

# Adapter lengths for all datasets

2025-03-19

## Load all data

```
a60_unmod <- read.table(file = "a60_unmod_polyA_position.tsv", sep = "\t", header = TRUE)
a60_30 <- read.table(file = "a60_30_polyA_position.tsv", sep = "\t", header = TRUE)
a60_60 <- read.table(file = "a60_60_polyA_position.tsv", sep = "\t", header = TRUE)

a120_unmod <- read.table(file = "a120_unmod_polyA_position.tsv", sep = "\t", header = TRUE)
a120_1mod <- read.table(file = "a120_1mod_polyA_position.tsv", sep = "\t", header = TRUE)
a120_2mod <- read.table(file = "a120_2mod_polyA_position.tsv", sep = "\t", header = TRUE)
a120_4mod <- read.table(file = "a120_4mod_polyA_position.tsv", sep = "\t", header = TRUE)
```

## Define lengths of adapter

```
a60_unmod["adapter_length"] <- a60_unmod$start - 1
a60_30["adapter_length"] <- a60_30$start - 1
a60_60["adapter_length"] <- a60_60$start - 1

a120_unmod["adapter_length"] <- a120_unmod$start - 1
a120_1mod["adapter_length"] <- a120_1mod$start - 1
a120_2mod["adapter_length"] <- a120_2mod$start - 1
a120_4mod["adapter_length"] <- a120_4mod$start - 1
```

