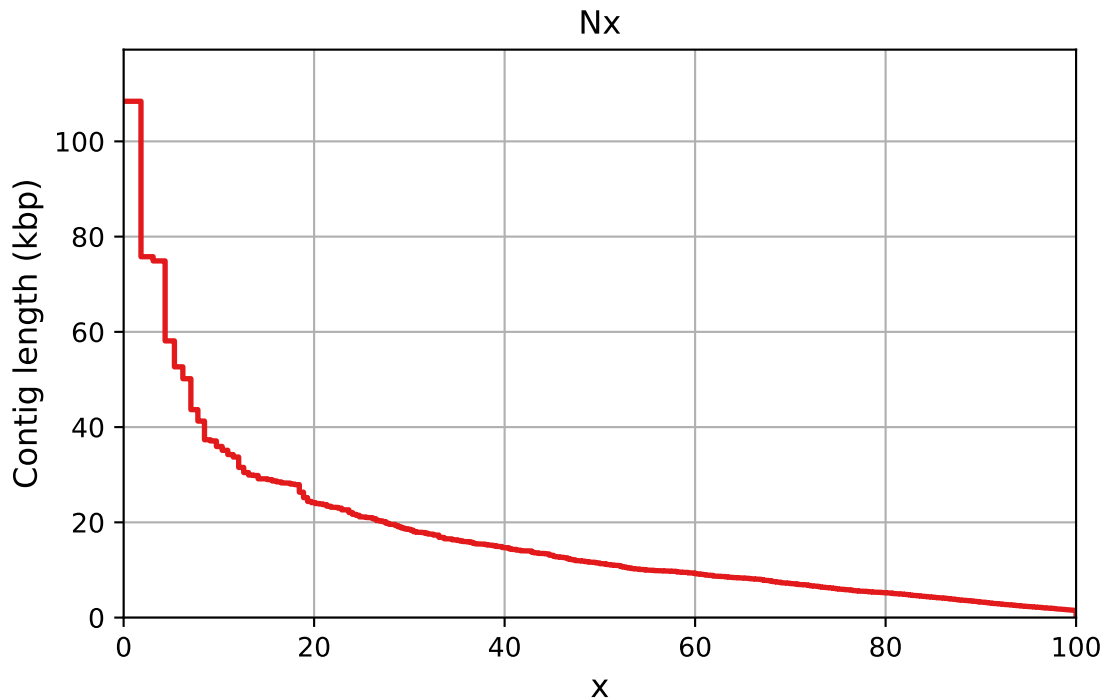


Report

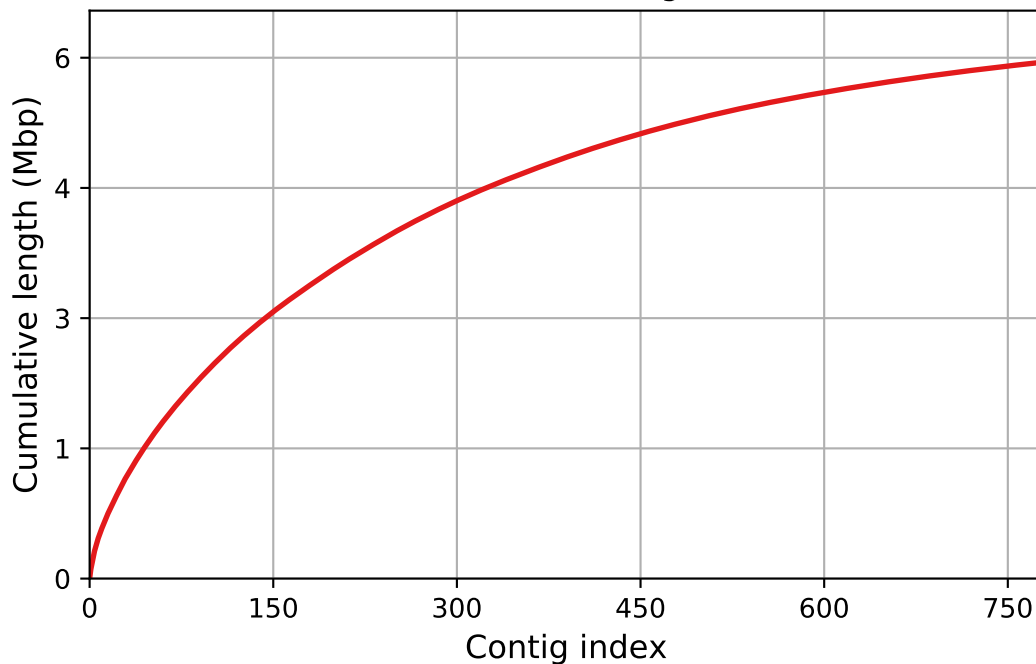
	MEGAHIT-MetaBAT2-3106237.8.fa
# contigs (>= 0 bp)	778
# contigs (>= 1000 bp)	778
# contigs (>= 5000 bp)	387
# contigs (>= 10000 bp)	168
# contigs (>= 25000 bp)	29
# contigs (>= 50000 bp)	6
Total length (>= 0 bp)	5947729
Total length (>= 1000 bp)	5947729
Total length (>= 5000 bp)	4841423
Total length (>= 10000 bp)	3261420
Total length (>= 25000 bp)	1148589
Total length (>= 50000 bp)	420025
# contigs	778
Largest contig	108422
Total length	5947729
GC (%)	42.16
N50	11327
N75	6007
L50	142
L75	318
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 2 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).



