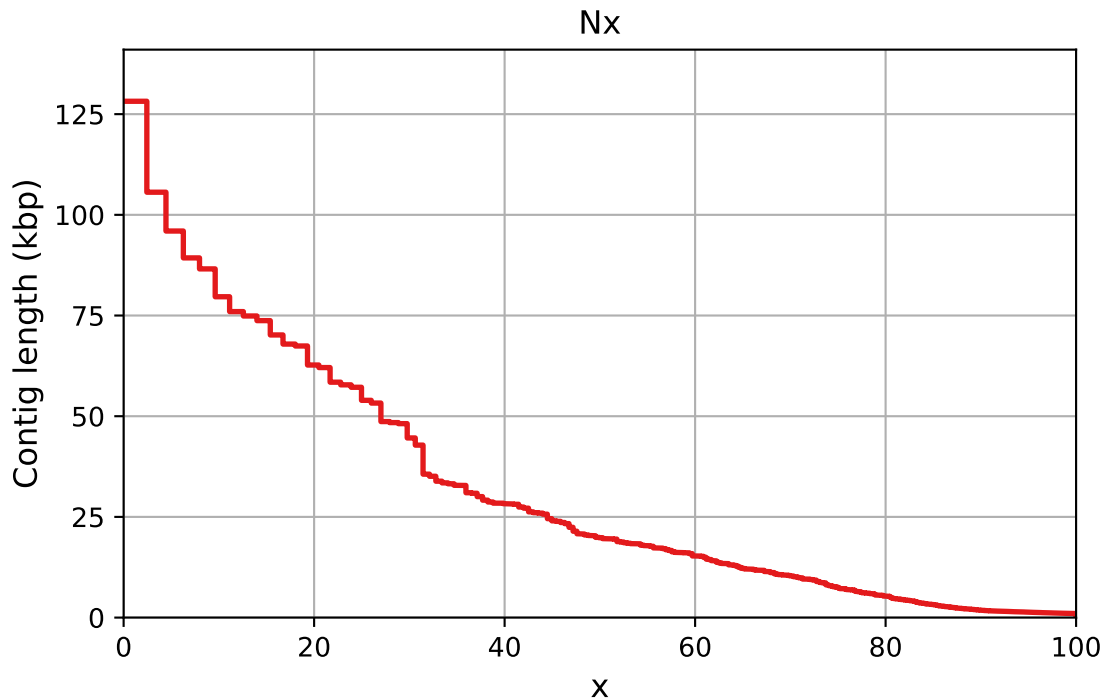


# Report

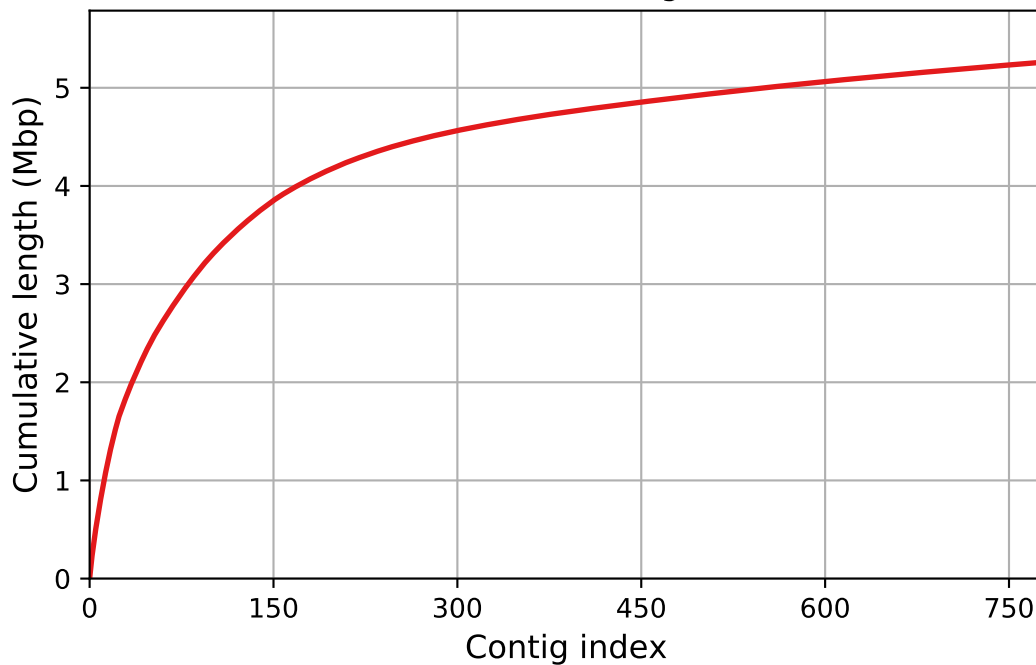
	MEGAHIT-MaxBin2-3101140.005.fa
# contigs (>= 0 bp)	777
# contigs (>= 1000 bp)	777
# contigs (>= 5000 bp)	209
# contigs (>= 10000 bp)	136
# contigs (>= 25000 bp)	47
# contigs (>= 50000 bp)	19
Total length (>= 0 bp)	5260026
Total length (>= 1000 bp)	5260026
Total length (>= 5000 bp)	4236777
Total length (>= 10000 bp)	3719922
Total length (>= 25000 bp)	2340140
Total length (>= 50000 bp)	1420870
# contigs	777
Largest contig	128197
Total length	5260026
GC (%)	50.83
N50	19834
N75	7491
L50	61
L75	162
# N's per 100 kbp	0.00
# predicted rRNA genes	4 + 9 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).



