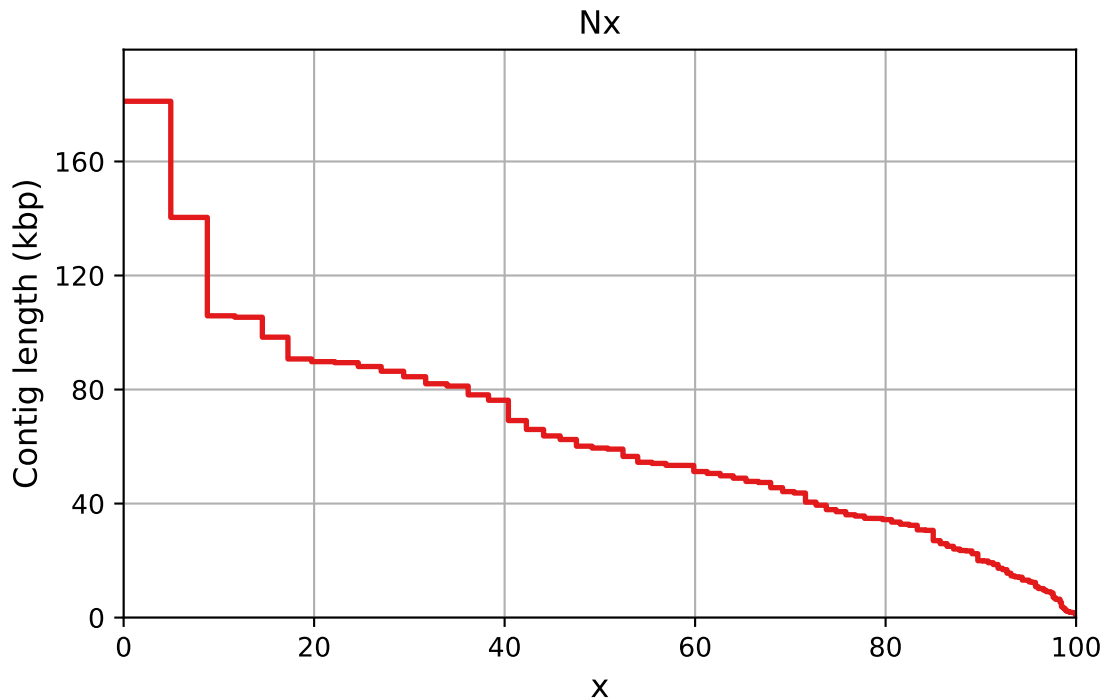


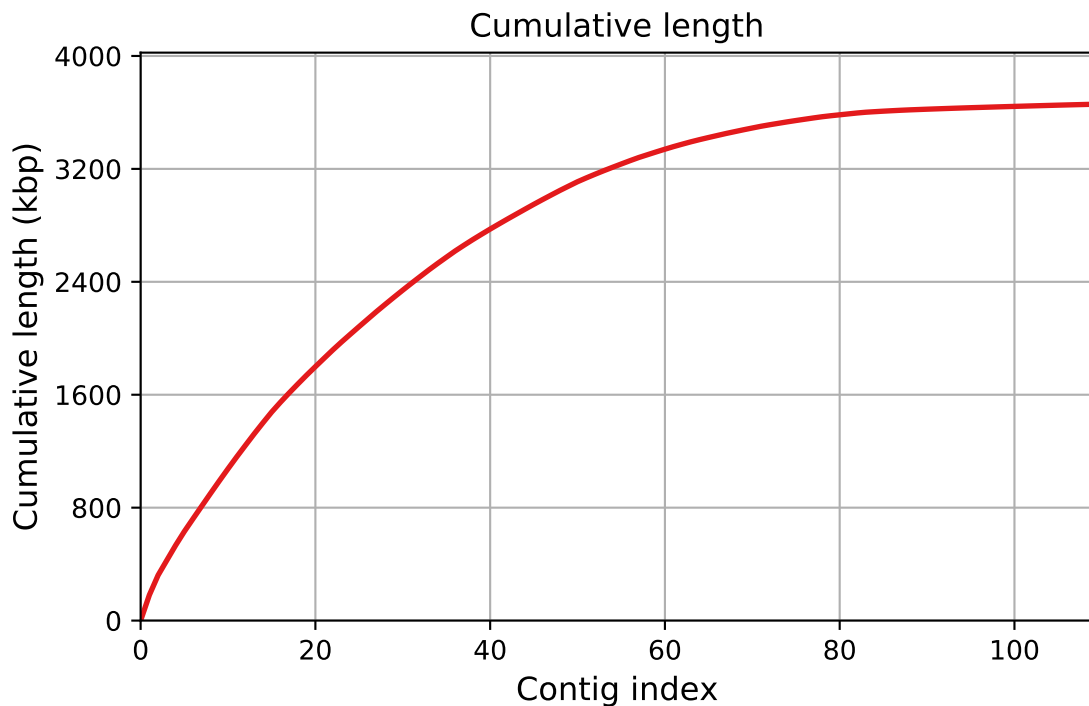
Report

	MEGAHIT-MetaBAT2-3108596.1.fa
# contigs (>= 0 bp)	109
# contigs (>= 1000 bp)	109
# contigs (>= 5000 bp)	83
# contigs (>= 10000 bp)	74
# contigs (>= 25000 bp)	52
# contigs (>= 50000 bp)	29
Total length (>= 0 bp)	3657781
Total length (>= 1000 bp)	3657781
Total length (>= 5000 bp)	3601422
Total length (>= 10000 bp)	3533148
Total length (>= 25000 bp)	3162048
Total length (>= 50000 bp)	2291275
# contigs	109
Largest contig	181120
Total length	3657781
GC (%)	41.16
N50	59469
N75	37157
L50	21
L75	40
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 1 part