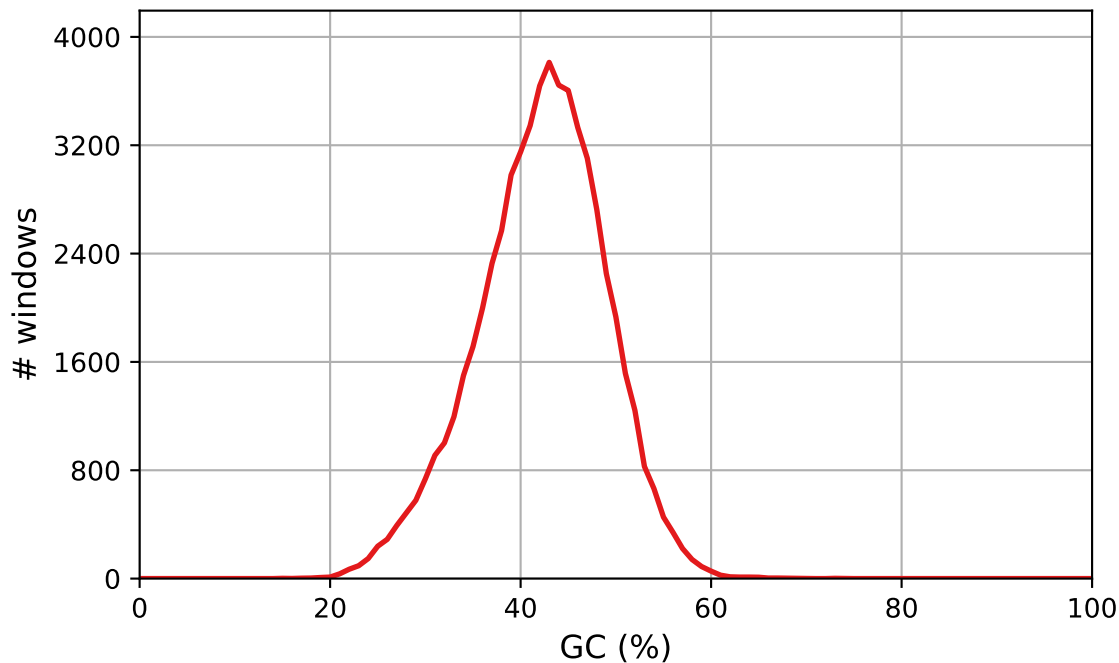
— MEGAHIT-MetaBAT2-3106237.8.fa

Cumulative length



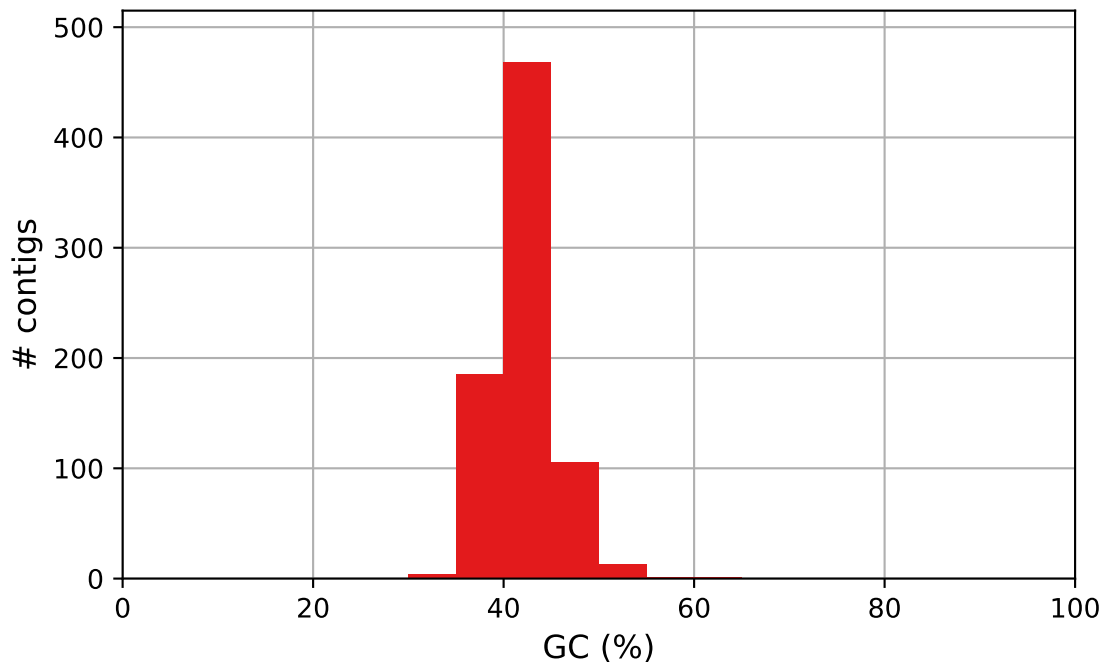
— MEGAHIT-MetaBAT2-3106237.8.fa

GC content



— MEGAHIT-MetaBAT2-3106237.8.fa

MEGAHIT-MetaBAT2-3106237.8.fa GC content



MEGAHit-MetaBAT2-3106237.8.fa