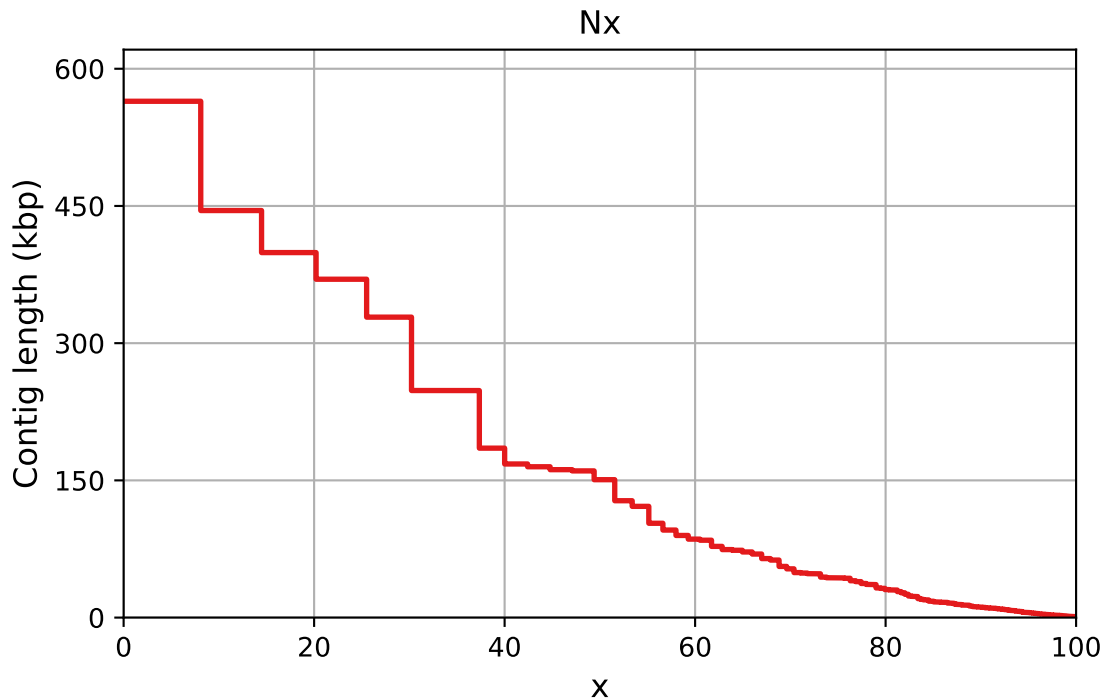


Report

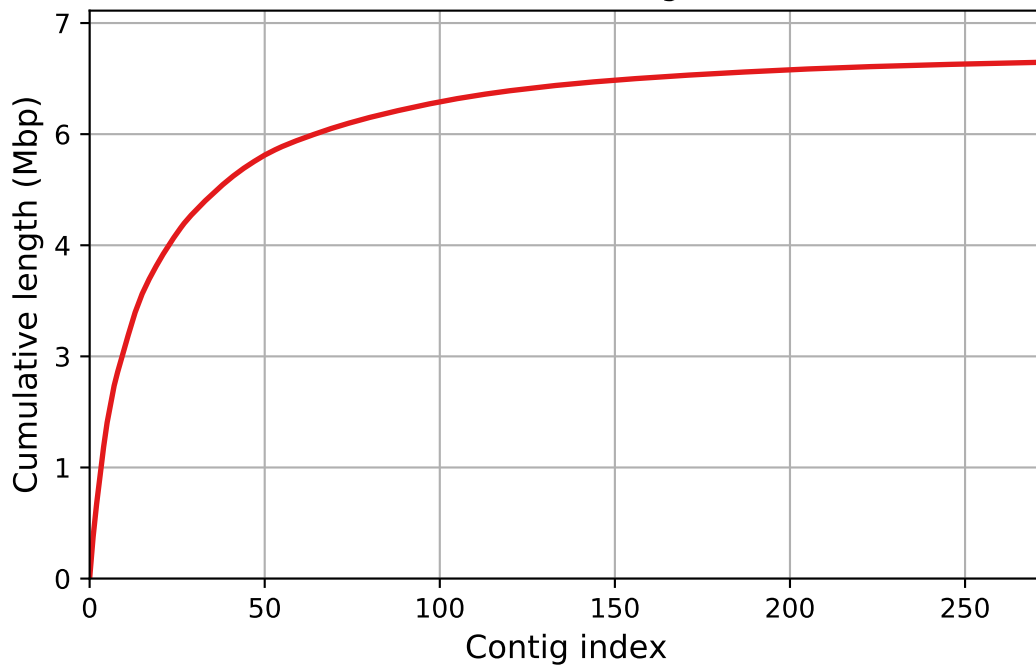
	MEGAHIT-MaxBin2-3106384.002.fa
# contigs (>= 0 bp)	272
# contigs (>= 1000 bp)	272
# contigs (>= 5000 bp)	133
# contigs (>= 10000 bp)	95
# contigs (>= 25000 bp)	51
# contigs (>= 50000 bp)	29
Total length (>= 0 bp)	6970471
Total length (>= 1000 bp)	6970471
Total length (>= 5000 bp)	6656749
Total length (>= 10000 bp)	6389294
Total length (>= 25000 bp)	5742848
Total length (>= 50000 bp)	4905959
# contigs	272
Largest contig	564450
Total length	6970471
GC (%)	45.18
N50	150751
N75	43480
L50	13
L75	36
# N's per 100 kbp	0.00
# predicted rRNA genes	4 + 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).



