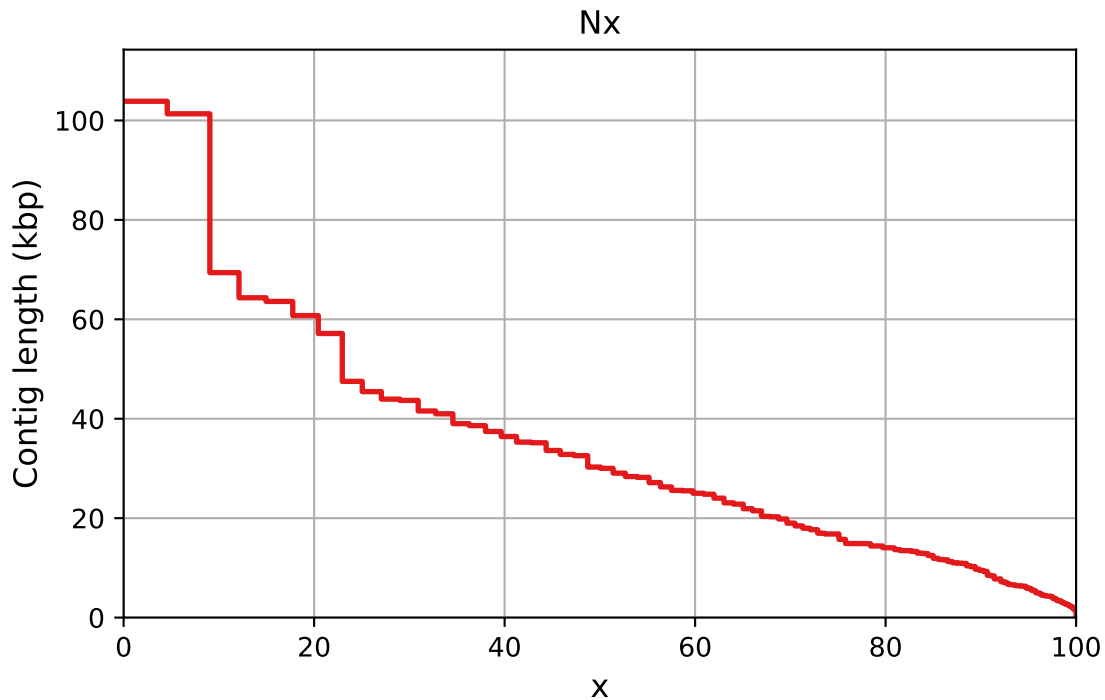


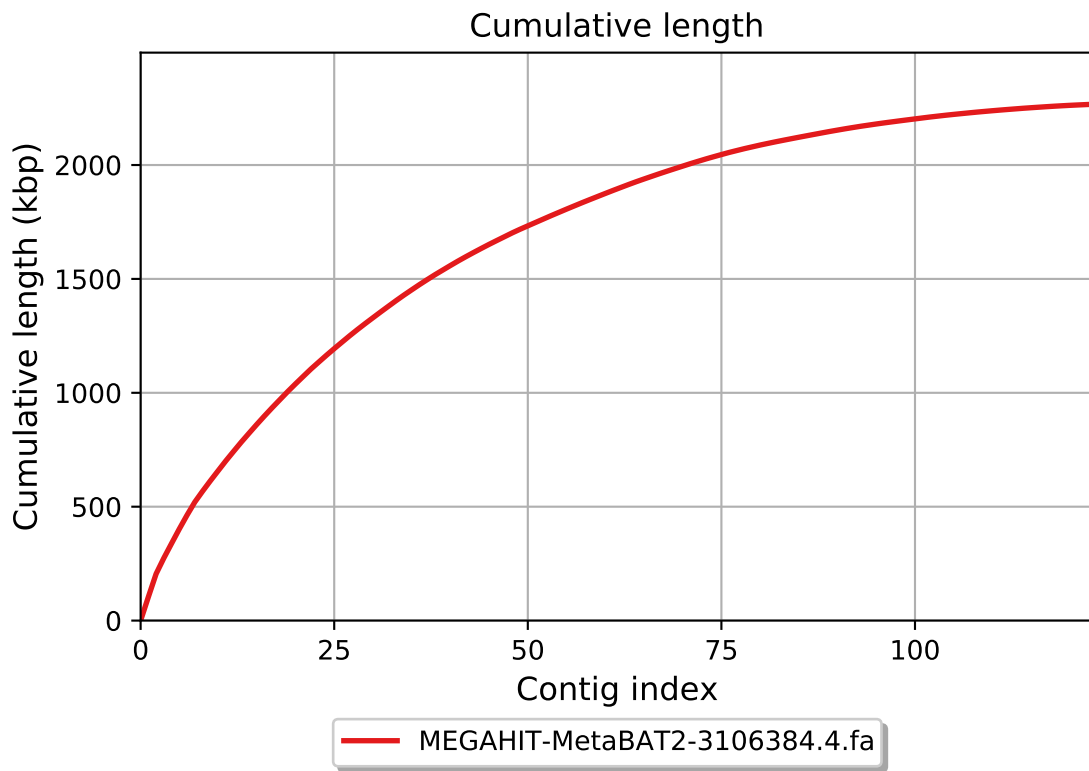
# Report

	MEGAHIT-MetaBAT2-3106384.4.fa
# contigs (>= 0 bp)	123
# contigs (>= 1000 bp)	123
# contigs (>= 5000 bp)	93
# contigs (>= 10000 bp)	73
# contigs (>= 25000 bp)	32
# contigs (>= 50000 bp)	7
Total length (>= 0 bp)	2266704
Total length (>= 1000 bp)	2266704
Total length (>= 5000 bp)	2170159
Total length (>= 10000 bp)	2026664
Total length (>= 25000 bp)	1379995
Total length (>= 50000 bp)	520412
# contigs	123
Largest contig	103859
Total length	2266704
GC (%)	59.94
N50	30277
N75	16817
L50	23
L75	48
# N's per 100 kbp	0.00
# predicted rRNA genes	3 + 0 part

