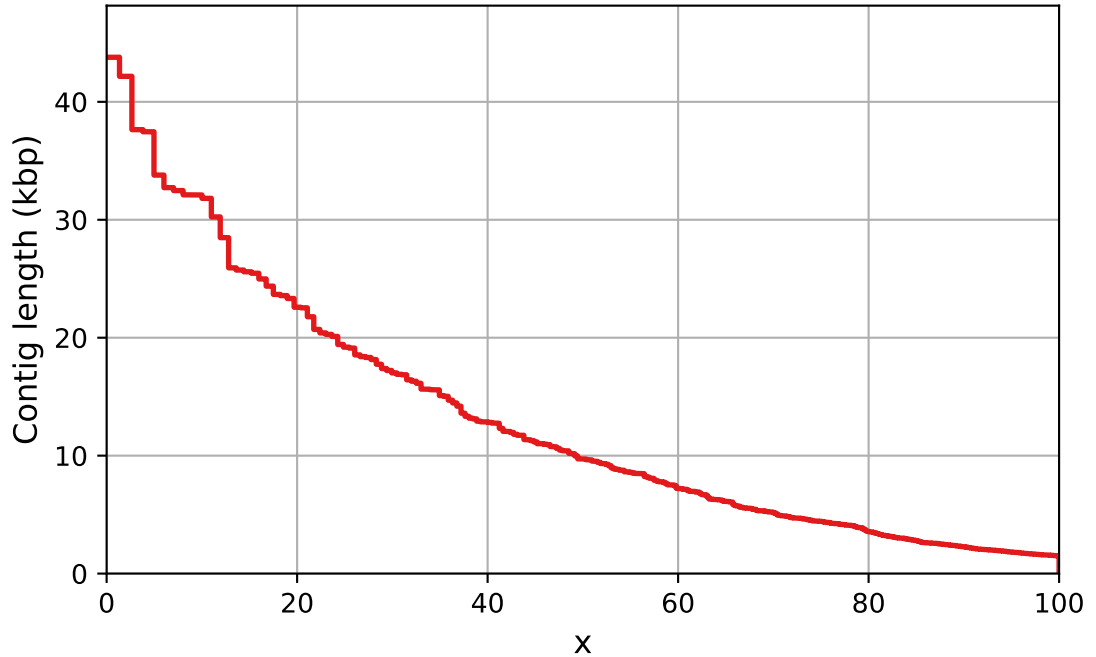


Report

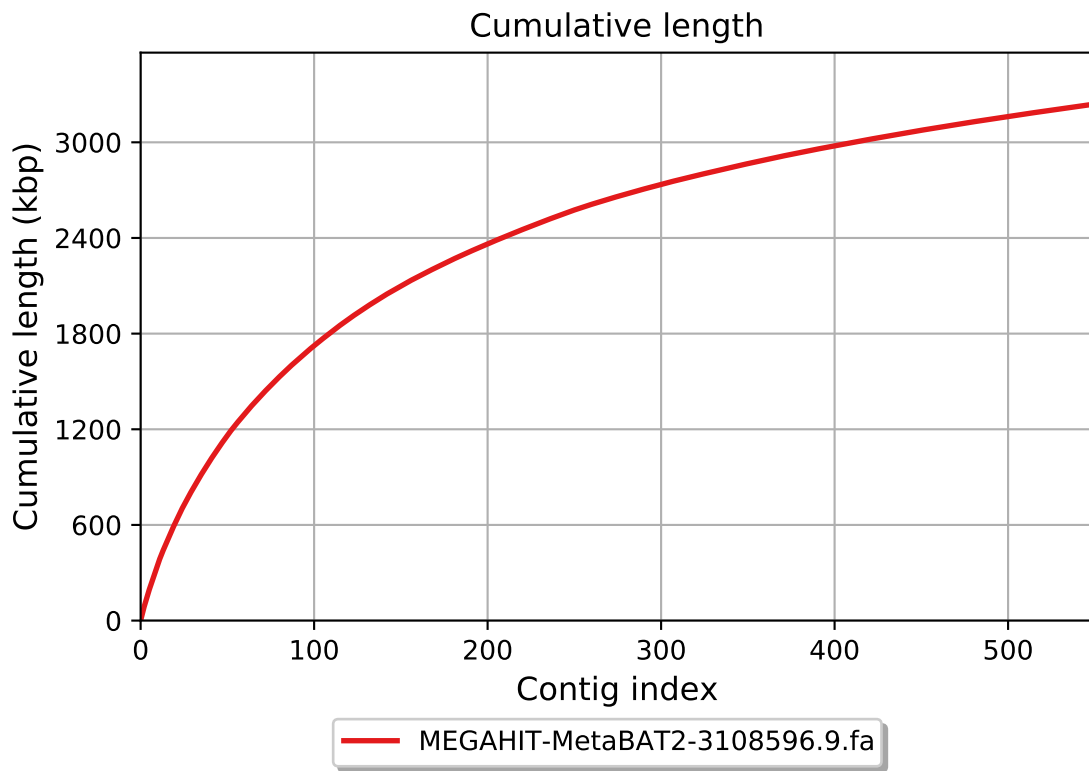
	MEGAHIT-MetaBAT2-3108596.9.fa
# contigs (>= 0 bp)	549
# contigs (>= 1000 bp)	549
# contigs (>= 5000 bp)	183
# contigs (>= 10000 bp)	86
# contigs (>= 25000 bp)	16
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3238623
Total length (>= 1000 bp)	3238623
Total length (>= 5000 bp)	2281071
Total length (>= 10000 bp)	1591658
Total length (>= 25000 bp)	517518
Total length (>= 50000 bp)	0
# contigs	549
Largest contig	43779
Total length	3238623
GC (%)	42.68
N50	9718
N75	4423
L50	89
L75	215
# 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).

