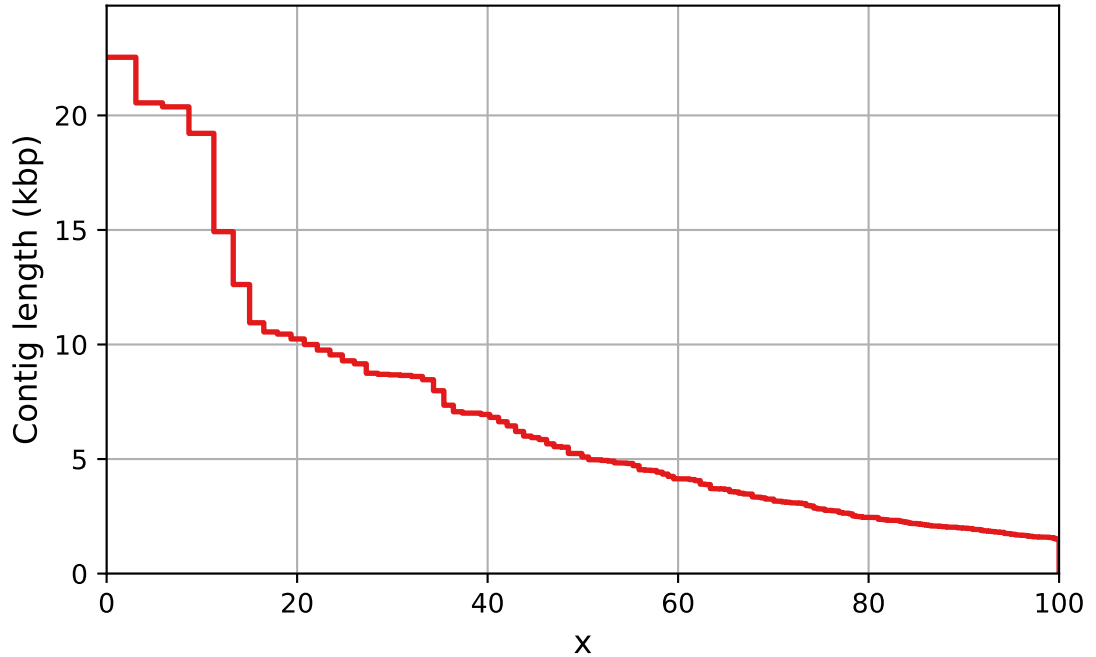


Report

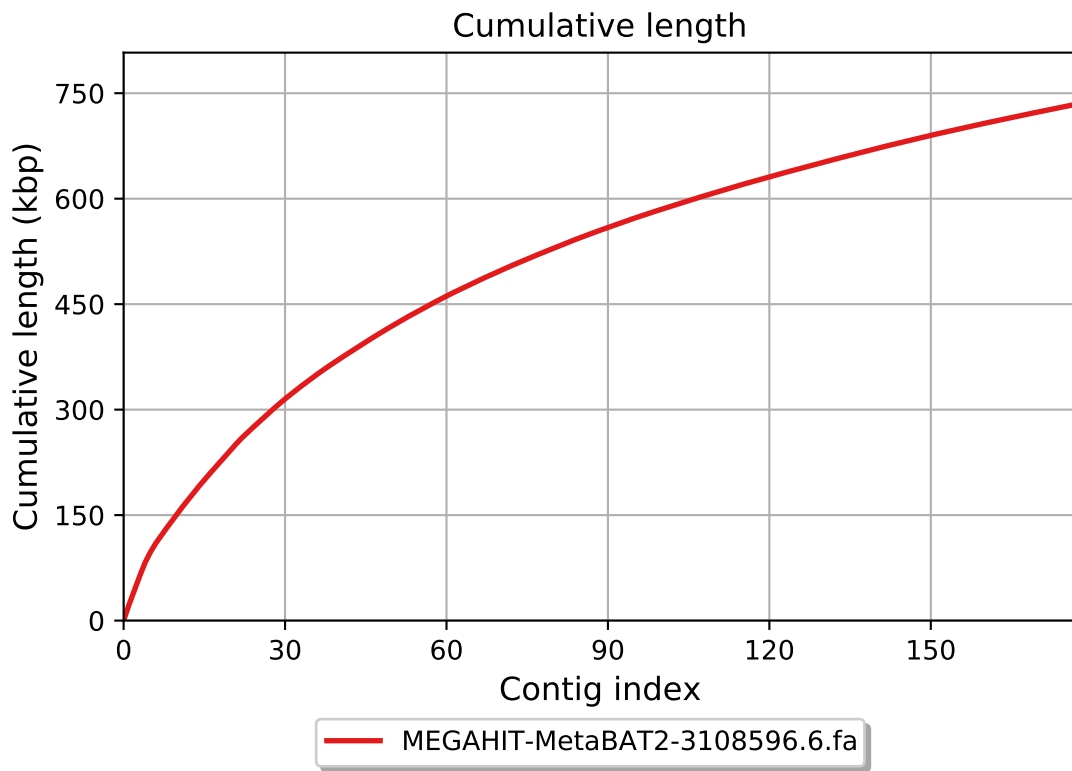
	MEGAHIT-MetaBAT2-3108596.6.fa
# contigs (>= 0 bp)	177
# contigs (>= 1000 bp)	177
# contigs (>= 5000 bp)	40
# contigs (>= 10000 bp)	10
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	734278
Total length (>= 1000 bp)	734278
Total length (>= 5000 bp)	371528
Total length (>= 10000 bp)	152414
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	177
Largest contig	22538
Total length	734278
GC (%)	39.48
N50	5089
N75	2824
L50	40
L75	87
# 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).