## Histograms of all the data in each set

```
datasets <- list(
  a60_30 = a60_30,
  a60_60 = a60_60,
  a60_unmod = a60_unmod,
  a120_1mod = a120_1mod,
  a120_2mod = a120_2mod,
  a120_4mod = a120_4mod,
  a120_unmod = a120_unmod
)

for (name in names(datasets)) {
  adapter_length <- datasets[[name]]$adapter_length

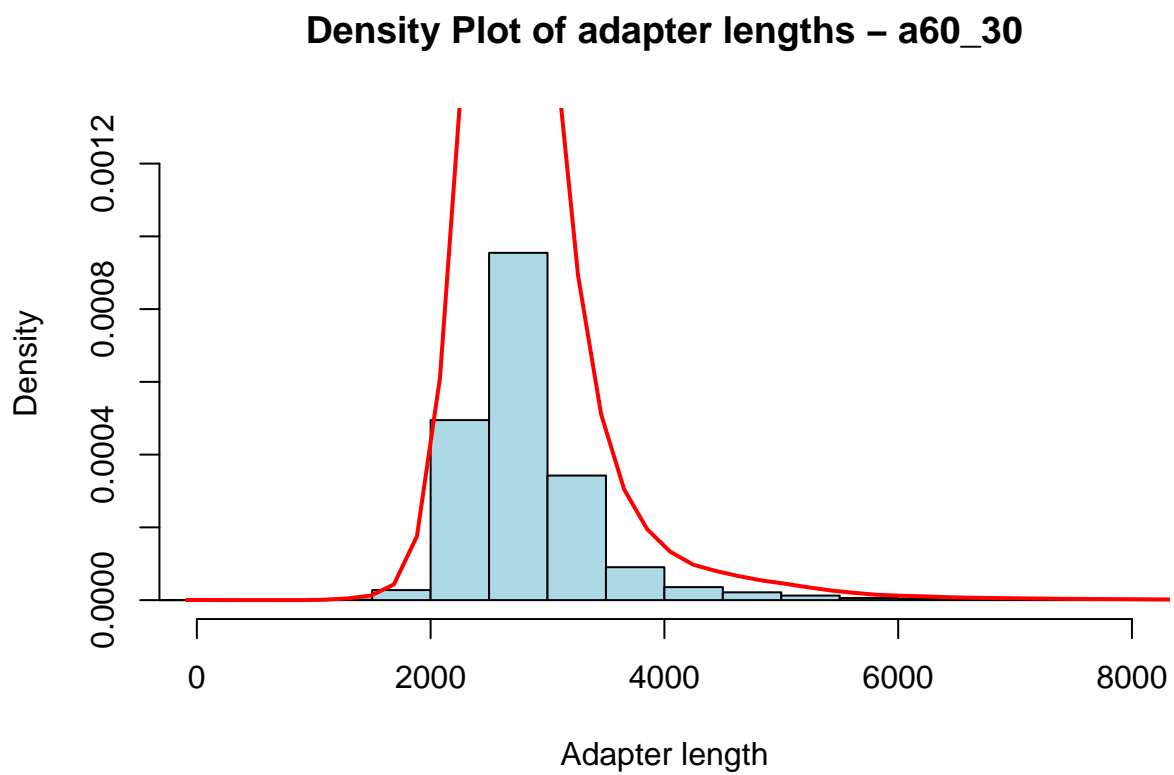
  hist(adapter_length,
