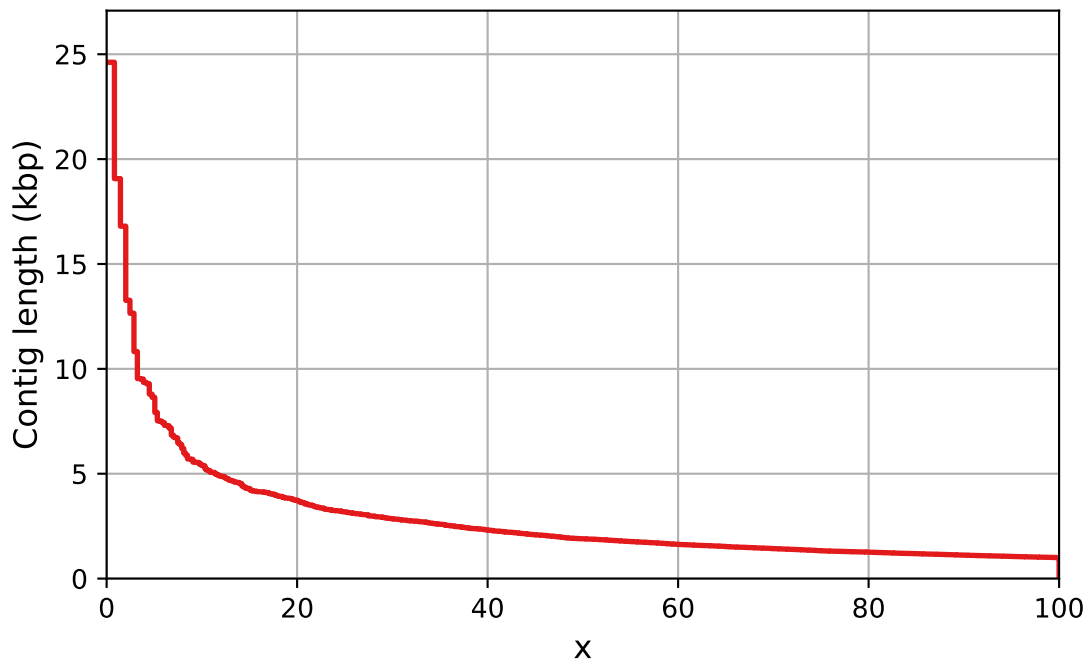


# Report

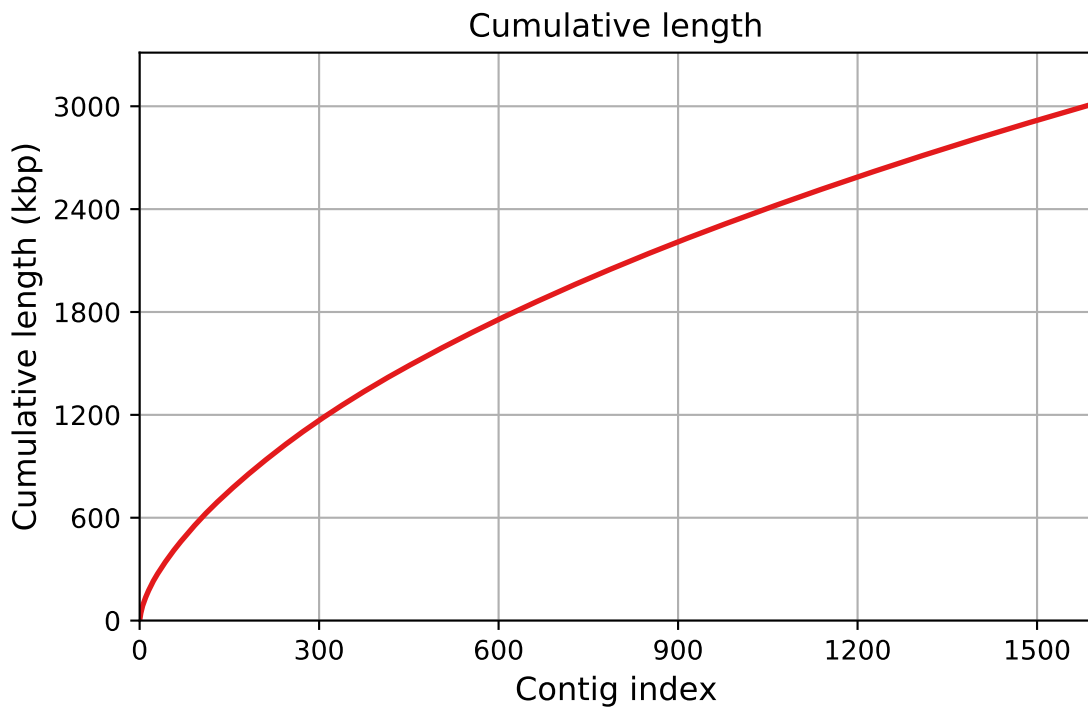
	MEGAHIT-MaxBin2-3108596.010.fa
# contigs (>= 0 bp)	1592
# contigs (>= 1000 bp)	1592
# contigs (>= 5000 bp)	43
# contigs (>= 10000 bp)	6
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3011651
Total length (>= 1000 bp)	3011651
Total length (>= 5000 bp)	341901
Total length (>= 10000 bp)	97229
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	1592
Largest contig	24617
Total length	3011651
GC (%)	52.40
N50	1897
N75	1326
L50	460
L75	937
# N's per 100 kbp	0.00
# predicted rRNA genes	4 + 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

