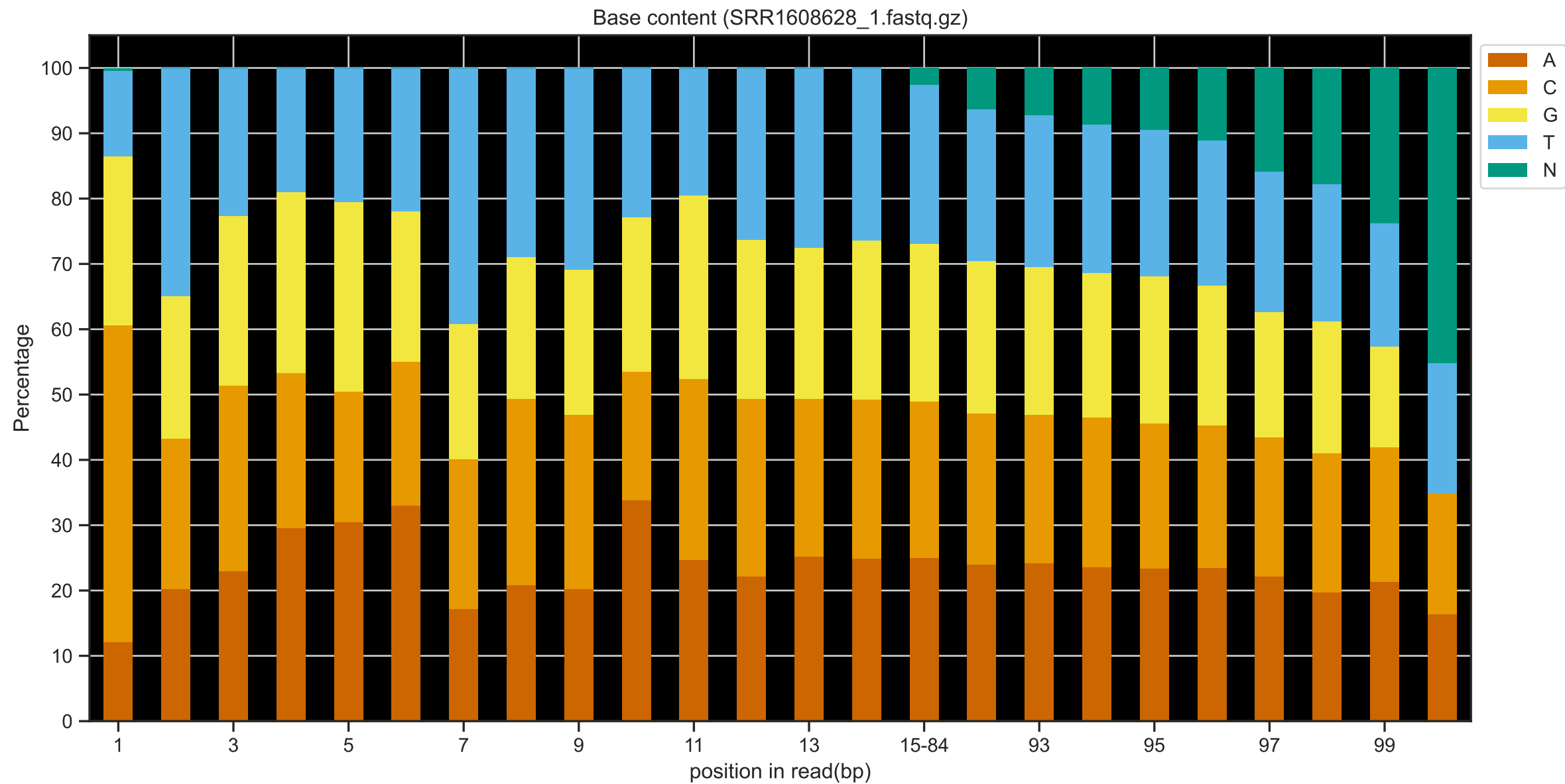
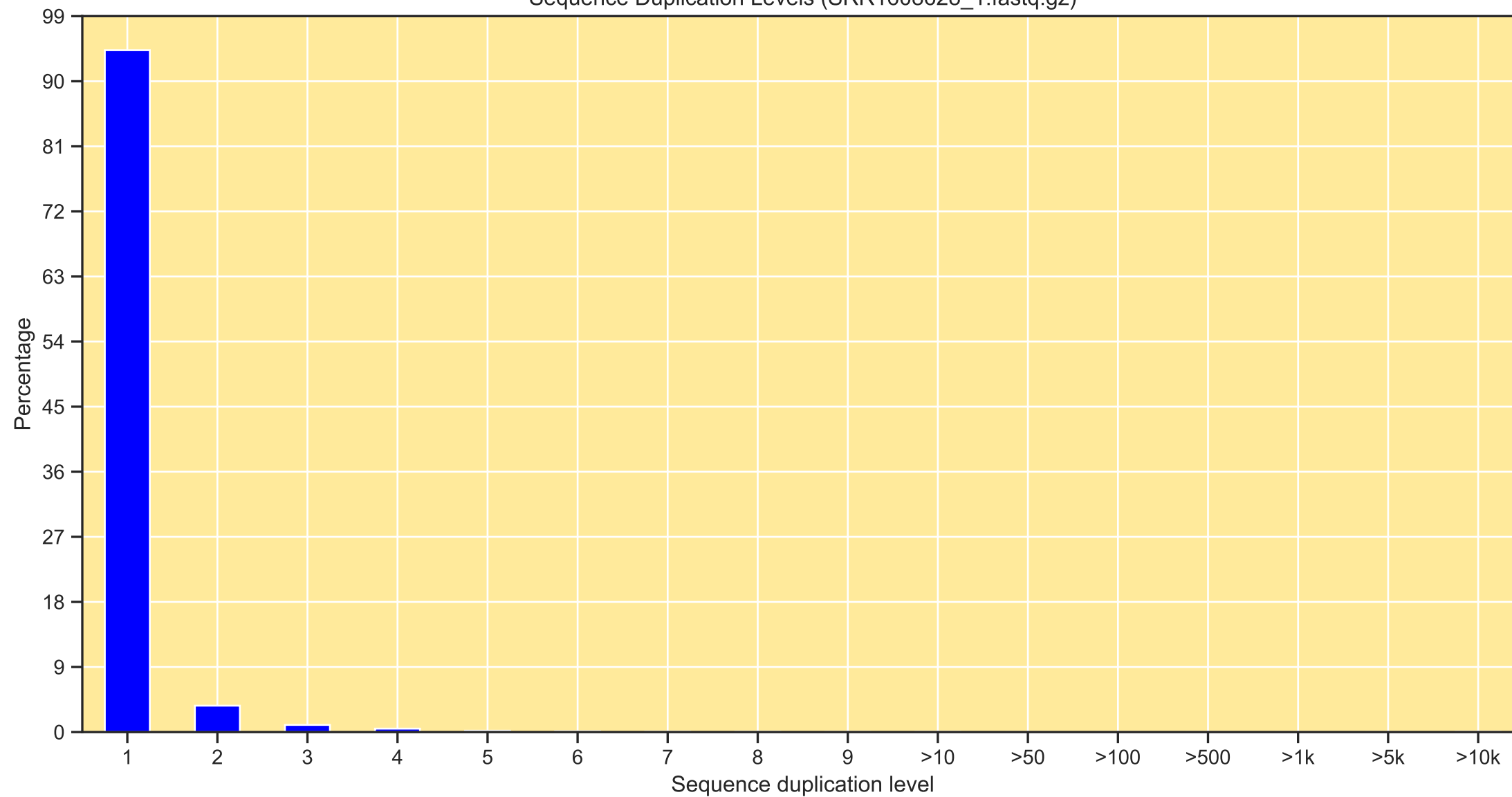