        probability = TRUE,
```

```

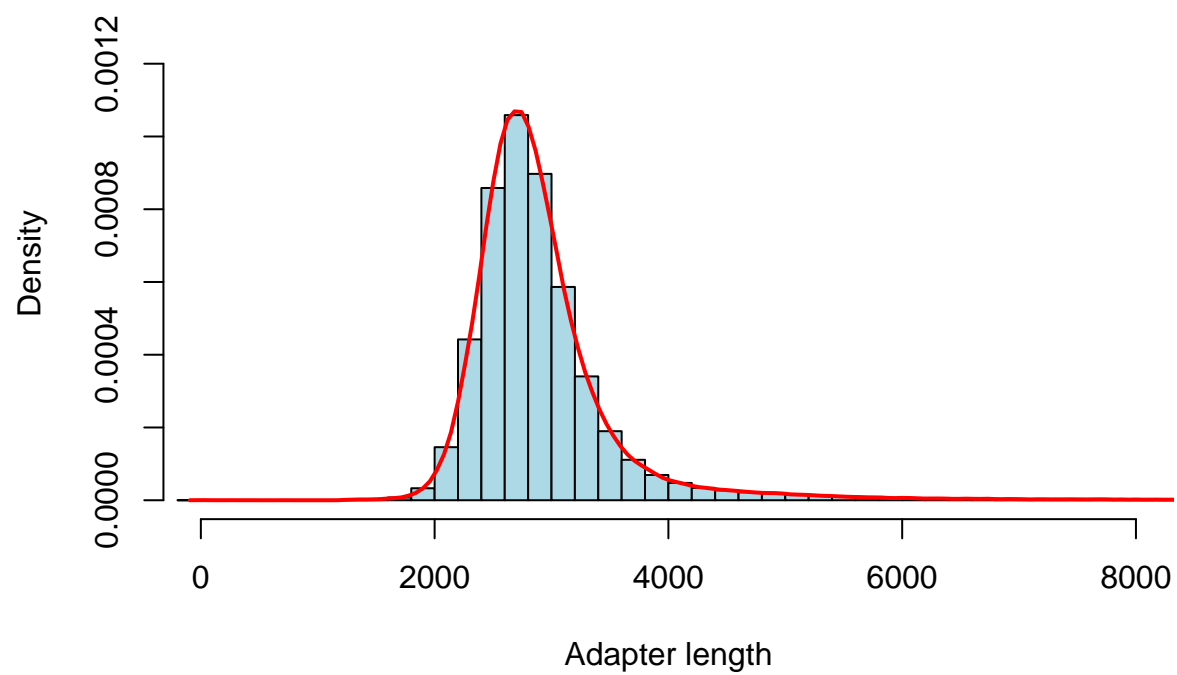
main = paste("Density Plot of adapter lengths -", name),
xlab = "Adapter length",
ylab = "Density",
xlim = c(0, 8000),
ylim = c(0, 0.0013),
col = "lightblue",
border = "black",
breaks = 200)

lines(density(adapter_length), col = "red", lwd = 2)
}

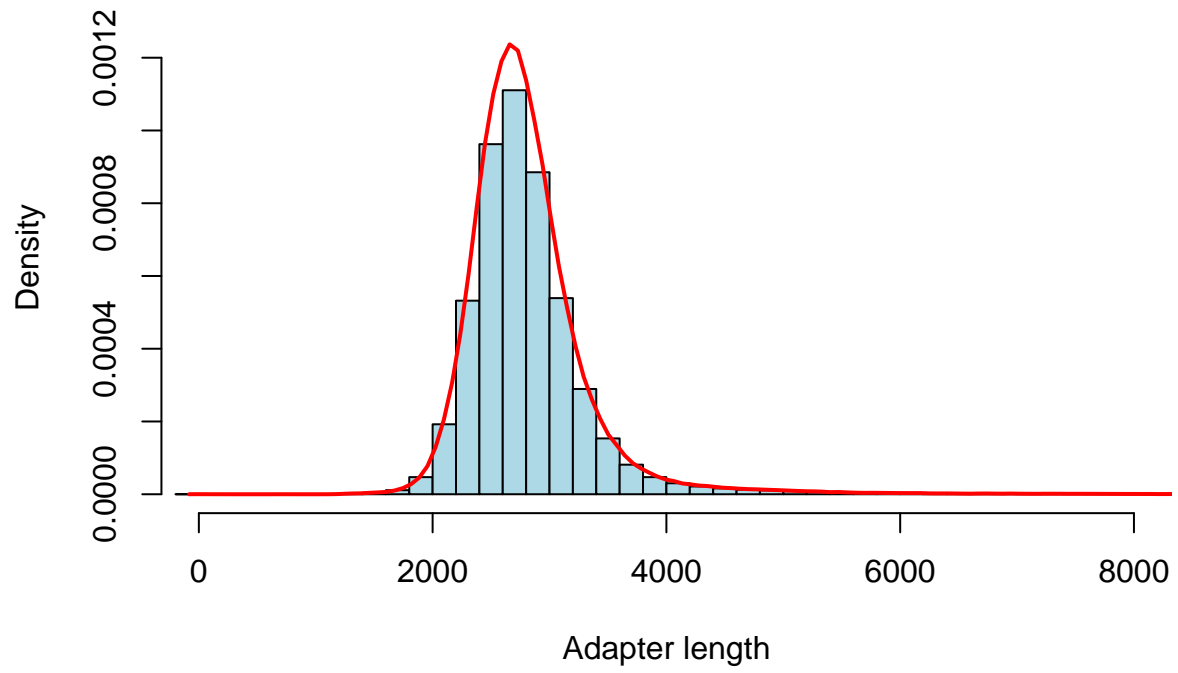
```



