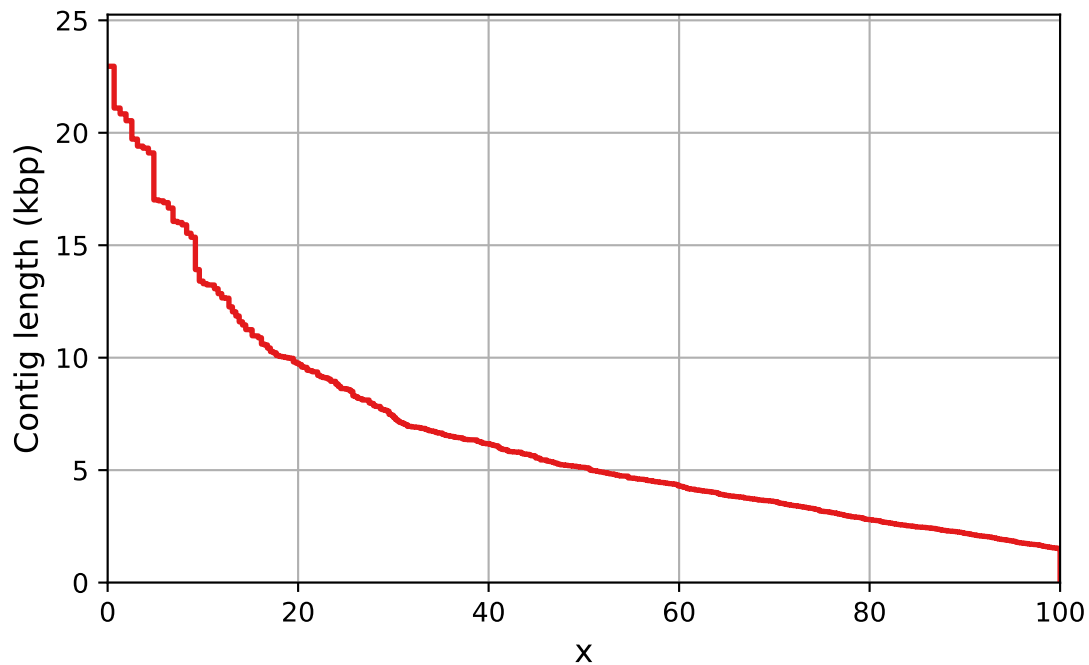


Report

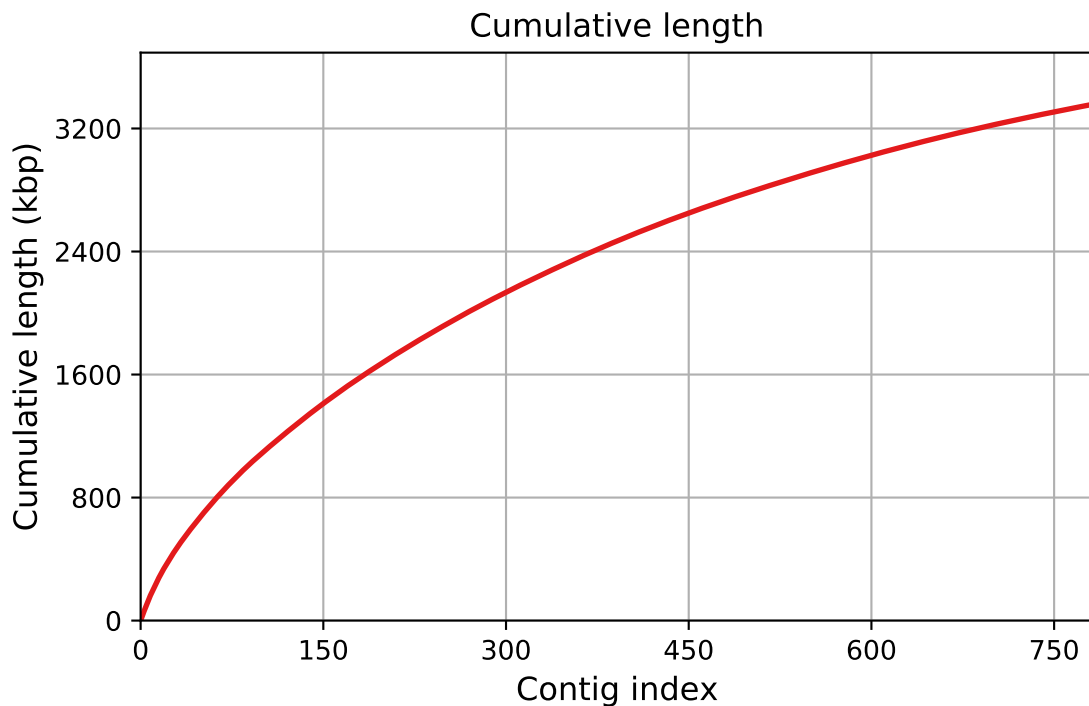
	MEGAHIT-MetaBAT2-3106237.7.fa
# contigs (>= 0 bp)	782
# contigs (>= 1000 bp)	782
# contigs (>= 5000 bp)	204
# contigs (>= 10000 bp)	45
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3357498
Total length (>= 1000 bp)	3357498
Total length (>= 5000 bp)	1704138
Total length (>= 10000 bp)	634495
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	782
Largest contig	22957
Total length	3357498
GC (%)	42.45
N50	5121
N75	3170
L50	200
L75	407
# 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

