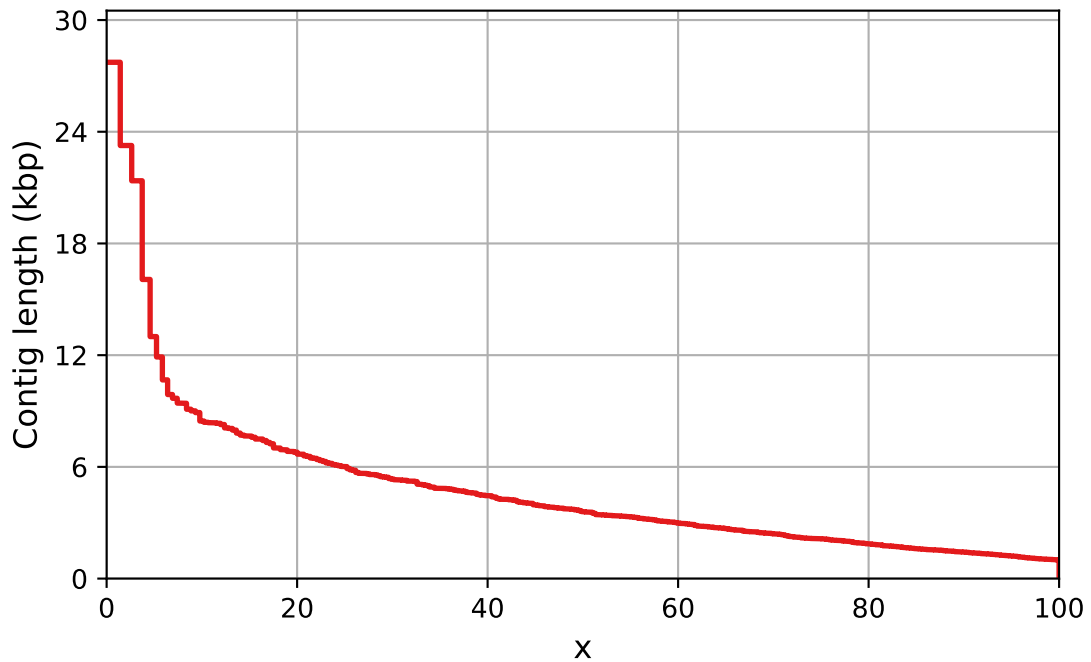


# Report

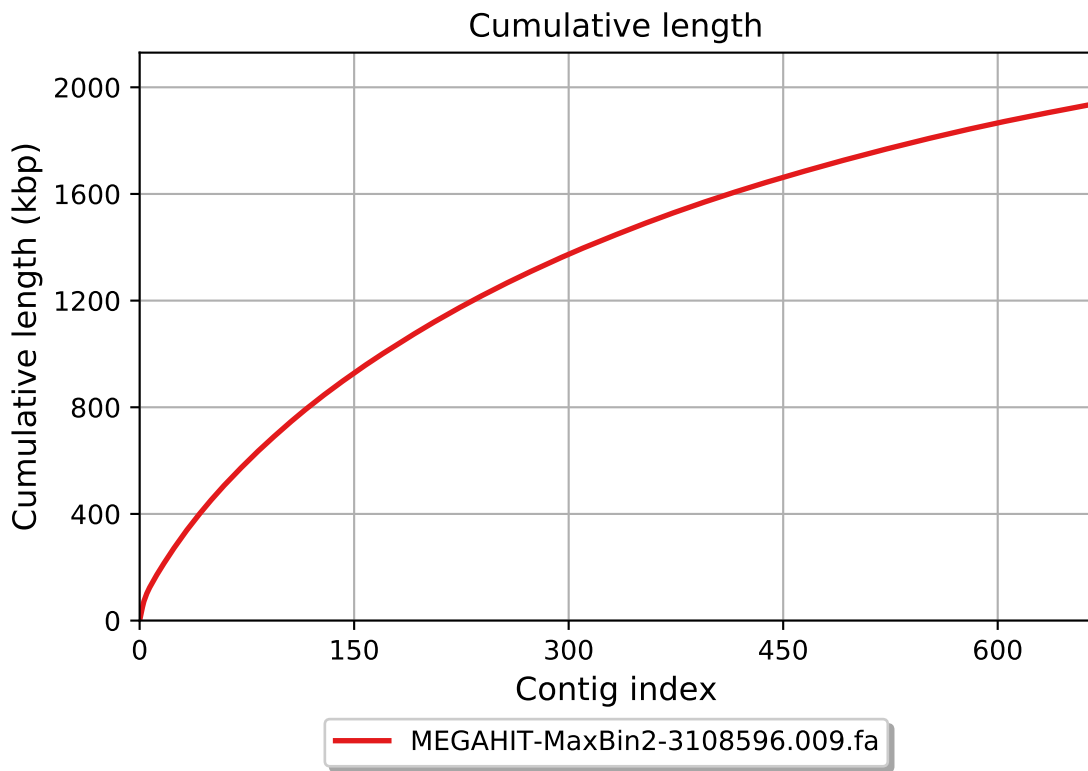
	MEGAHIT-MaxBin2-3108596.009.fa
# contigs (>= 0 bp)	666
# contigs (>= 1000 bp)	666
# contigs (>= 5000 bp)	86
# contigs (>= 10000 bp)	7
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1936332
Total length (>= 1000 bp)	1936332
Total length (>= 5000 bp)	651437
Total length (>= 10000 bp)	124000
Total length (>= 25000 bp)	27731
Total length (>= 50000 bp)	0
# contigs	666
Largest contig	27731
Total length	1936332
GC (%)	59.51
N50	3598
N75	2139
L50	161
L75	336
# N's per 100 kbp	0.00
# predicted rRNA genes	1 + 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).