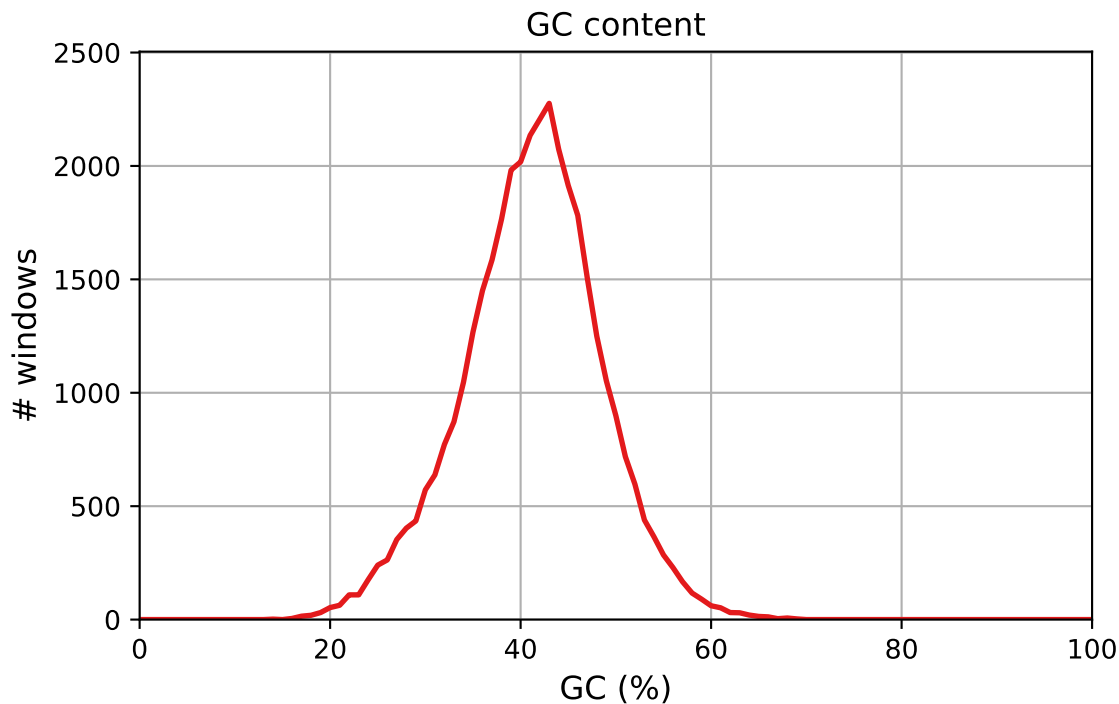
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— MEGAHIT-MetaBAT2-3108596.1.fa

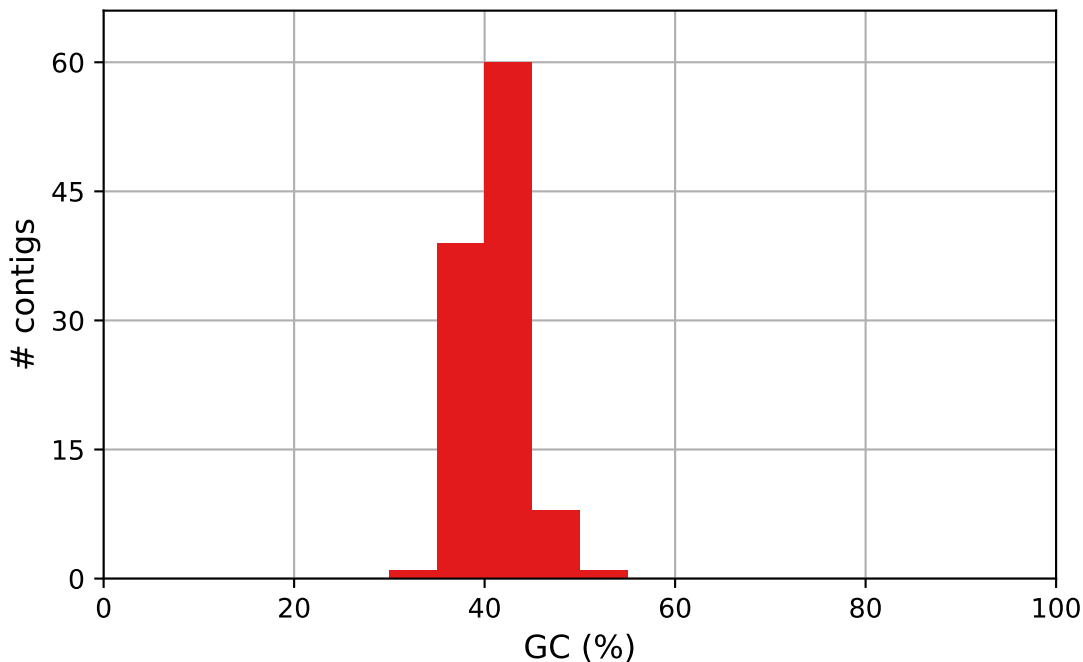


— MEGAHIT-MetaBAT2-3108596.1.fa



— MEGAHit-MetaBAT2-3108596.1.fa

MEGAHIT-MetaBAT2-3108596.1.fa GC content



MEGAHit-MetaBAT2-3108596.1.fa