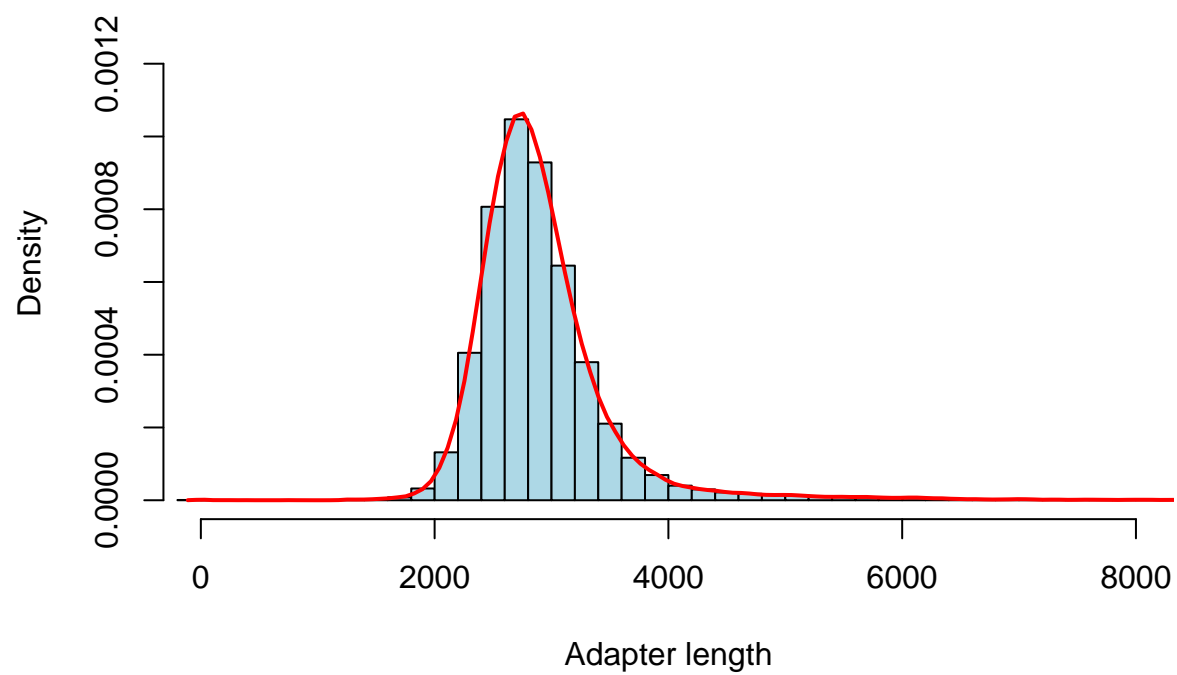
**Density Plot of adapter lengths – a60\_60**



