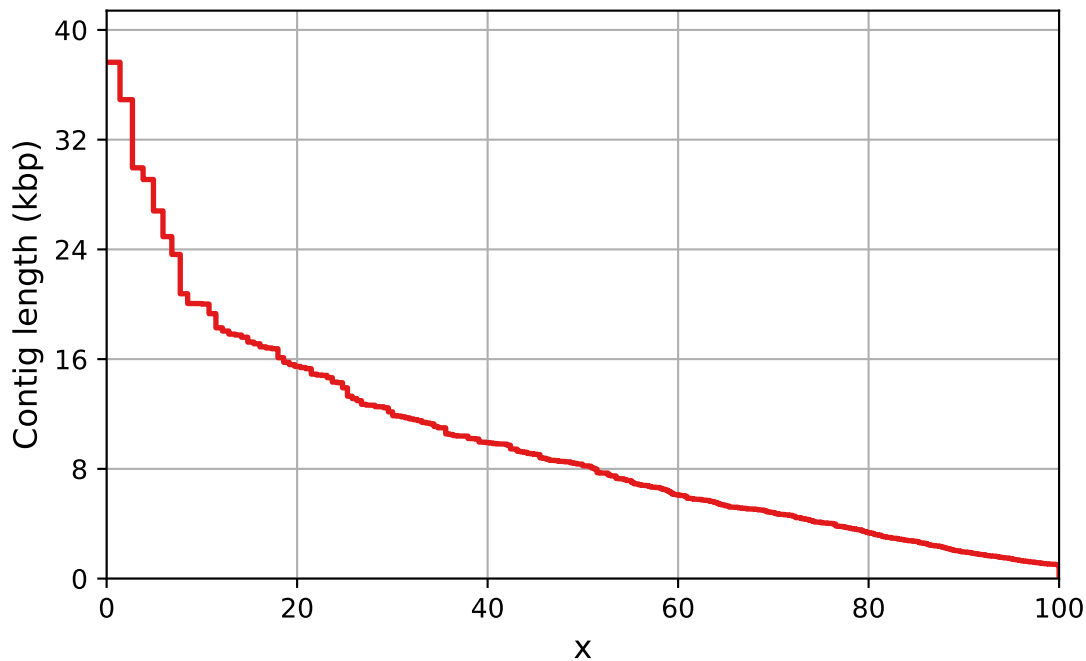


Report

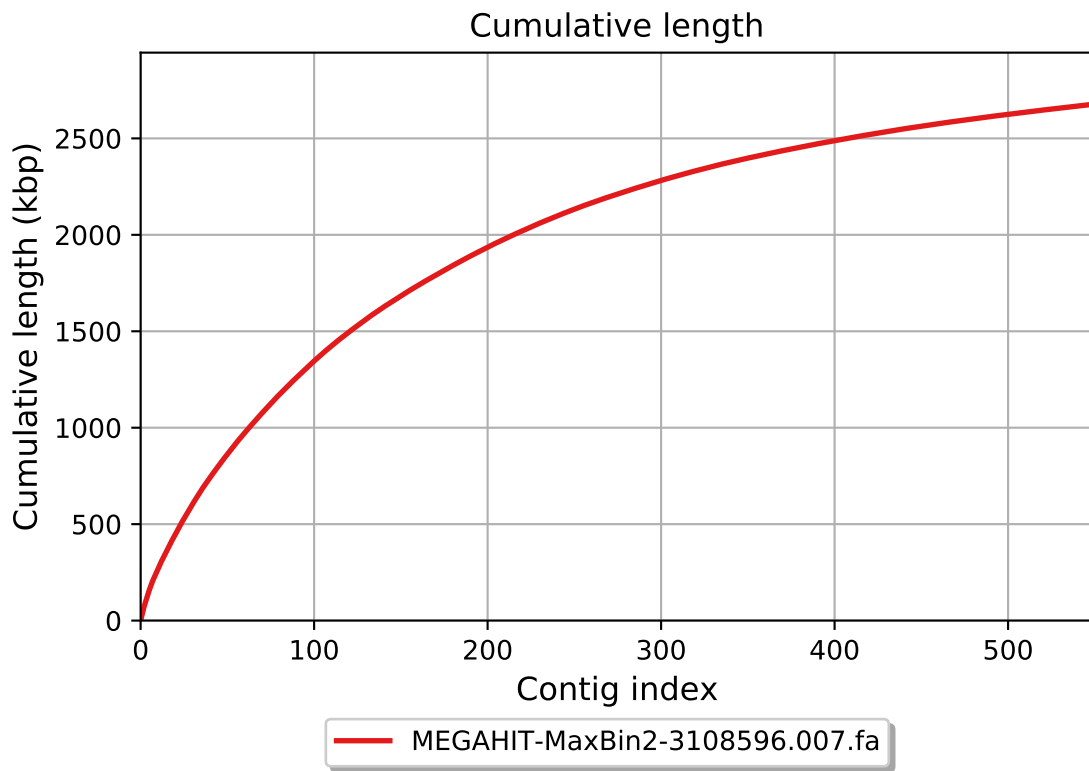
	MEGAHIT-MaxBin2-3108596.007.fa
# contigs (>= 0 bp)	549
# contigs (>= 1000 bp)	549
# contigs (>= 5000 bp)	179
# contigs (>= 10000 bp)	67
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2676952
Total length (>= 1000 bp)	2676952
Total length (>= 5000 bp)	1836305
Total length (>= 10000 bp)	1046242
Total length (>= 25000 bp)	158406
Total length (>= 50000 bp)	0
# contigs	549
Largest contig	37642
Total length	2676952
GC (%)	54.71
N50	8228
N75	4109
L50	100
L75	217
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 1 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).