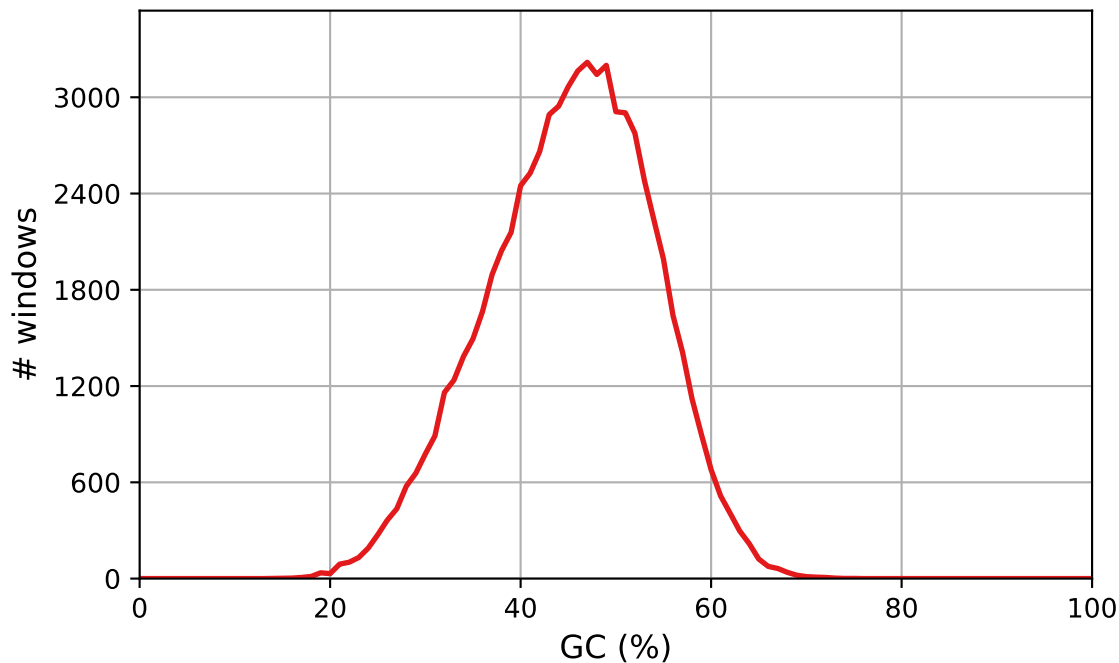
MEGAHIT-MaxBin2-3106384.002.fa

Cumulative length



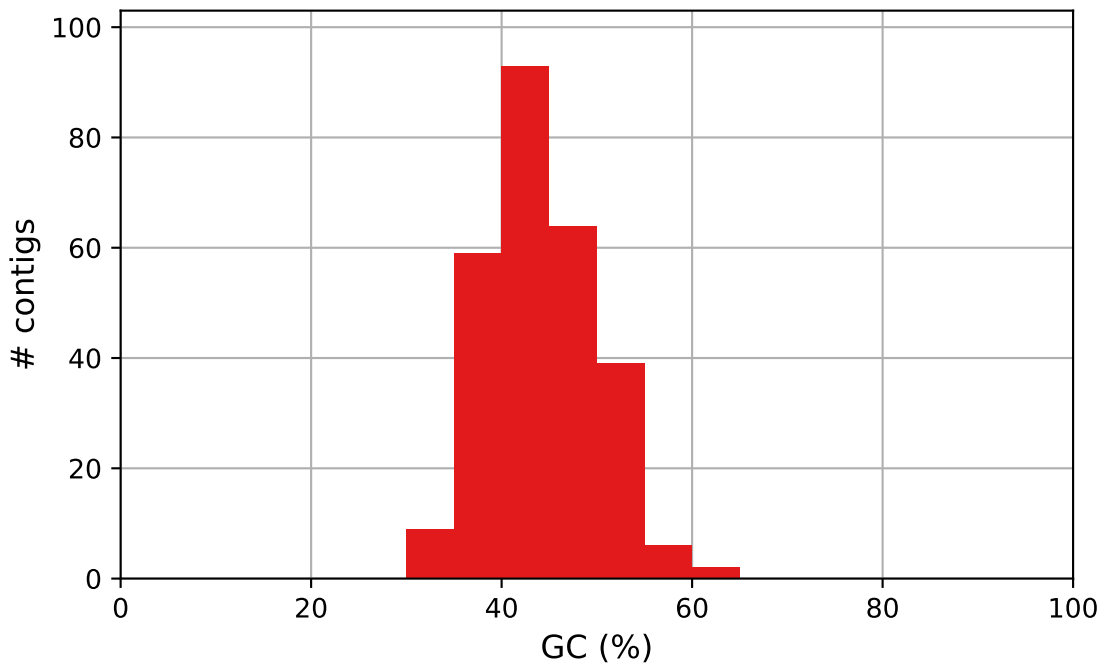
MEGAHIT-MaxBin2-3106384.002.fa

GC content



MEGAHIT-MaxBin2-3106384.002.fa

MEGAHIT-MaxBin2-3106384.002.fa GC content



MEGAHit-MaxBin2-3106384.002.fa