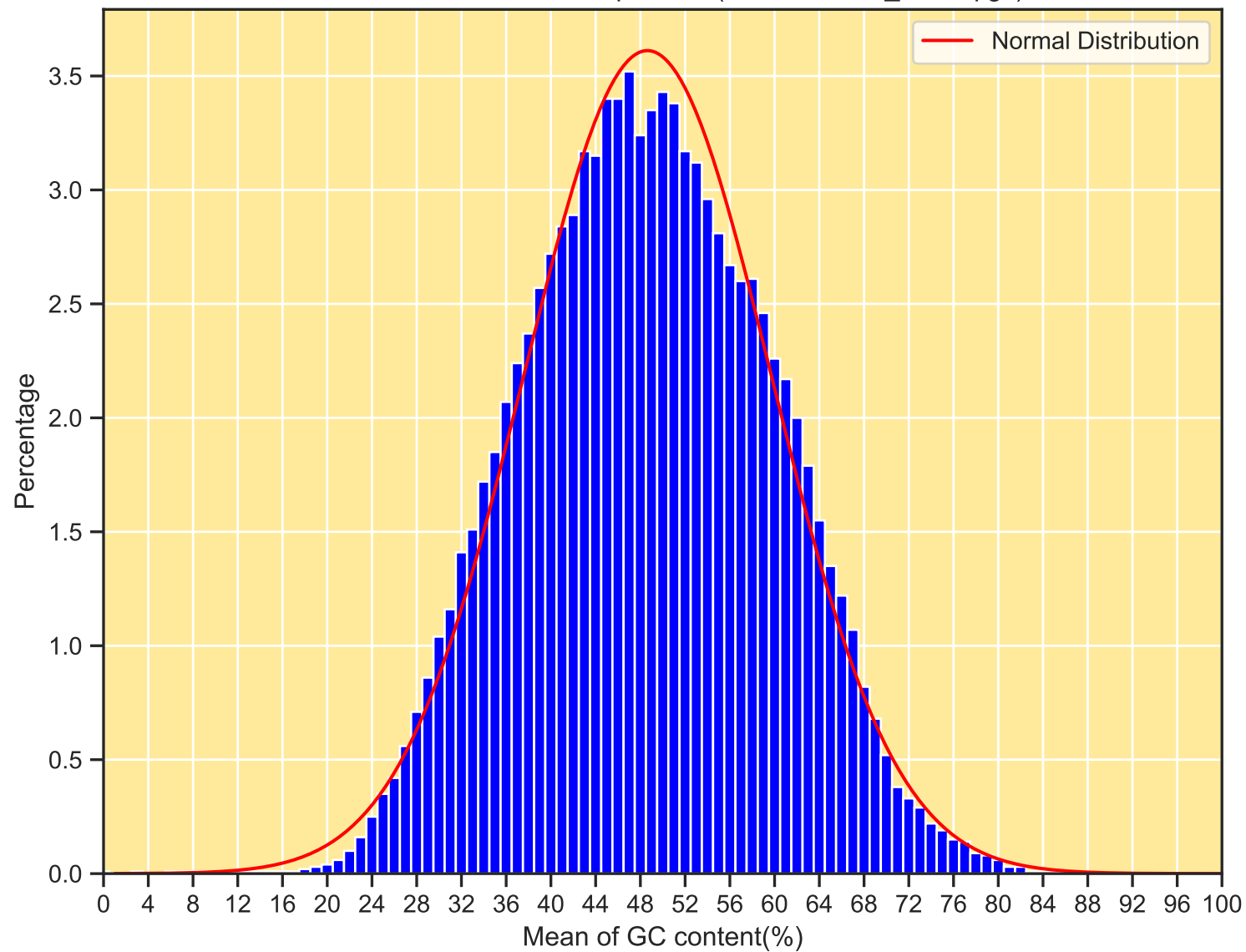
There are no Adapters in SRR1608628_1.fastq.gz



