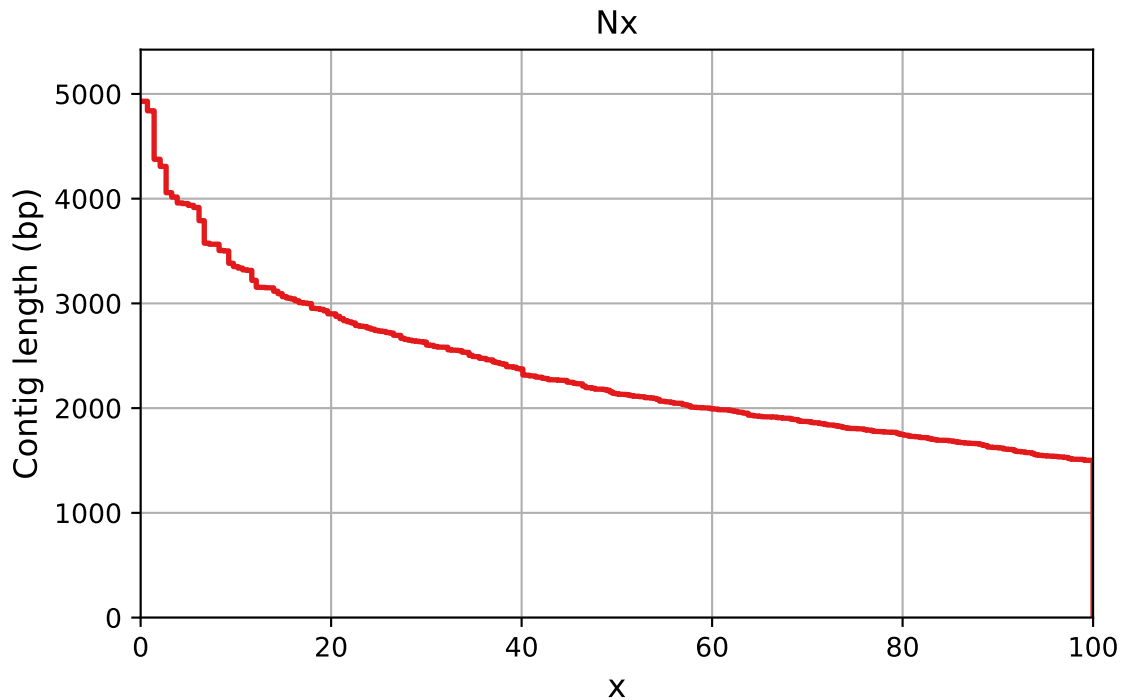


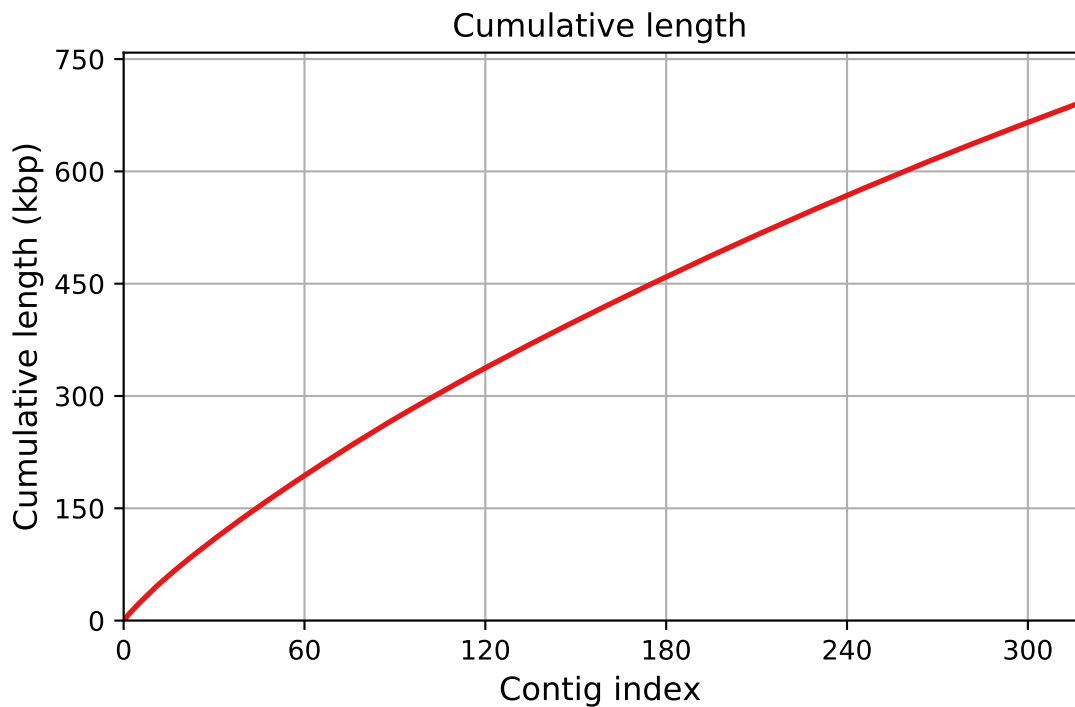
# Report

	MEGAHIT-MetaBAT2-3114327.1.fa
# contigs (>= 0 bp)	316
# contigs (>= 1000 bp)	316
# contigs (>= 5000 bp)	0
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	689632
Total length (>= 1000 bp)	689632
Total length (>= 5000 bp)	0
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	316
Largest contig	4930
Total length	689632
GC (%)	41.68
N50	2137
N75	1805
L50	124
L75	212
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 0 part

