

QC SUMMARY REPORT

General Statistics

Copy table

Configure Columns

Plot

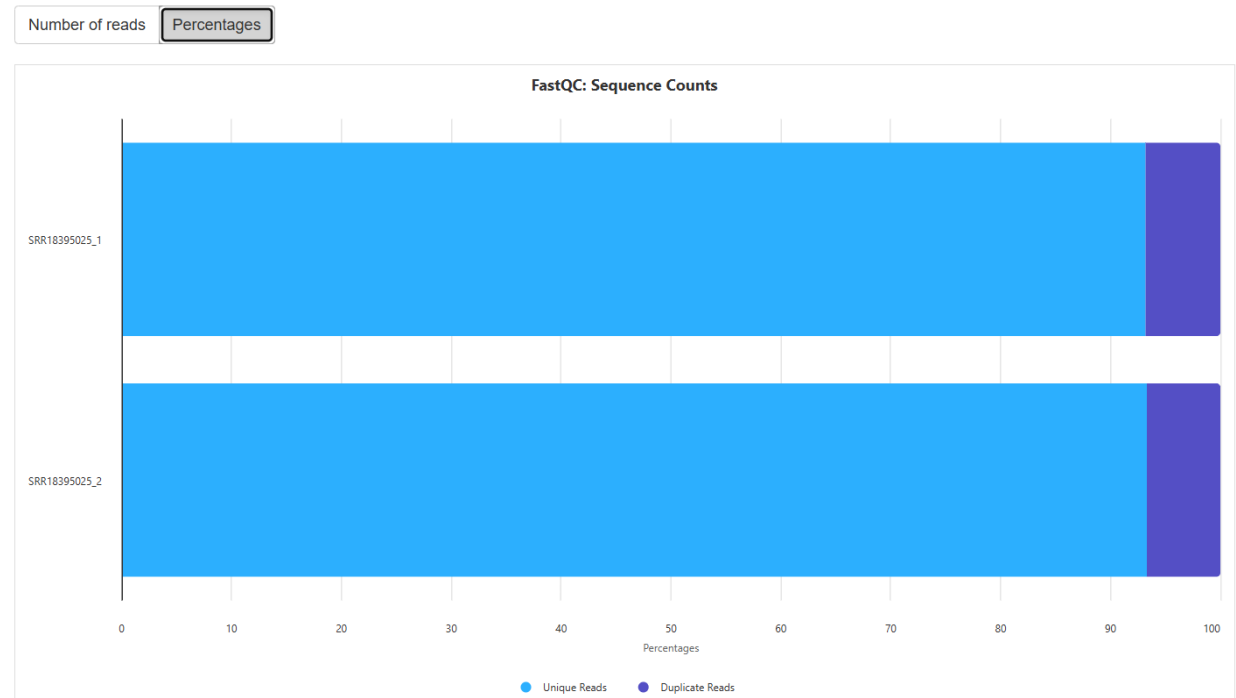
Showing $\frac{2}{3}$ rows and $\frac{3}{6}$ columns.

Sample Name	% Dups	% GC	M Seqs
SRR18395025_1	6.8%	53%	0.1
SRR18395025_2	6.6%	53%	0.1

Sequence Counts

Help

Sequence counts for each sample. Duplicate read counts are an estimate only.