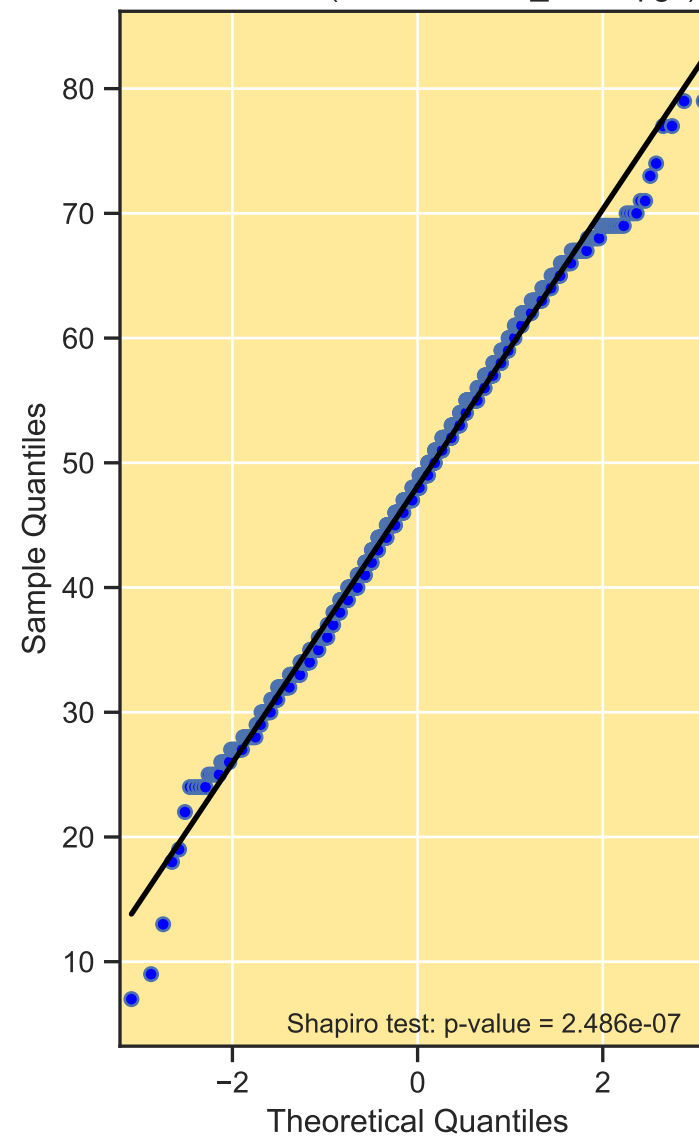
Sequence Duplication Levels (SRR1608628_1.fastq.gz)