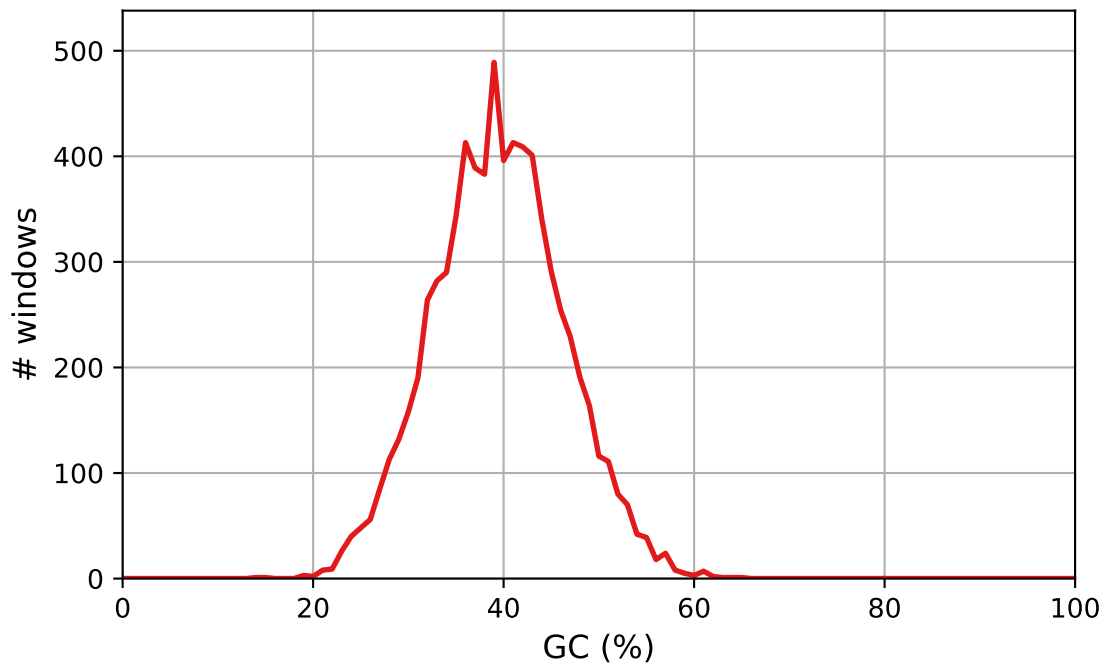
Nx



MEGAHIT-MetaBAT2-3108596.6.fa

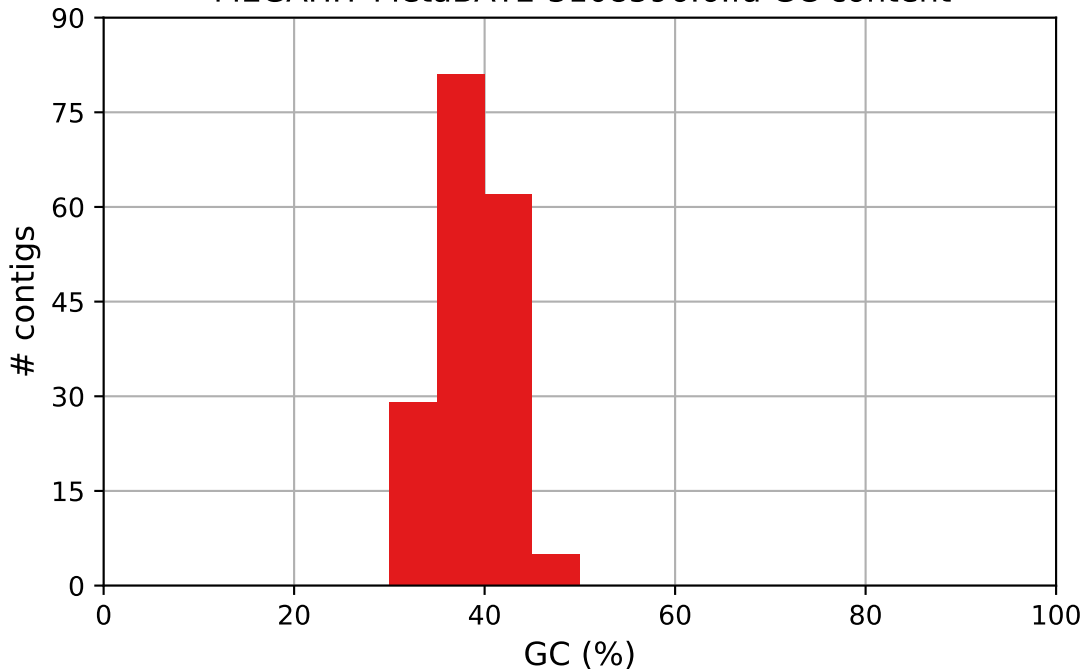


GC content



— MEGAHIT-MetaBAT2-3108596.6.fa

MEGAHIT-MetaBAT2-3108596.6.fa GC content



MEGAHit-MetaBAT2-3108596.6.fa