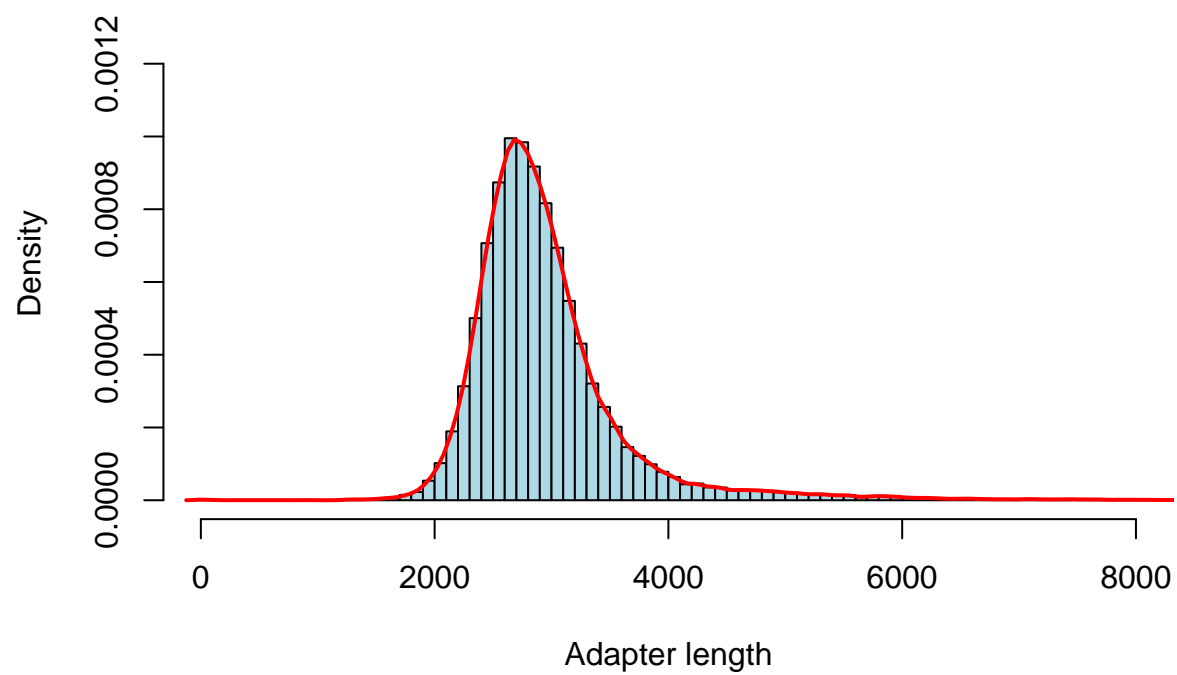
**Density Plot of adapter lengths – a60\_unmod**