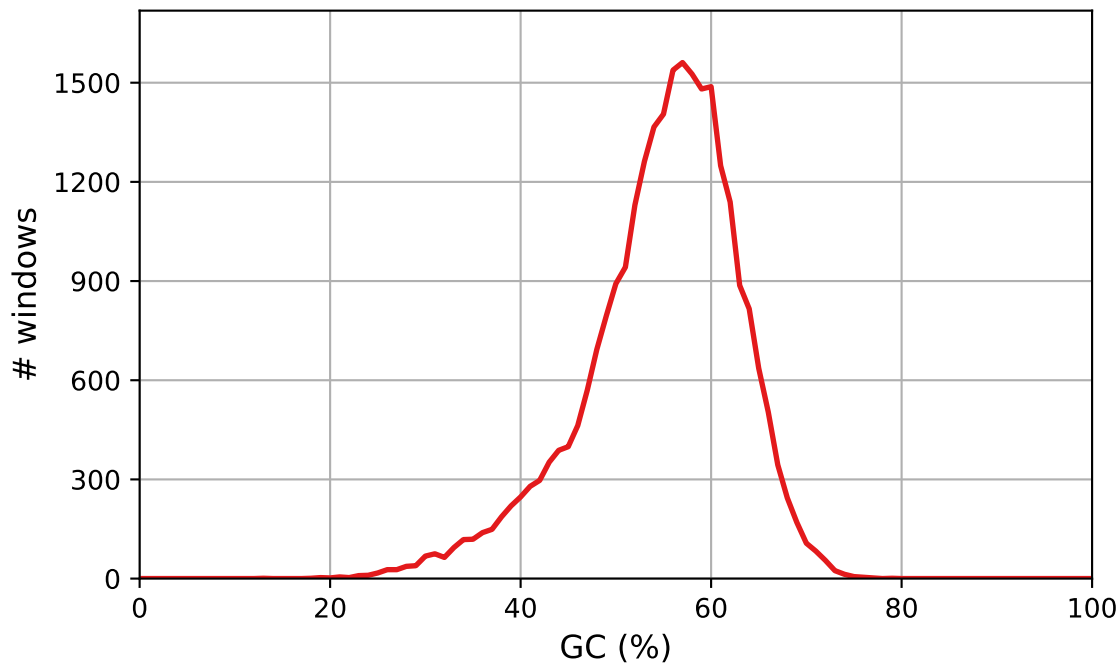
Nx



MEGAHIT-MaxBin2-3108596.007.fa

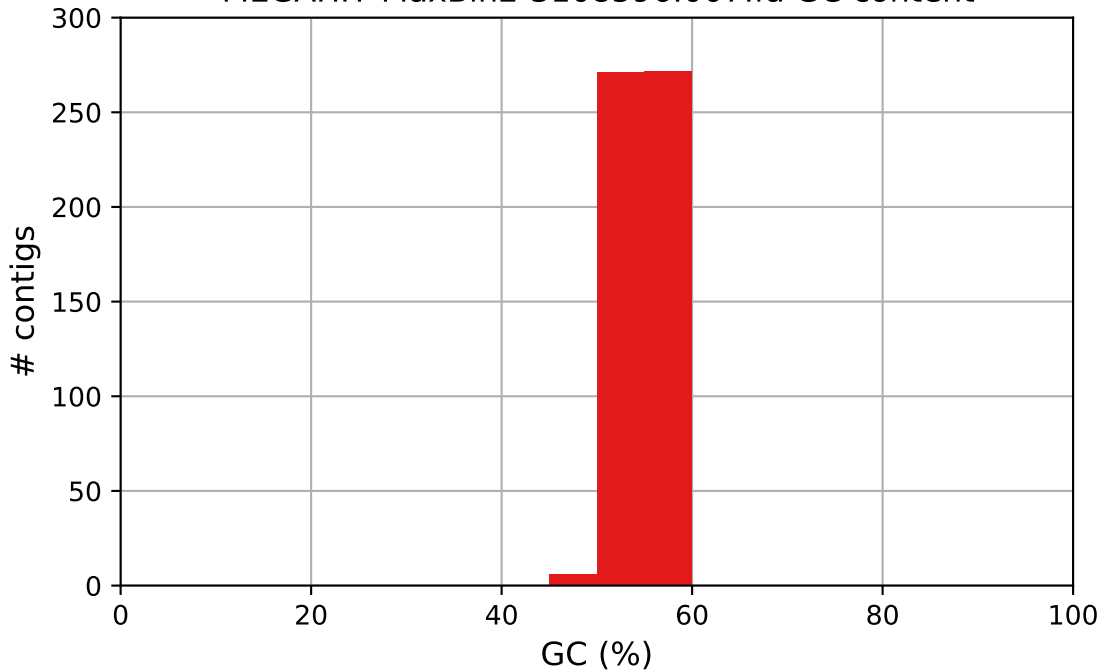


GC content



MEGAHit-MaxBin2-3108596.007.fa

MEGAHIT-MaxBin2-3108596.007.fa GC content



MEGAhit-MaxBin2-3108596.007.fa