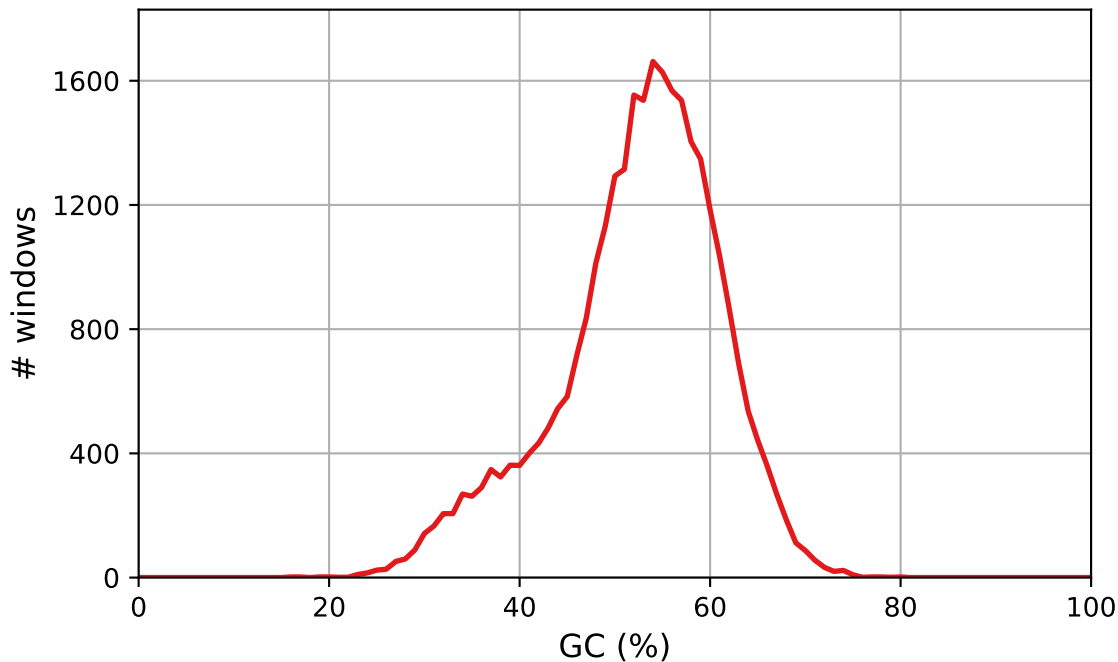


MEGAHIT-MaxBin2-3108596.010.fa



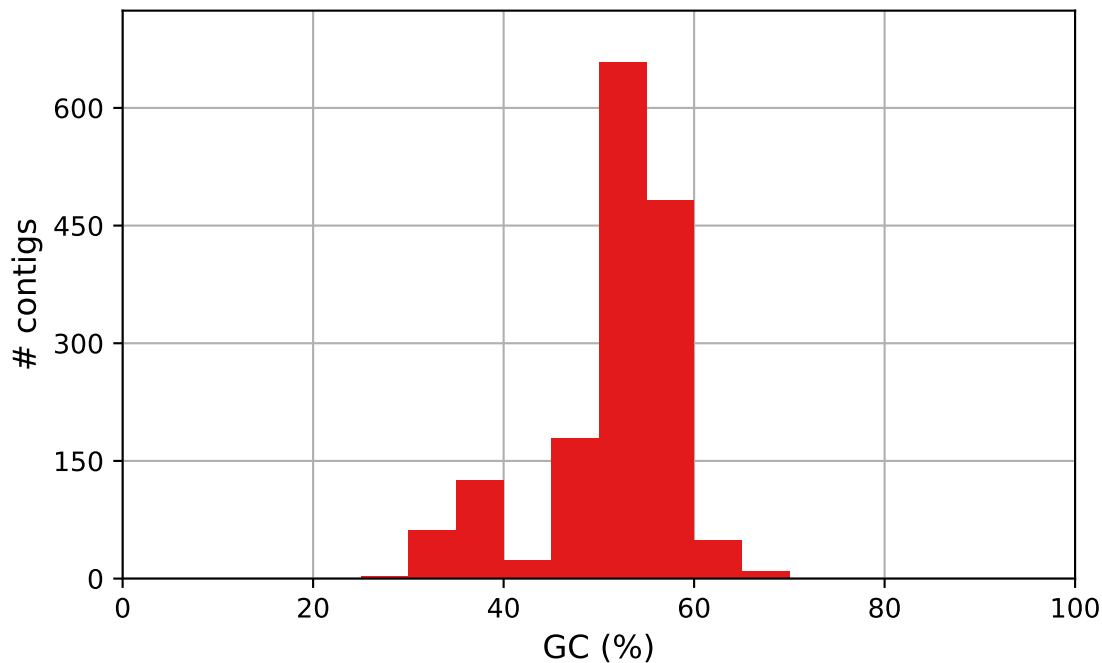
— MEGAHIT-MaxBin2-3108596.010.fa

## GC content



MEGAHIT-MaxBin2-3108596.010.fa

MEGAHIT-MaxBin2-3108596.010.fa GC content



MEGAHit-MaxBin2-3108596.010.fa