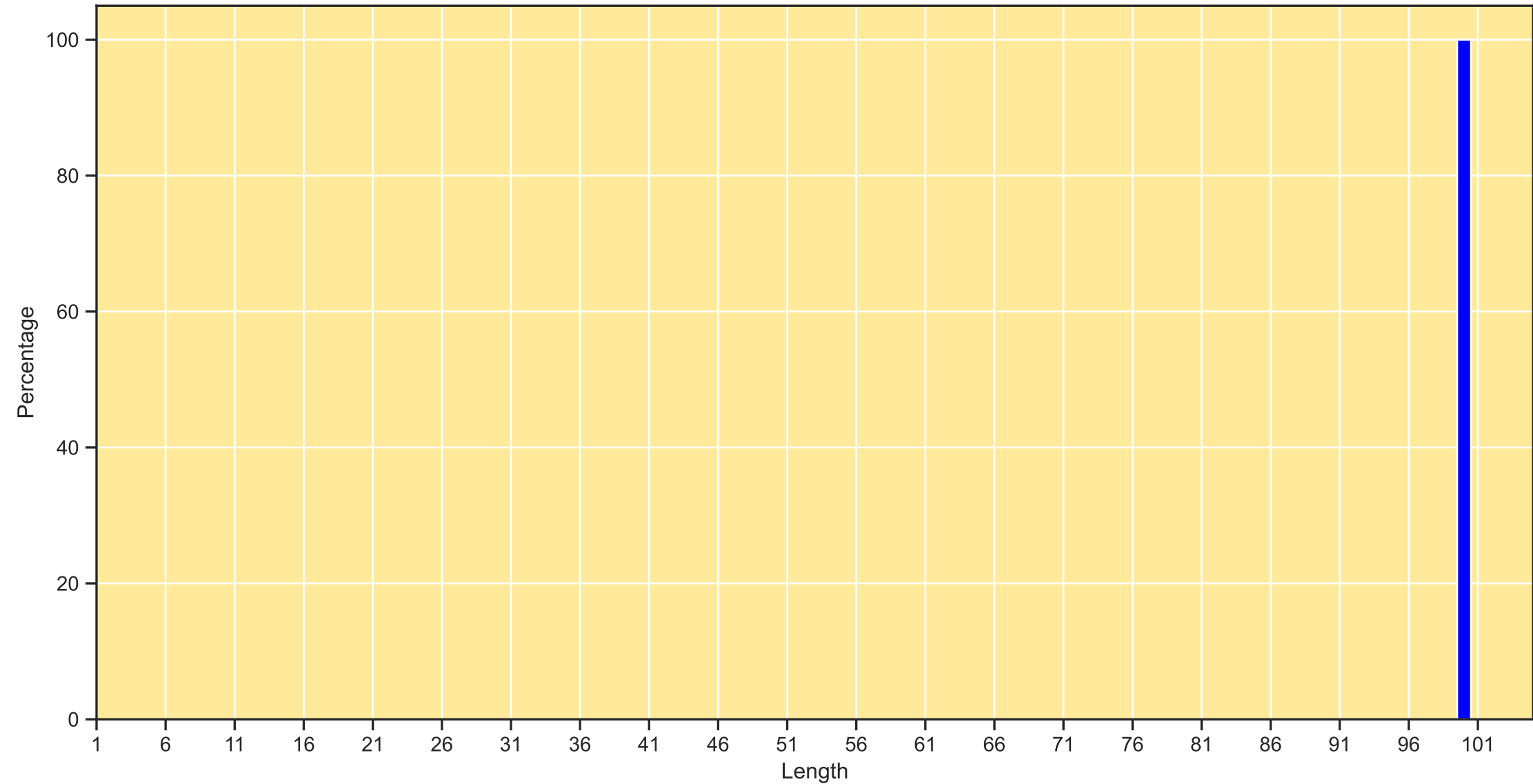
GC distribution over all sequences (SRR1608628_1.fastq.gz)



