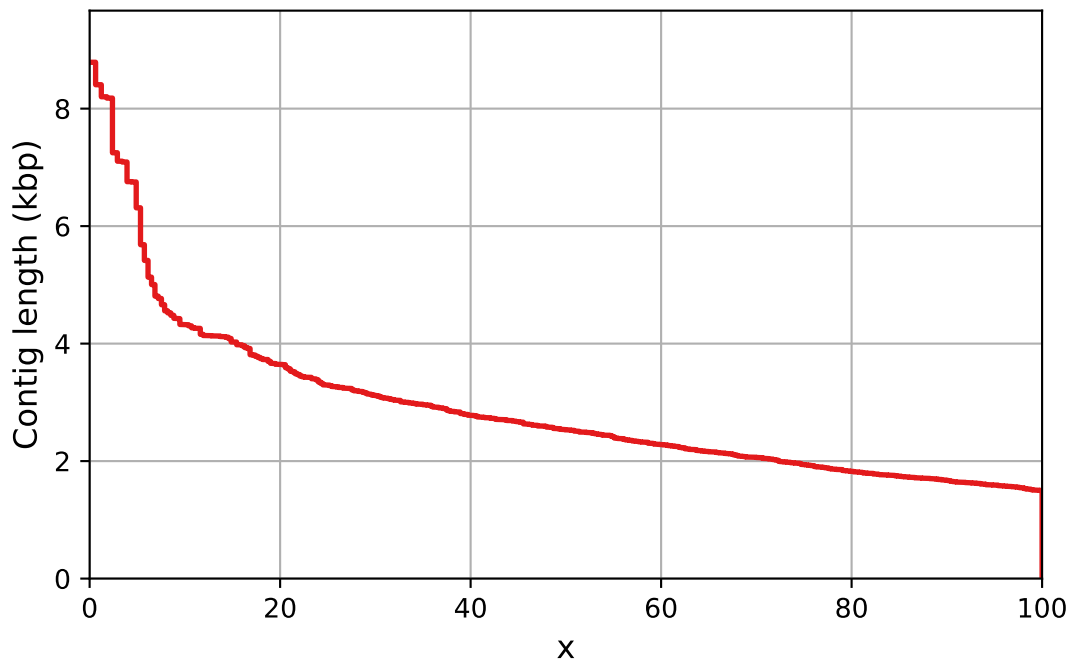


Report

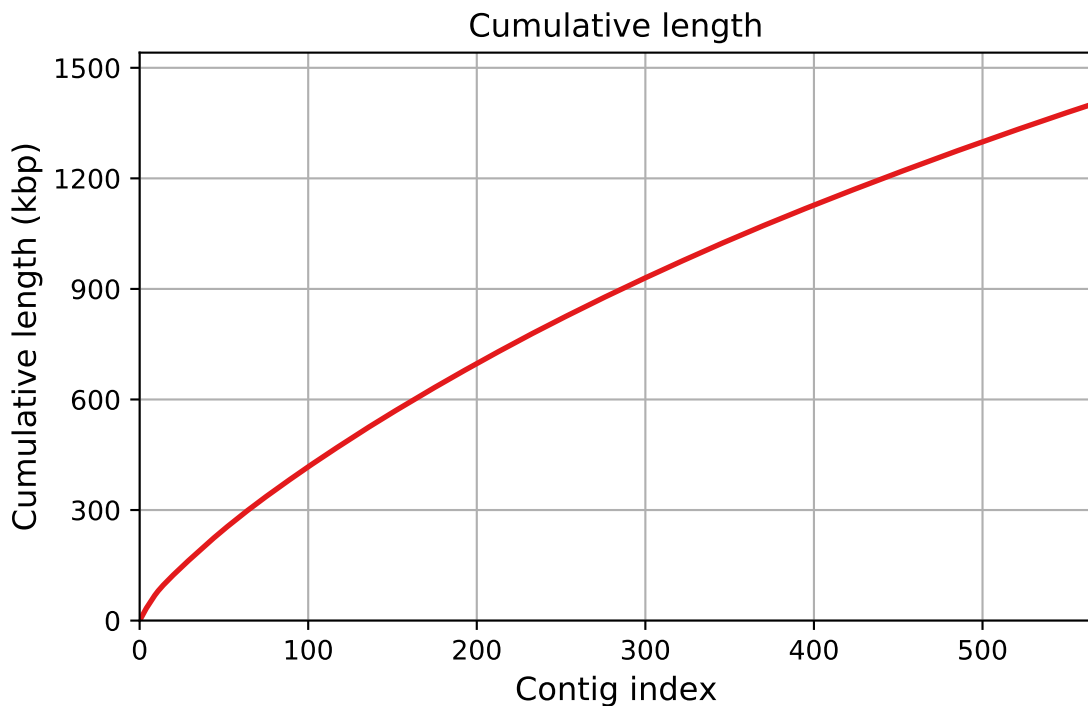
	MEGAHIT-MetaBAT2-3101140.5.fa
# contigs (>= 0 bp)	565
# contigs (>= 1000 bp)	565
# contigs (>= 5000 bp)	14
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1401059
Total length (>= 1000 bp)	1401059
Total length (>= 5000 bp)	96078
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	565
Largest contig	8789
Total length	1401059
GC (%)	51.63
N50	2533
N75	1939
L50	202
L75	360
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 0 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

