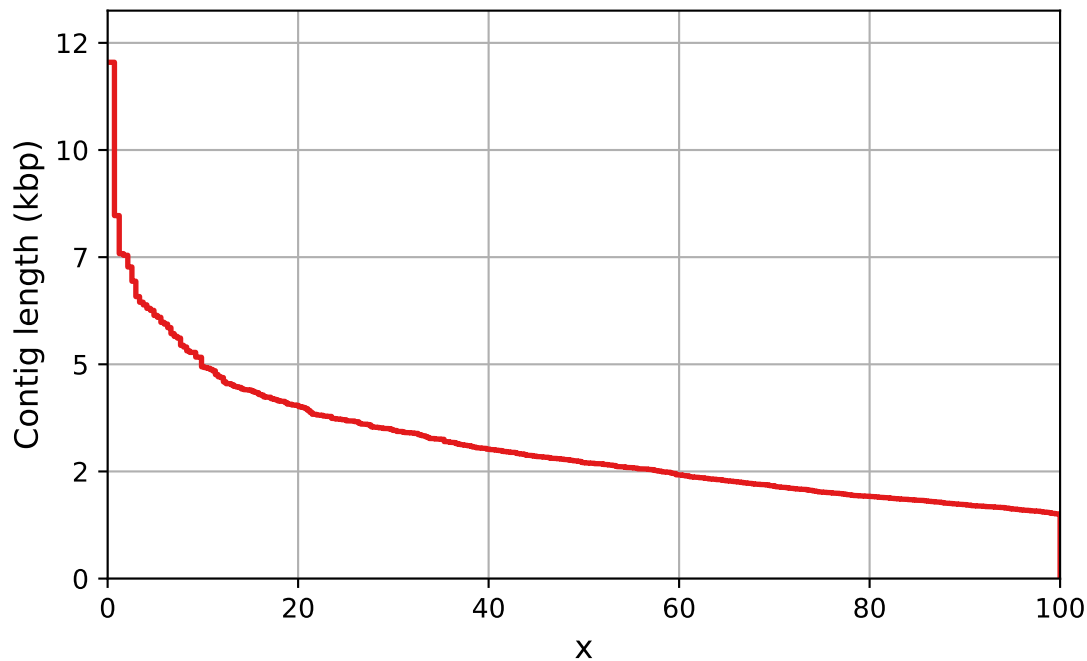


# Report

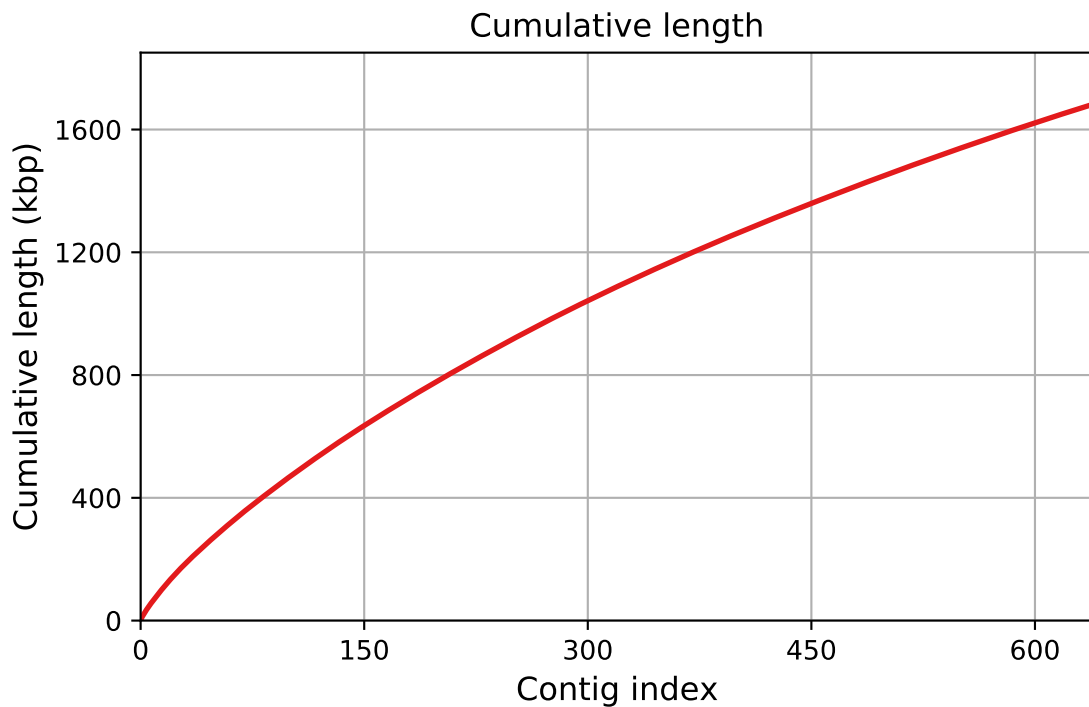
	MEGAHIT-MetaBAT2-3108596.11.fa
# contigs (>= 0 bp)	639
# contigs (>= 1000 bp)	639
# contigs (>= 5000 bp)	26
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1682314
Total length (>= 1000 bp)	1682314
Total length (>= 5000 bp)	165862
Total length (>= 10000 bp)	12047
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	639
Largest contig	12047
Total length	1682314
GC (%)	44.34
N50	2702
N75	2018
L50	222
L75	401
# 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