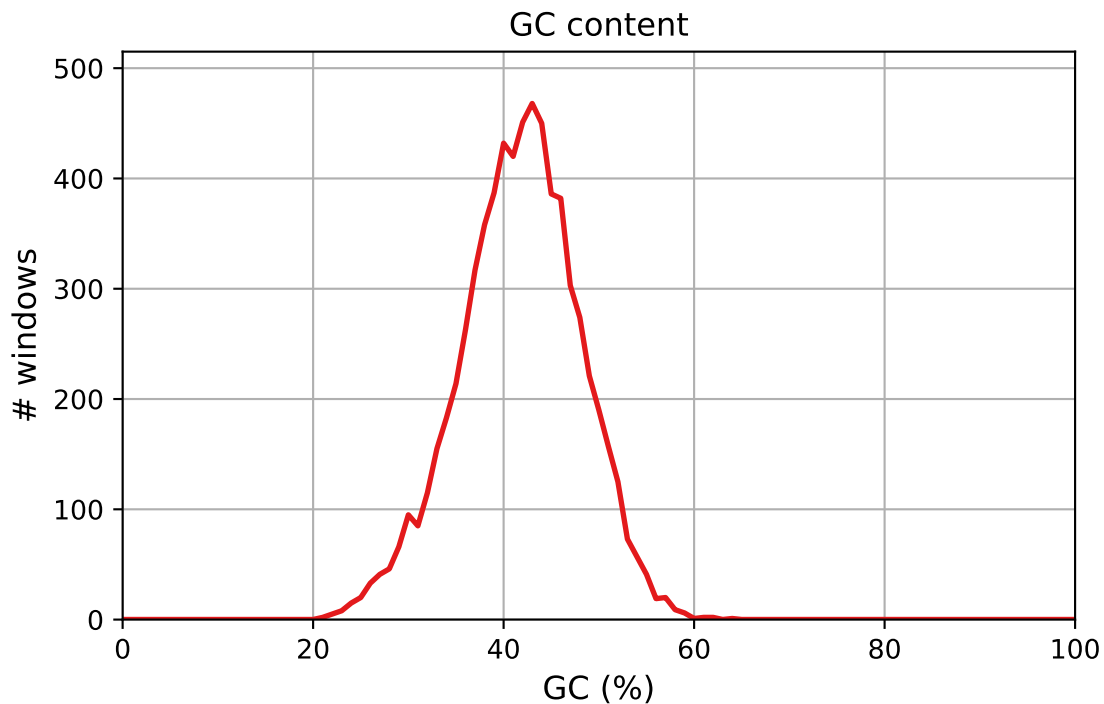
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— MEGAHIT-MetaBAT2-3114327.1.fa

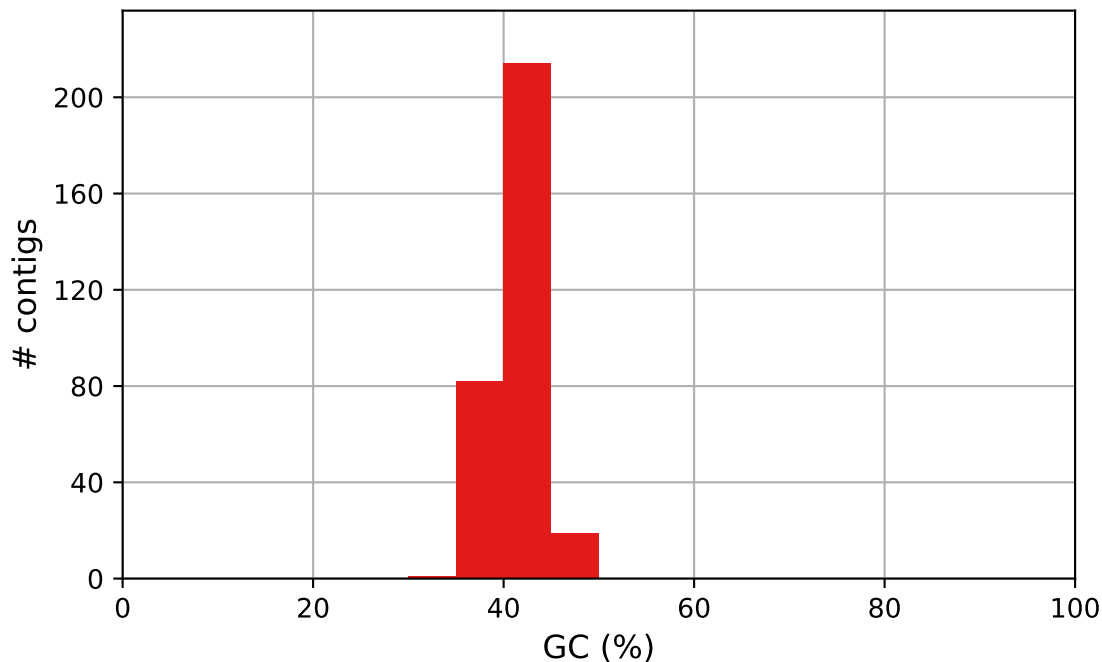


— MEGAHIT-MetaBAT2-3114327.1.fa



— MEGAHit-MetaBAT2-3114327.1.fa

MEGAHIT-MetaBAT2-3114327.1.fa GC content



MEGAHit-MetaBAT2-3114327.1.fa