Sequence Quality Histograms

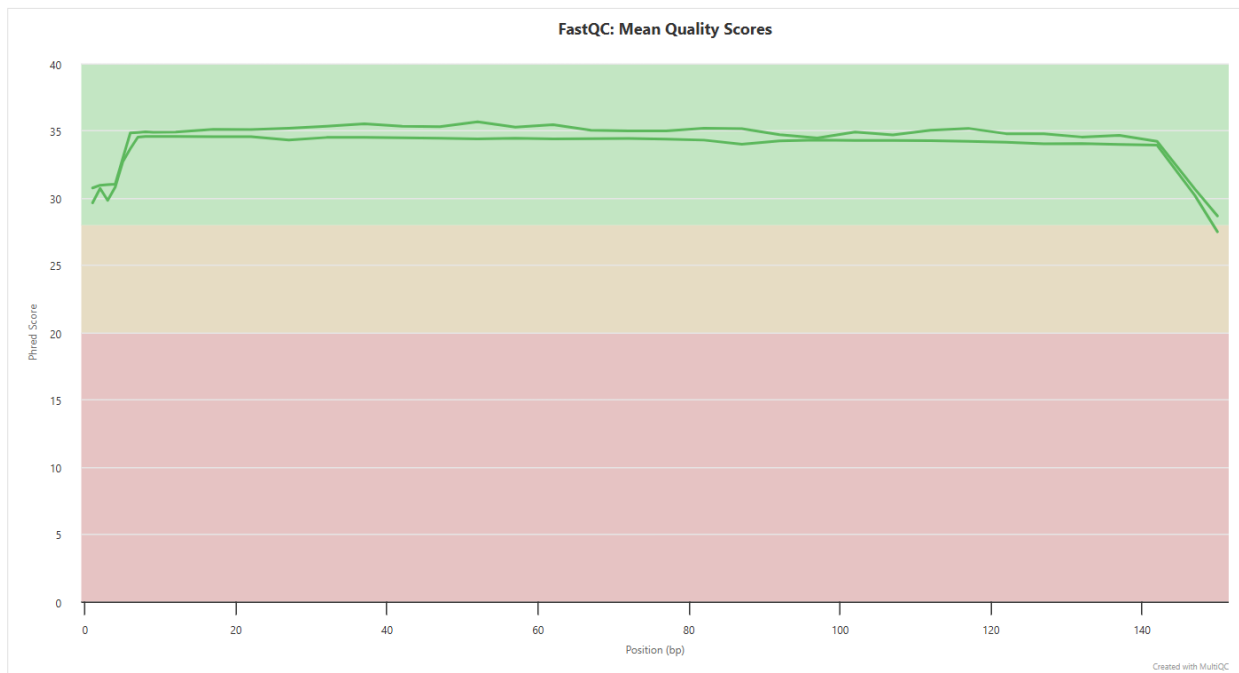
2

2

[Help](#)

The mean quality value across each base position in the read.

