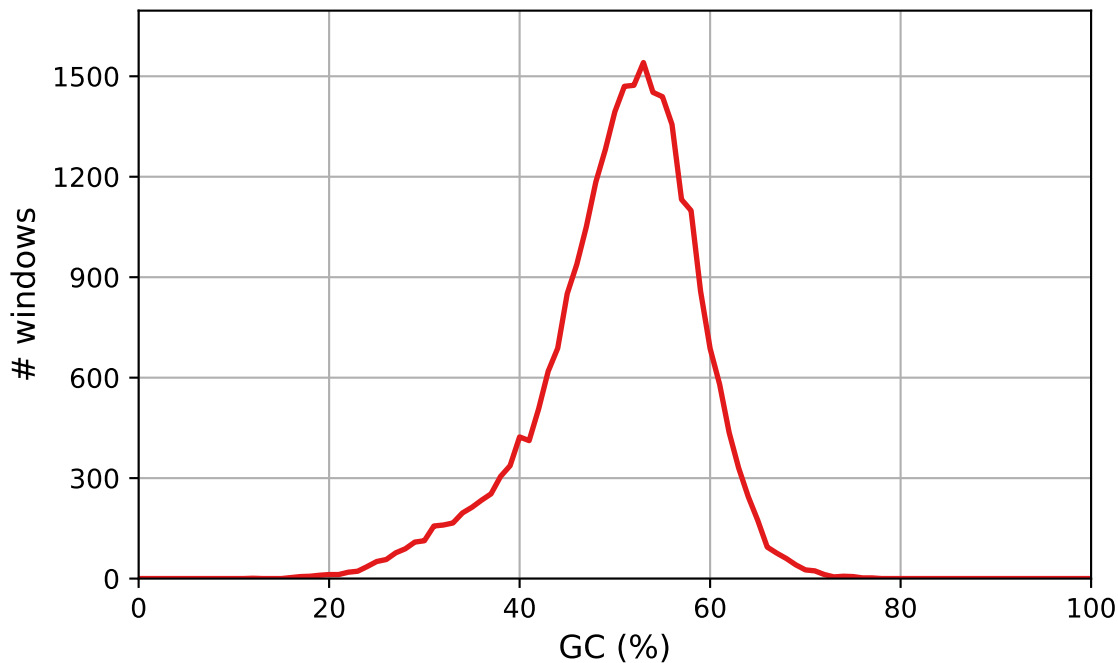
MEGAHIT-MaxBin2-3101140.007.fa

Cumulative length



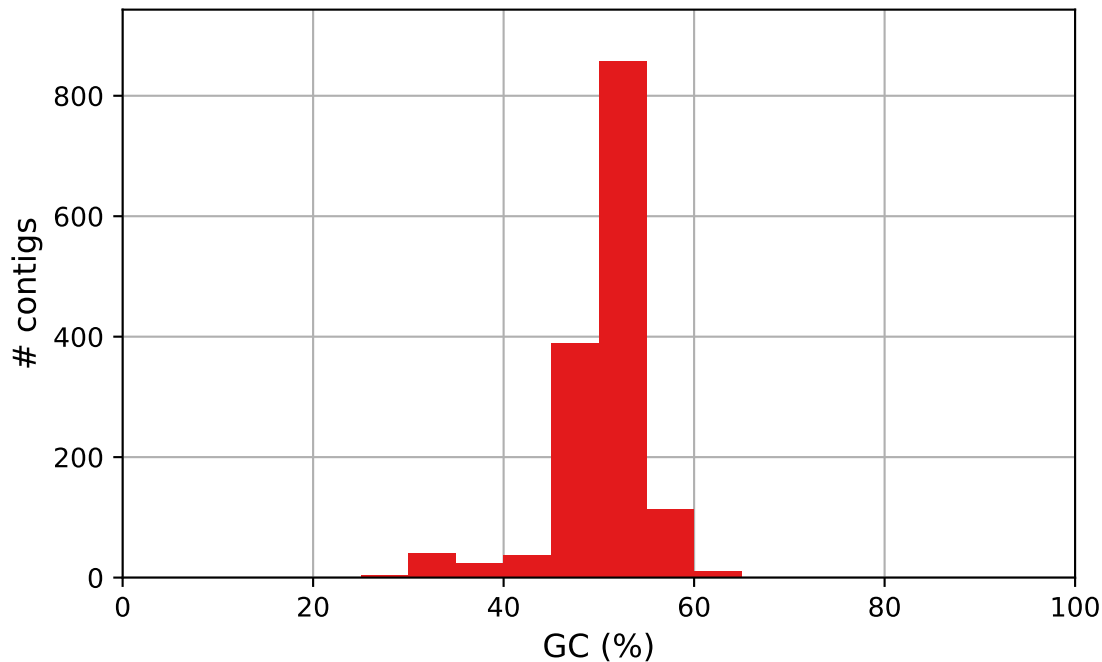
MEGAHIT-MaxBin2-3101140.007.fa

GC content



MEGAHIT-MaxBin2-3101140.007.fa

MEGAHIT-MaxBin2-3101140.007.fa GC content



MEGAHit-MaxBin2-3101140.007.fa