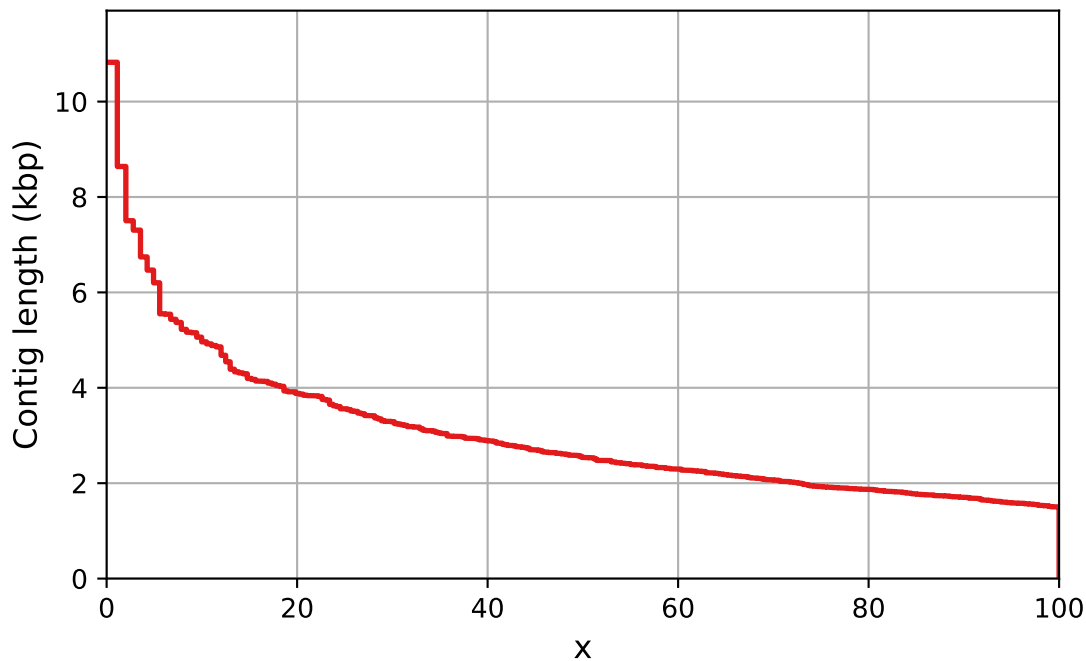


Report

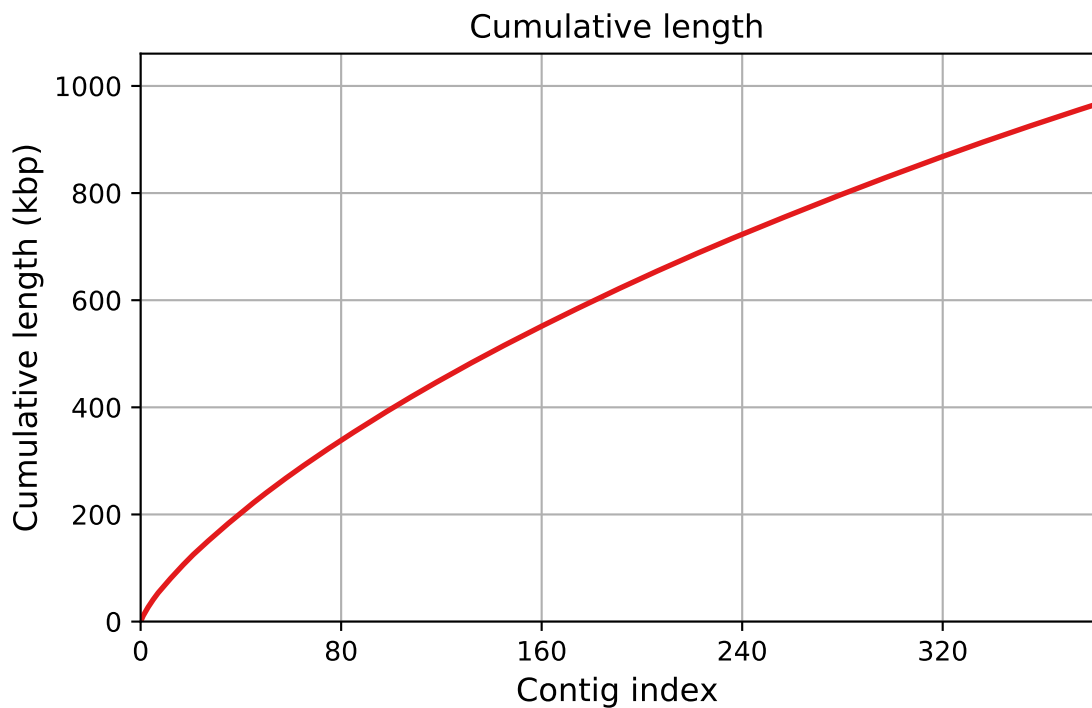
	MEGAHIT-MetaBAT2-3108596.5.fa
# contigs (>= 0 bp)	380
# contigs (>= 1000 bp)	380
# contigs (>= 5000 bp)	15
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	963930
Total length (>= 1000 bp)	963930
Total length (>= 5000 bp)	96187
Total length (>= 10000 bp)	10824
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	380
Largest contig	10824
Total length	963930
GC (%)	56.06
N50	2541
N75	1929
L50	132
L75	240
# 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

