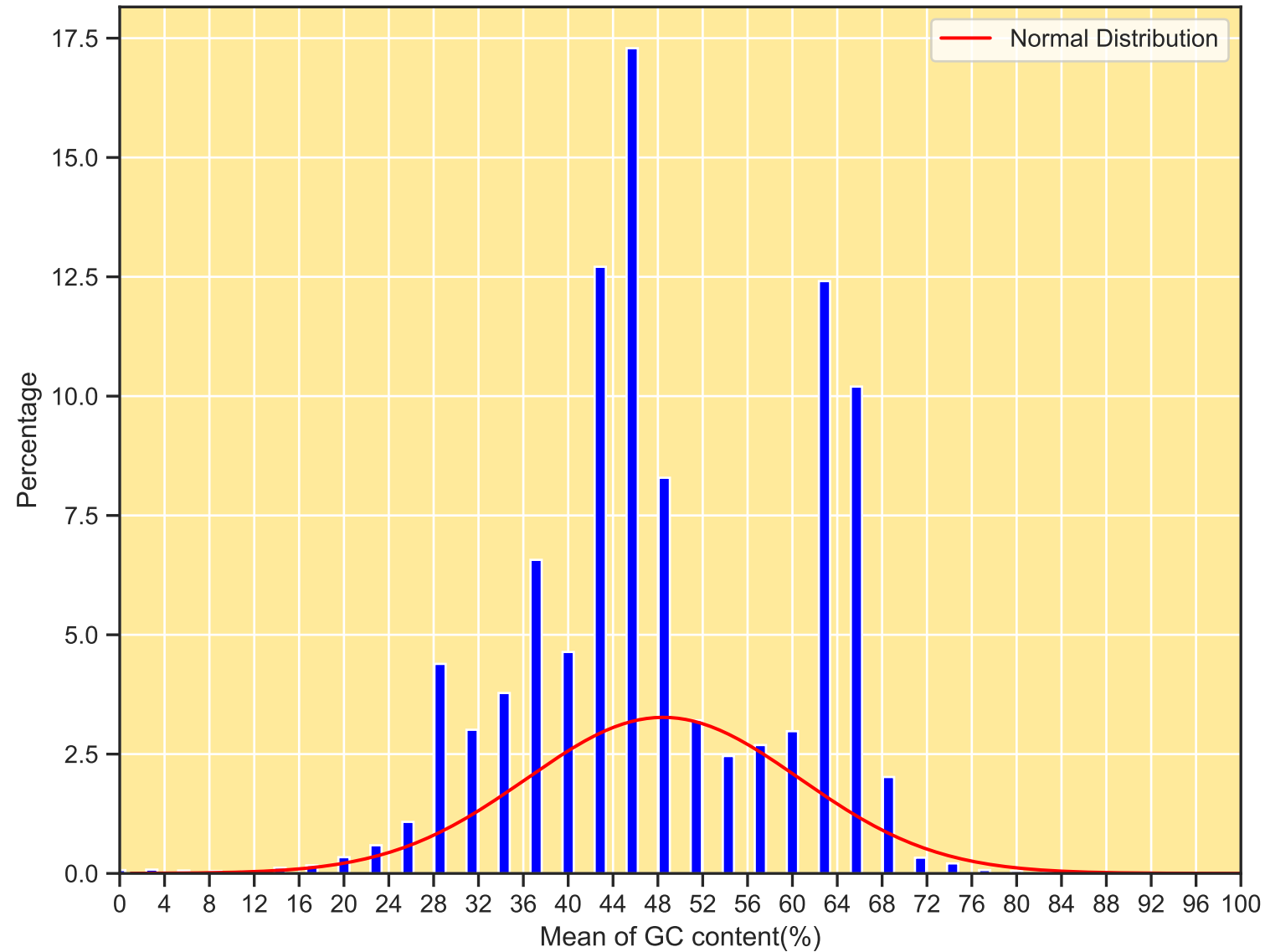


GC distribution over all sequences (file.fastq.gz)



GC Q-Q Plot (file.fastq.gz)