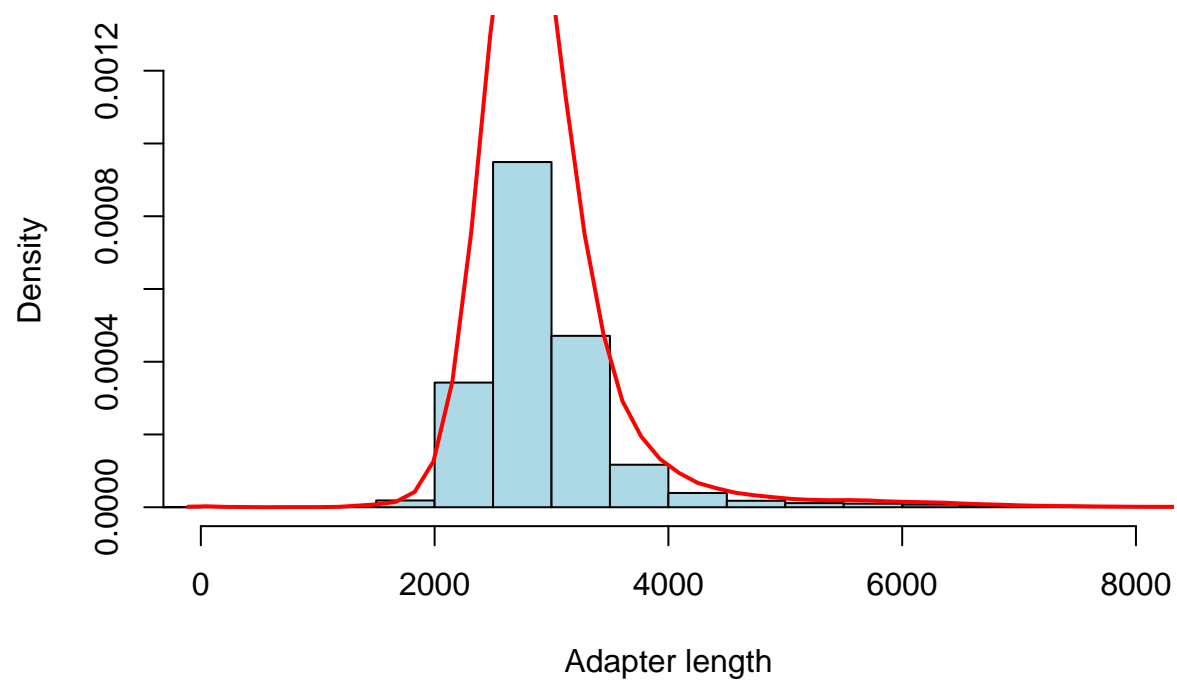
**Density Plot of adapter lengths – a120\_1mod**



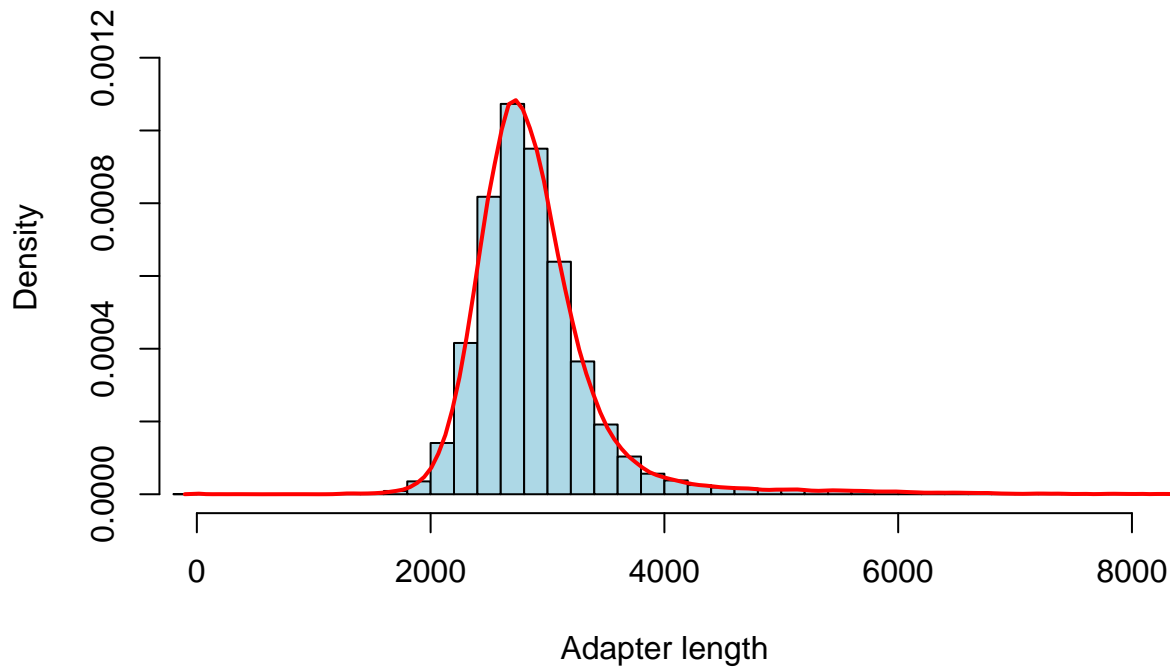
**Density Plot of adapter lengths – a120\_2mod**