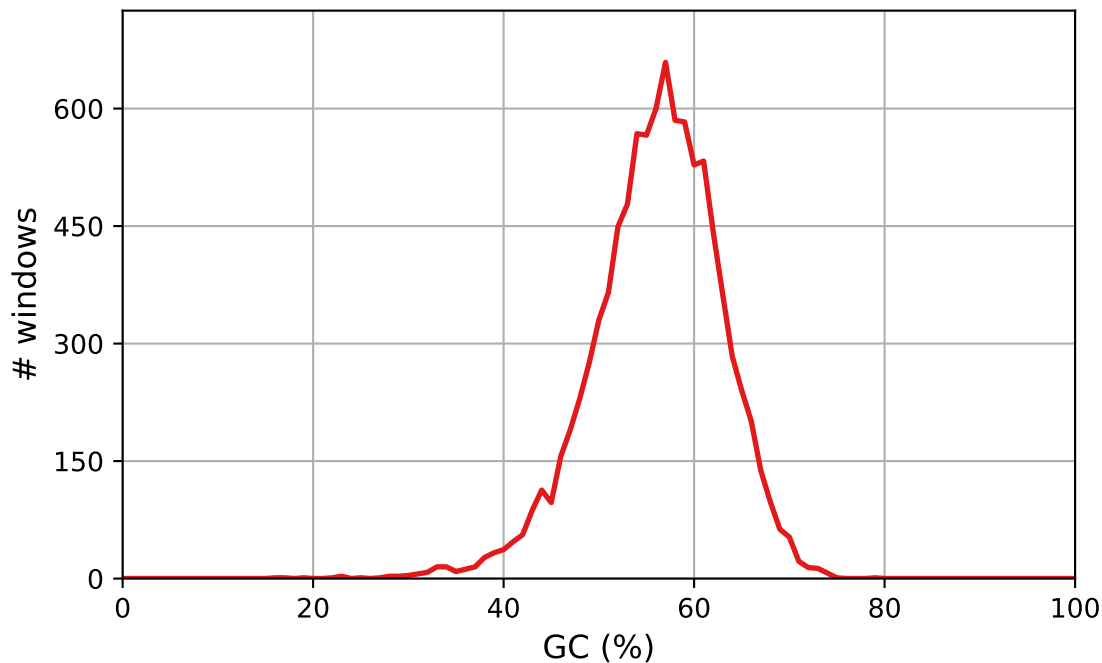


MEGAHIT-MetaBAT2-3108596.5.fa



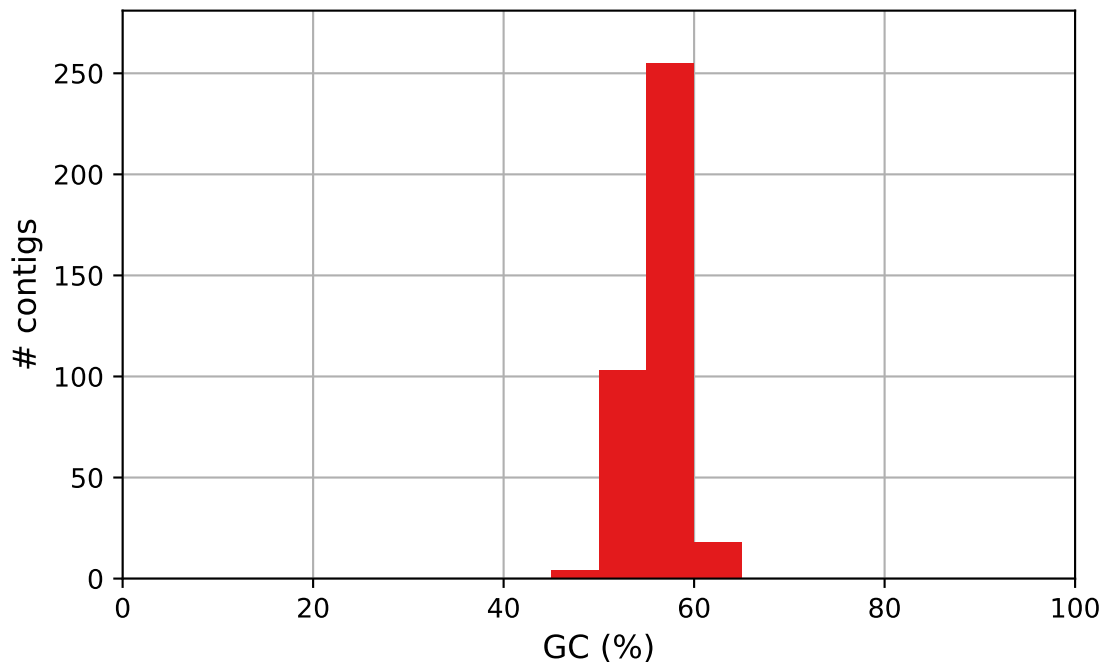
— MEGAHIT-MetaBAT2-3108596.5.fa

GC content



— MEGAHIT-MetaBAT2-3108596.5.fa

MEGAHIT-MetaBAT2-3108596.5.fa GC content



MEGAHIT-MetaBAT2-3108596.5.fa