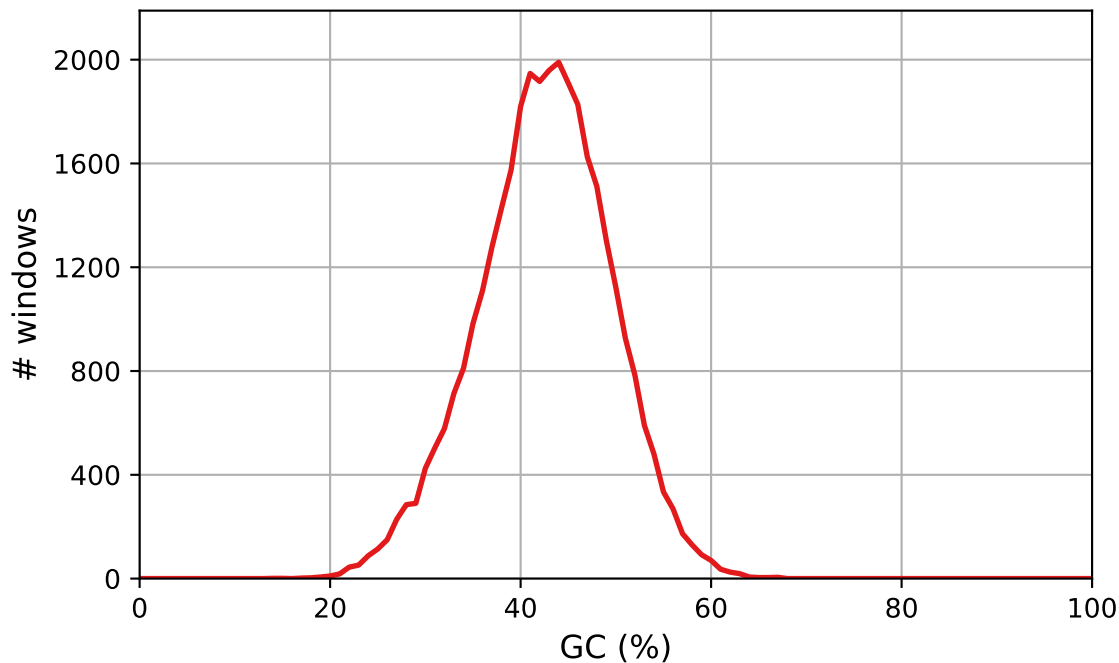


MEGAHIT-MetaBAT2-3106237.7.fa



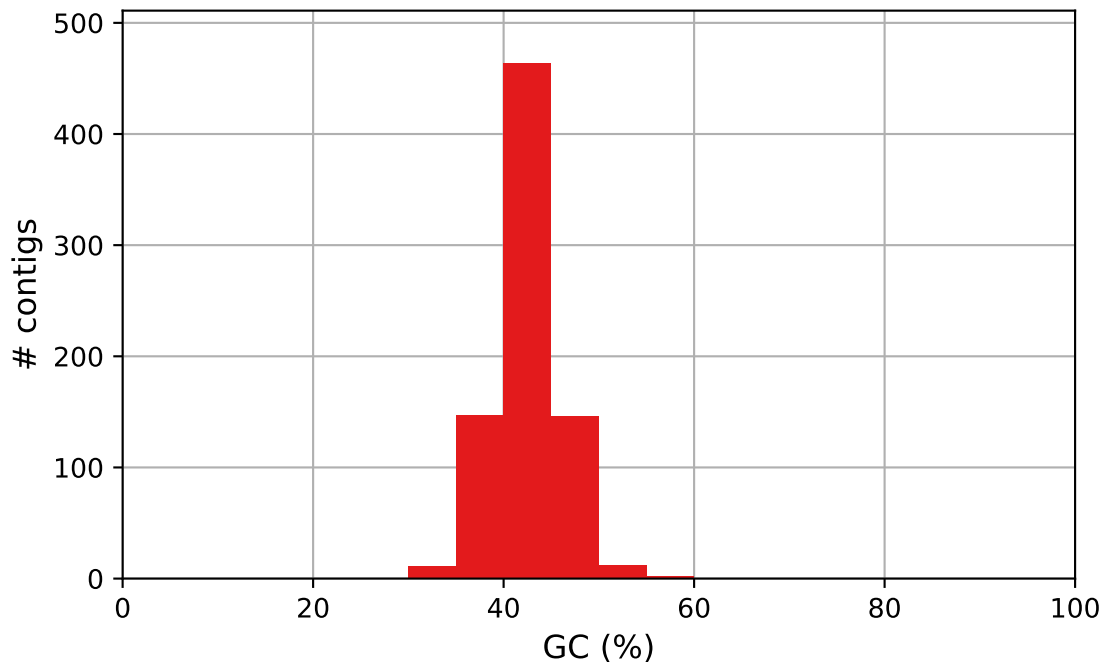
— MEGAHIT-MetaBAT2-3106237.7.fa

GC content



— MEGAHIT-MetaBAT2-3106237.7.fa

MEGAHIT-MetaBAT2-3106237.7.fa GC content



MEGAHIT-MetaBAT2-3106237.7.fa