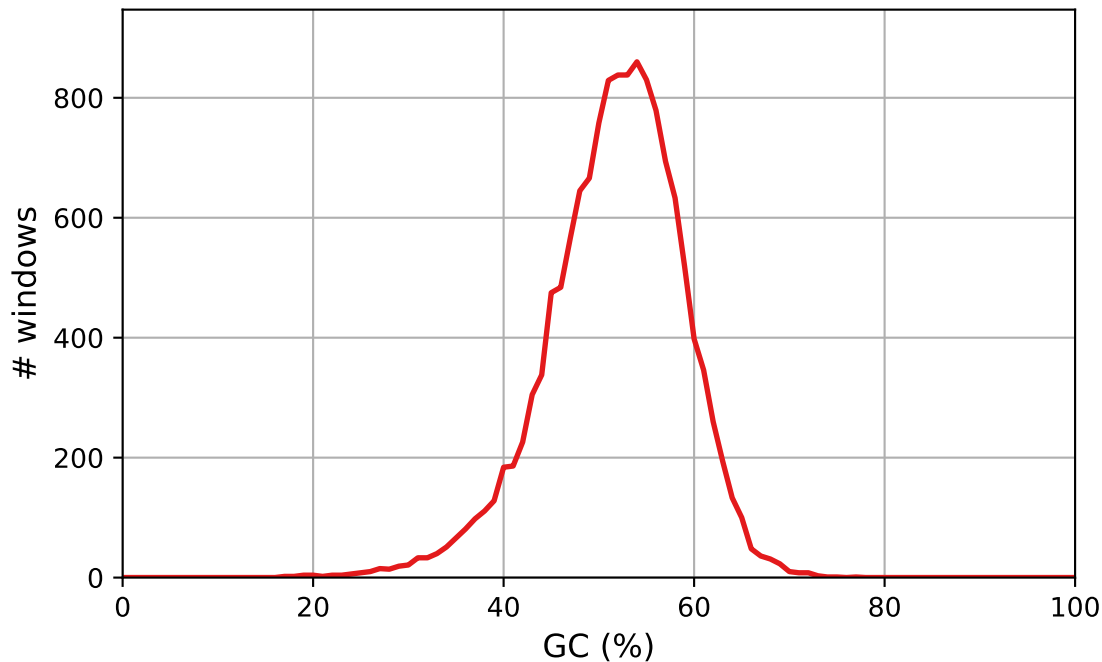


— MEGAHIT-MetaBAT2-3101140.5.fa



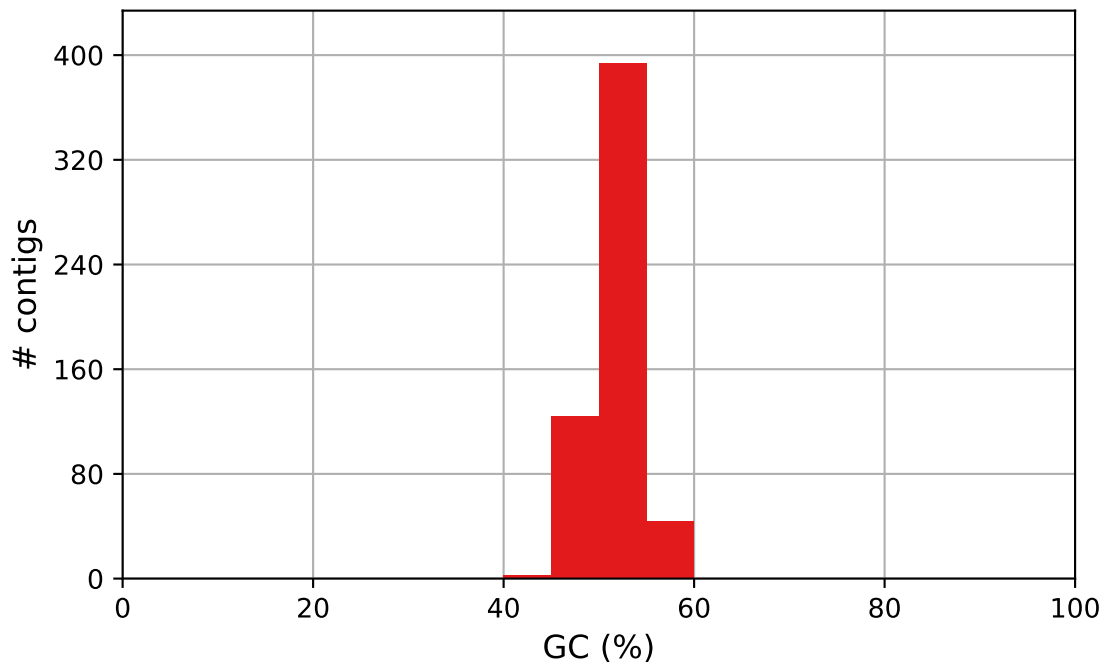
— MEGAHIT-MetaBAT2-3101140.5.fa

GC content



— MEGAHIT-MetaBAT2-3101140.5.fa

MEGAHIT-MetaBAT2-3101140.5.fa GC content



MEGAHit-MetaBAT2-3101140.5.fa