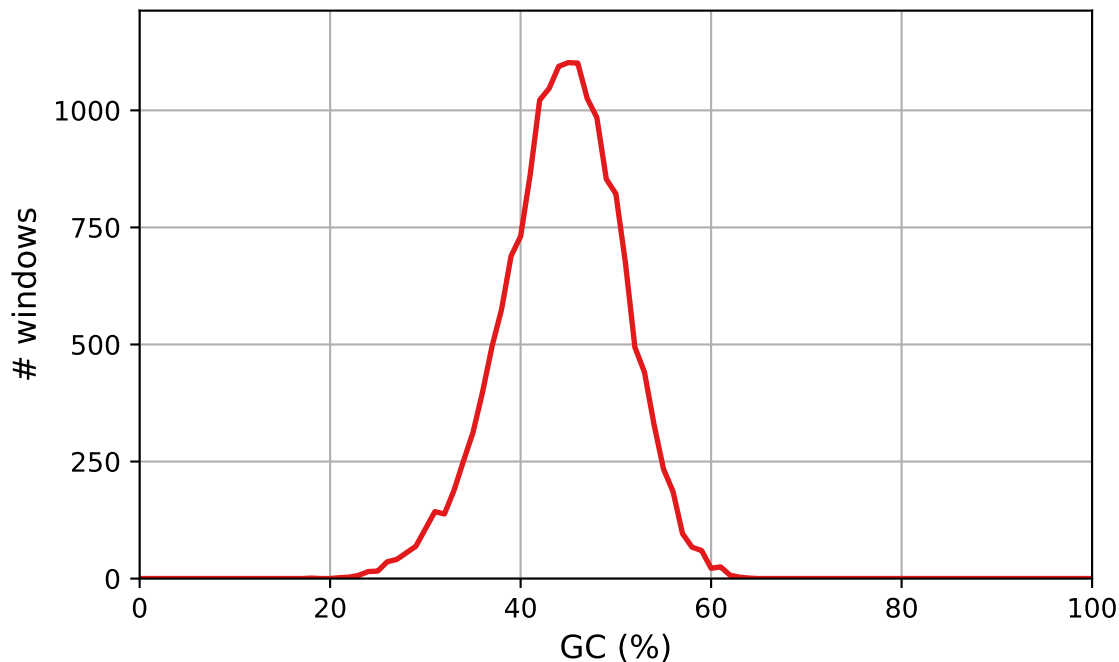


MEGAHIT-MetaBAT2-3108596.11.fa



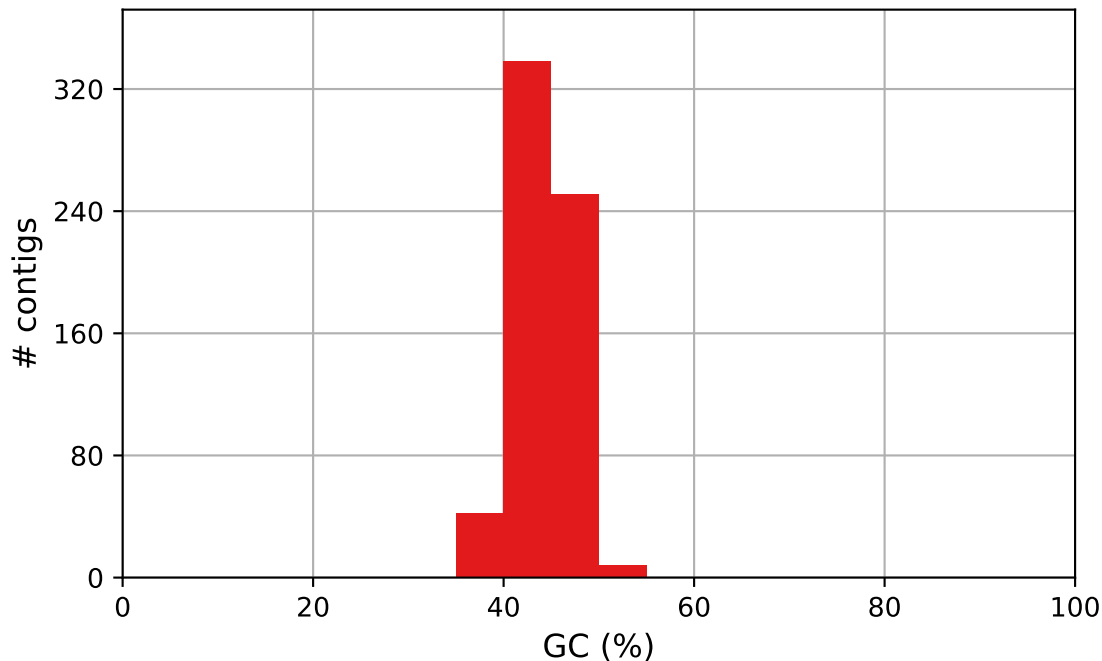
— MEGAHIT-MetaBAT2-3108596.11.fa

## GC content



MEGAHit-MetaBAT2-3108596.11.fa

MEGAHIT-MetaBAT2-3108596.11.fa GC content



MEGAHit-MetaBAT2-3108596.11.fa