GC Q-Q Plot (SRR1608628_1.fastq.gz)

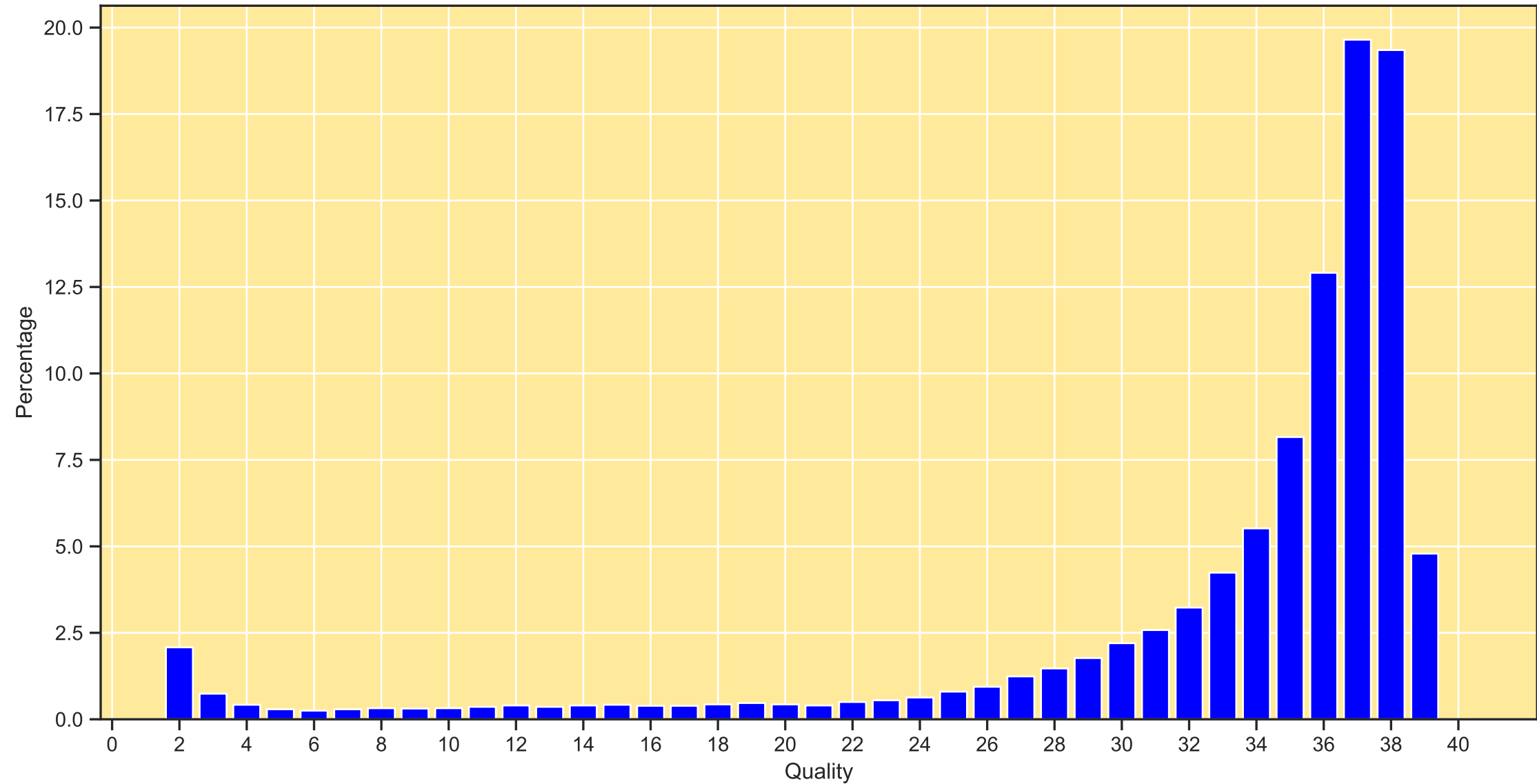


Length Distribution (SRR1608628_1.fastq.gz)



There are no overrepresented sequences in SRR1608628_1.fastq.gz

Per sequence quality scores (SRR1608628_1.fastq.gz)



Quality scores across all bases (SRR1608628_1.fastq.gz)