**Density Plot of adapter lengths – a120\_4mod**



## Density Plot of adapter lengths – a120\_unmod

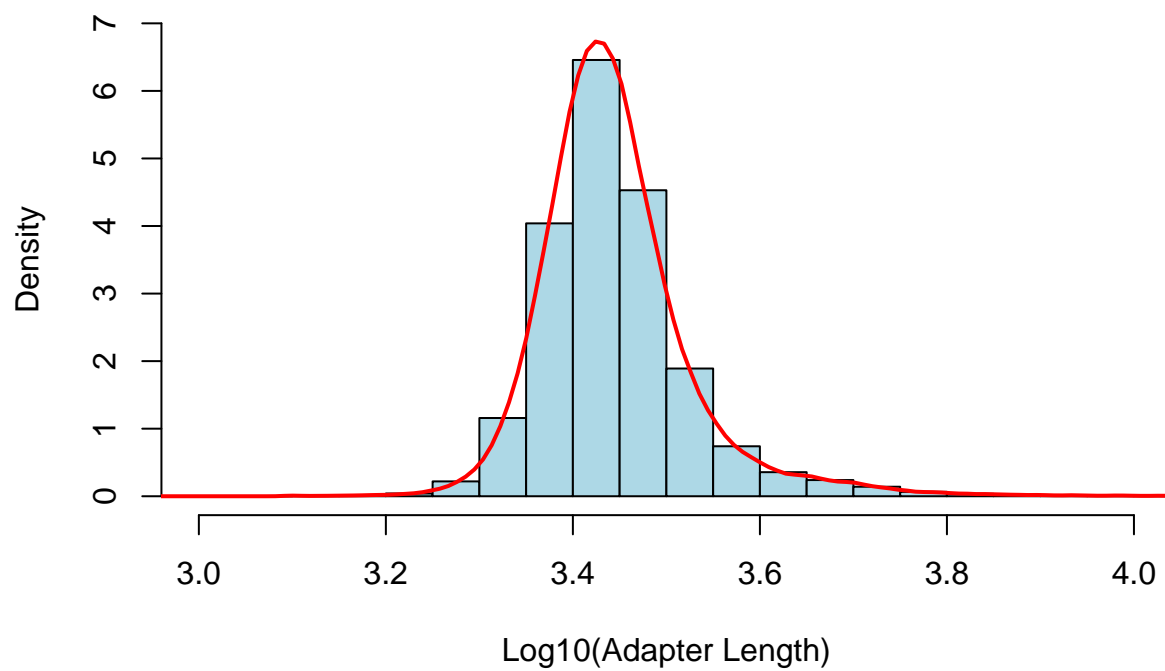


Histograms of all the data with log transformation

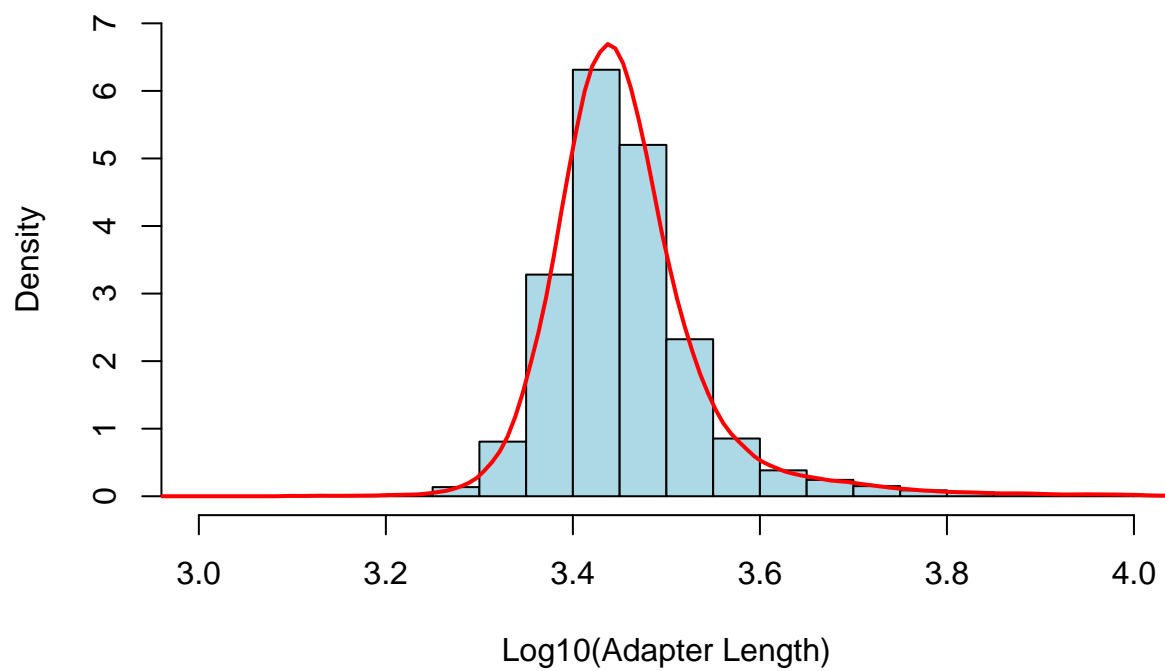
```
for (name in names(datasets)) {  
  adapter_length <- datasets[[name]]$adapter_length  
  
  # Remove values <= 0  
  adapter_length <- adapter_length[adapter_length > 0]  
  
  log_adapter_length <- log10(adapter_length)  
  
  hist(log_adapter_length,  
        probability = TRUE,  
        main = paste("Density Plot of log-transformed adapter lengths -", name),  
        xlab = "Log10(Adapter Length)",  
        ylab = "Density",  
        border = "black",  
        col = "lightblue",  
        xlim = c(3, 4),  
        ylim = c(0, 7),  
        breaks = 100)  
  
  lines(density(log_adapter_length), col = "red", lwd = 2)  
}
```