Y-Limits: ☒ on



Per Sequence Quality Scores

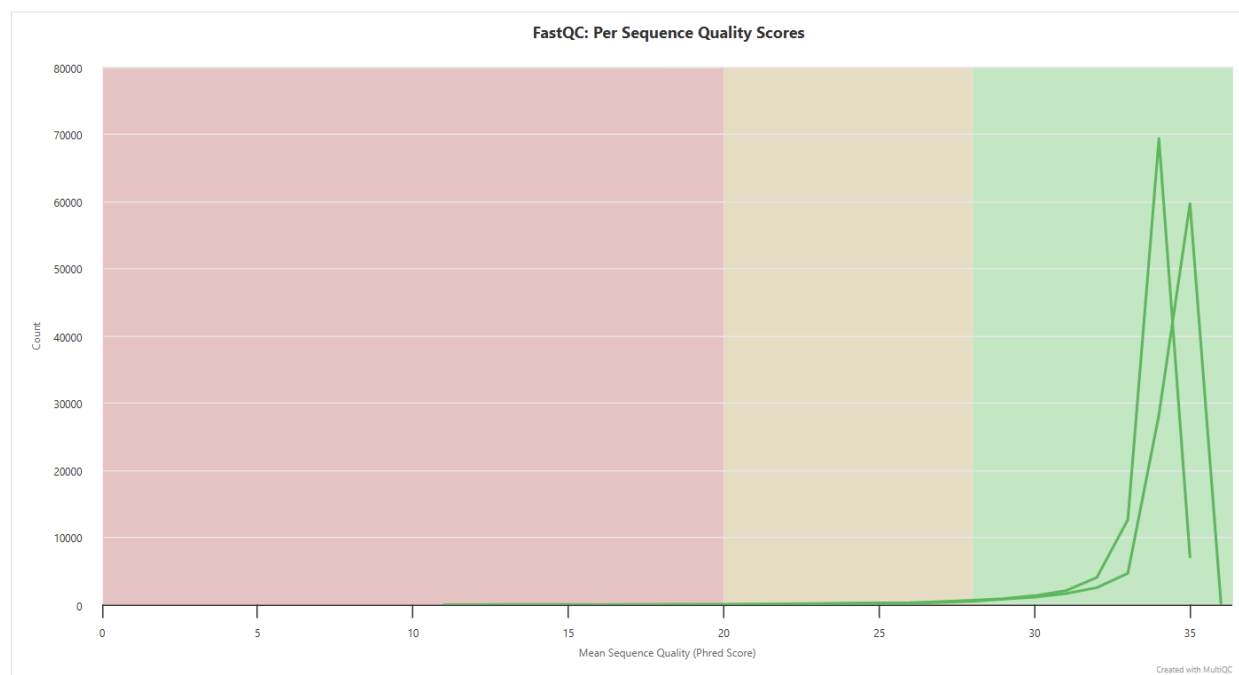
2

2

[Help](#)

The number of reads with average quality scores. Shows if a subset of reads has poor quality.

Y-Limits: ☒ on



Per Sequence GC Content

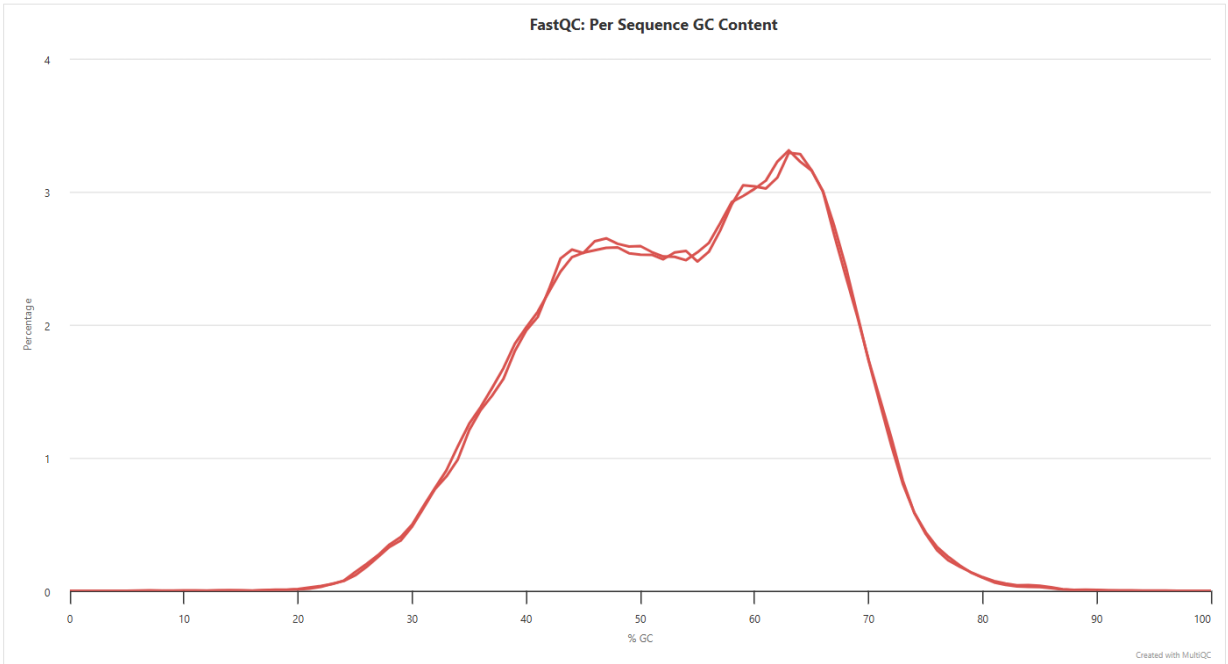
2 2

Help

The average GC content of reads. Normal random library typically have a roughly normal distribution of GC content.

Y-Limits: on

Percentages Counts



Adapter Content

2 2

Help

The cumulative percentage count of the proportion of your library which has seen each of the adapter sequences at each position.

Y-Limits: on