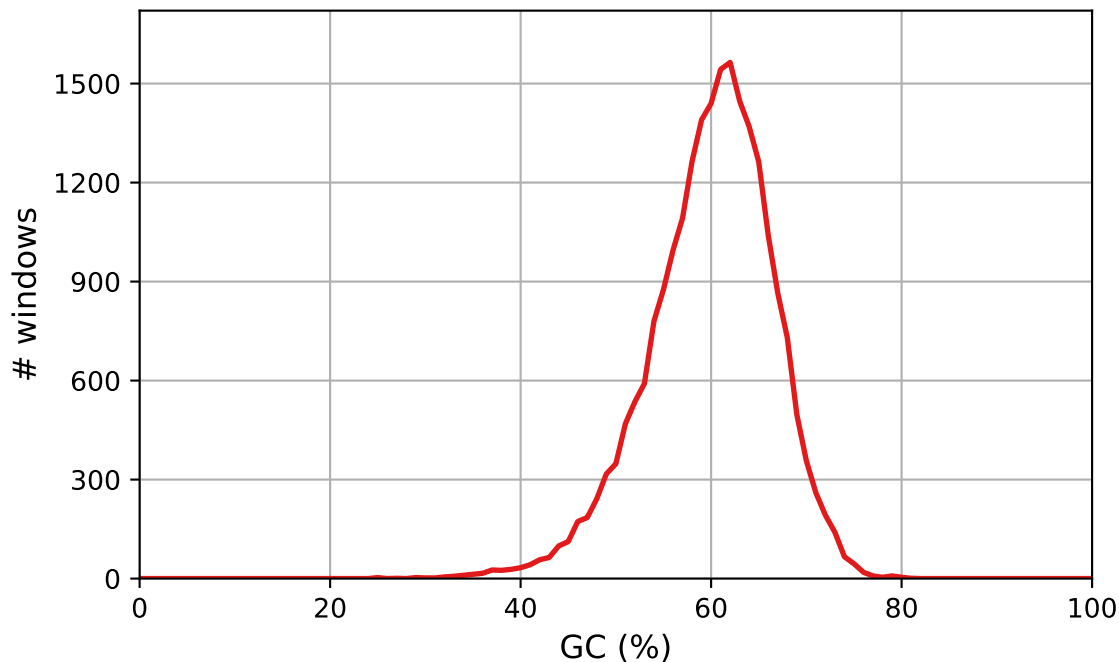
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— MEGAHIT-MetaBAT2-3106384.4.fa

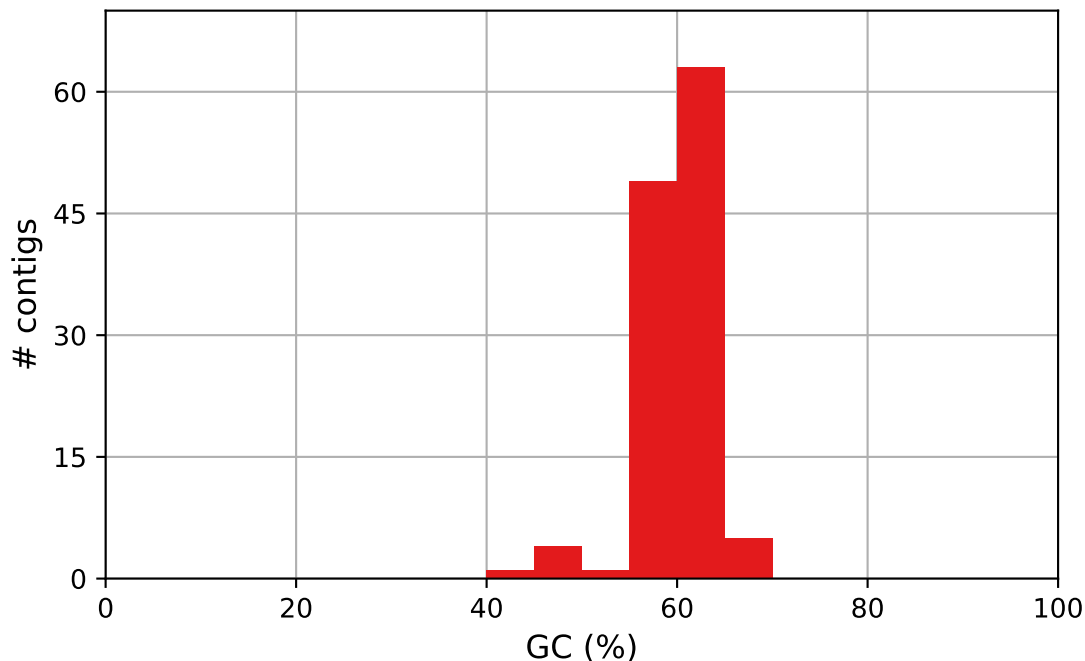


## GC content



— MEGAHIT-MetaBAT2-3106384.4.fa

MEGAHIT-MetaBAT2-3106384.4.fa GC content



MEGAhit-MetaBAT2-3106384.4.fa