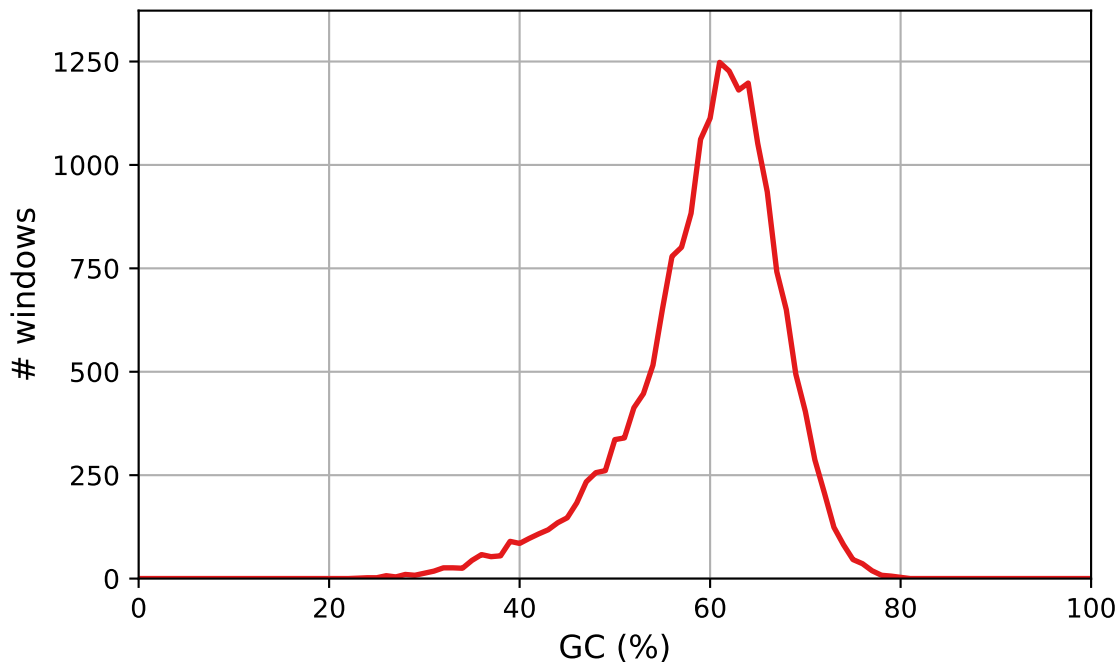
Nx



MEGAHIT-MaxBin2-3108596.009.fa

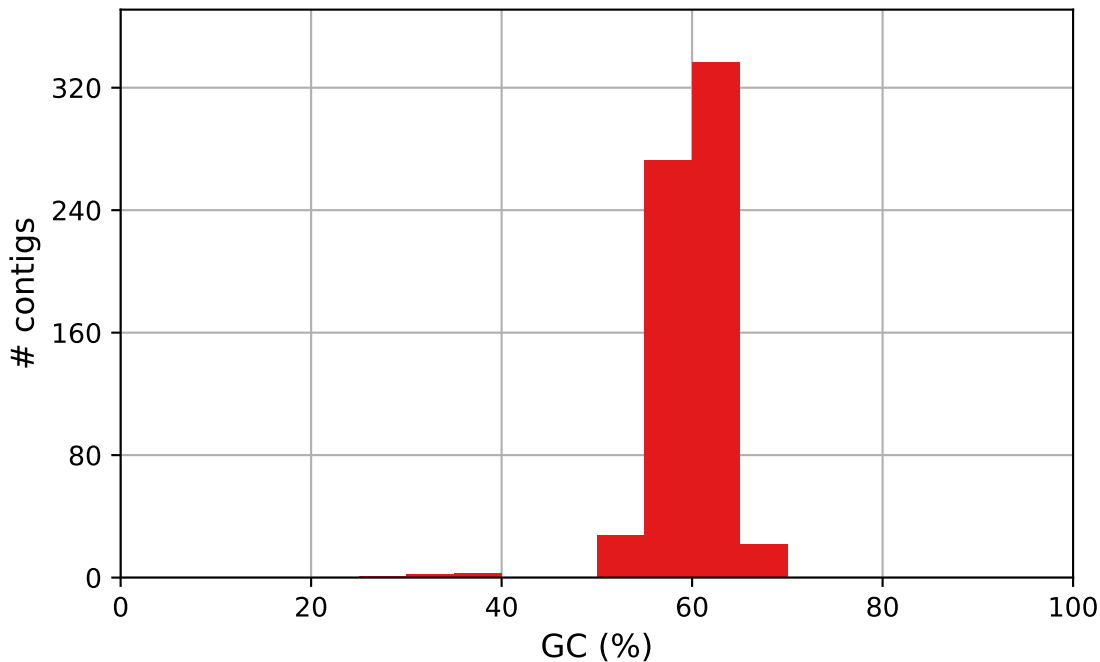


## GC content



MEGAHit-MaxBin2-3108596.009.fa

MEGAHIT-MaxBin2-3108596.009.fa GC content



MEGAHit-MaxBin2-3108596.009.fa