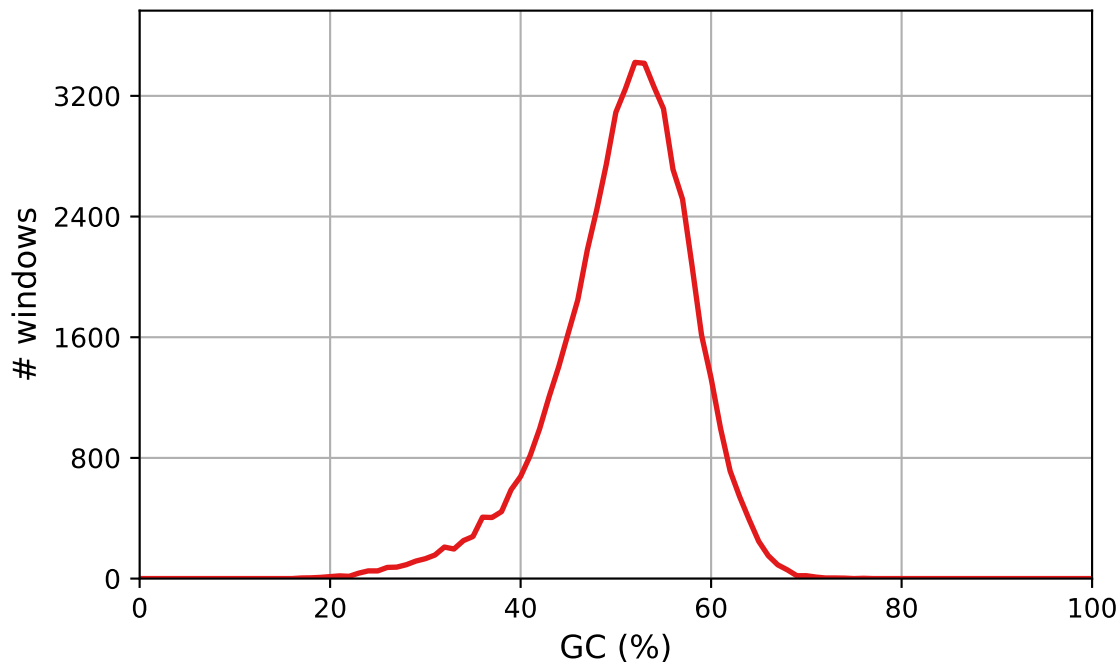
MEGAHIT-MaxBin2-3101140.005.fa

Cumulative length



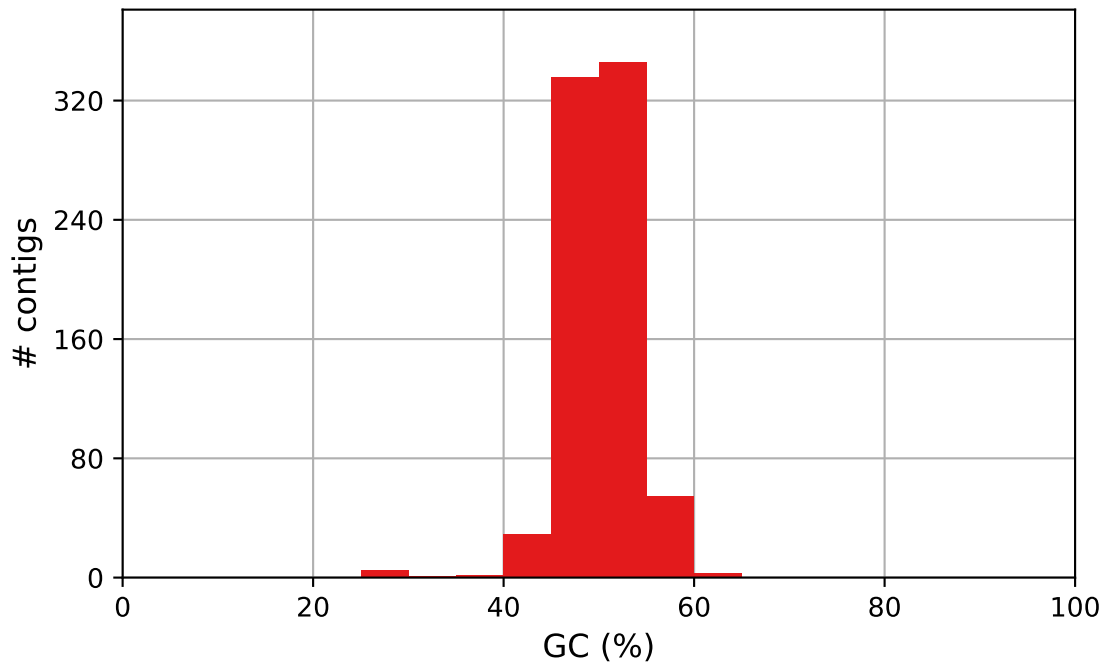
MEGAHIT-MaxBin2-3101140.005.fa

# GC content



MEGAHit-MaxBin2-3101140.005.fa

MEGAHIT-MaxBin2-3101140.005.fa GC content



MEGAHit-MaxBin2-3101140.005.fa