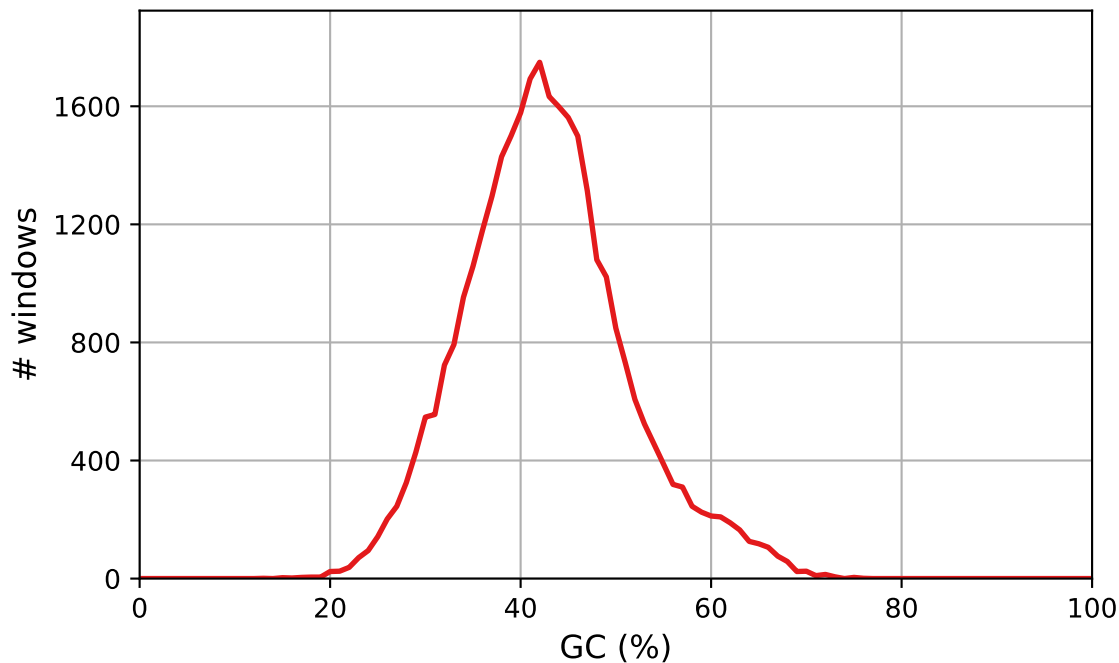
Nx



MEGAHIT-MetaBAT2-3108596.9.fa

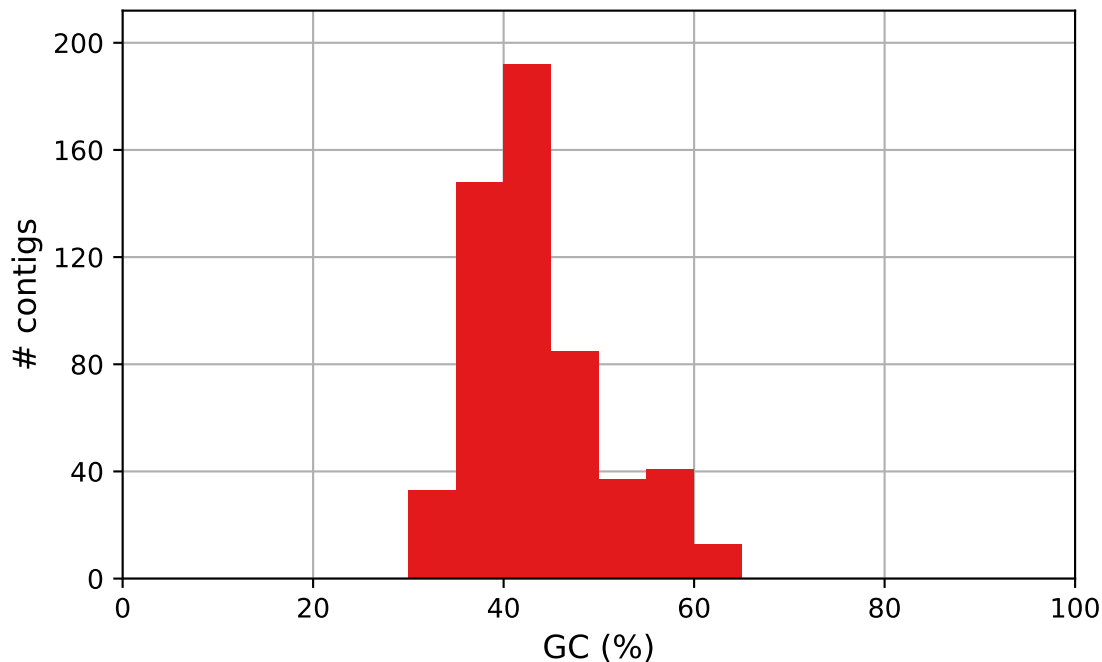


GC content



MEGAHIT-MetaBAT2-3108596.9.fa

MEGAHIT-MetaBAT2-3108596.9.fa GC content



MEGAHIT-MetaBAT2-3108596.9.fa