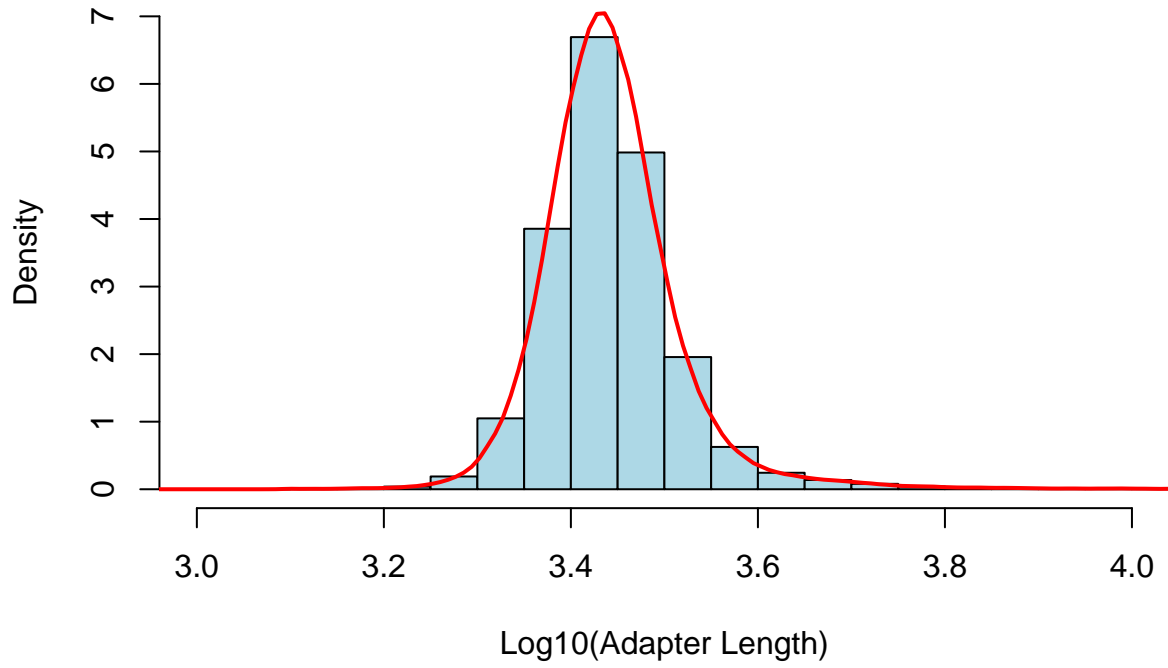
**Density Plot of log-transformed adapter lengths – a60\_30**



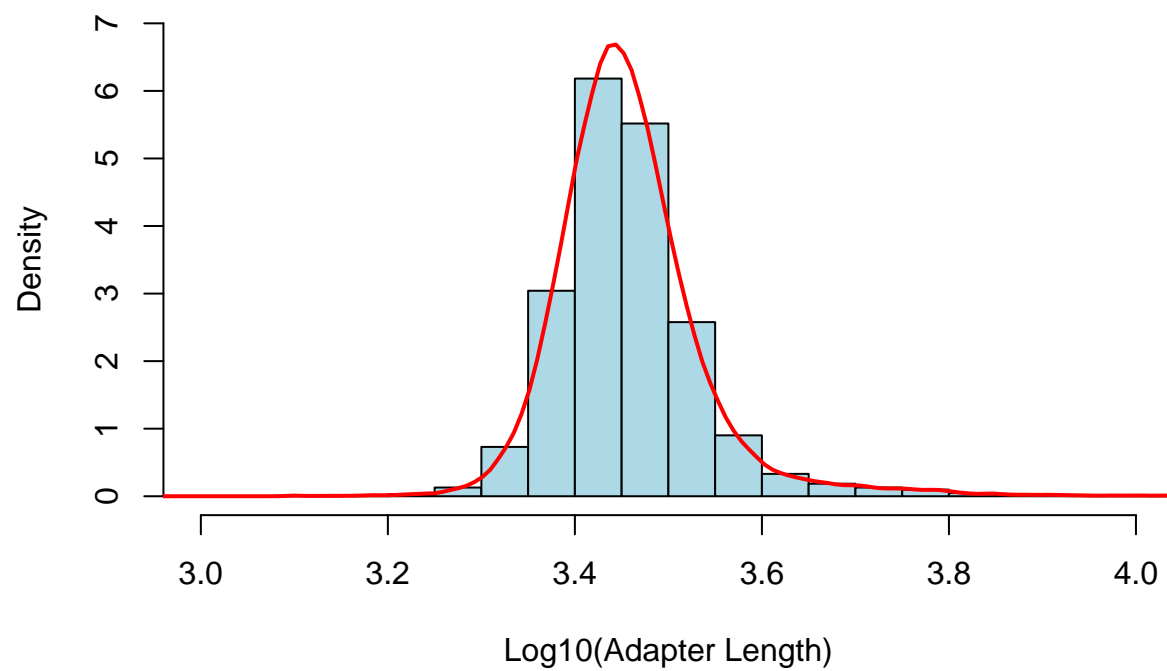
**Density Plot of log-transformed adapter lengths – a60\_60**



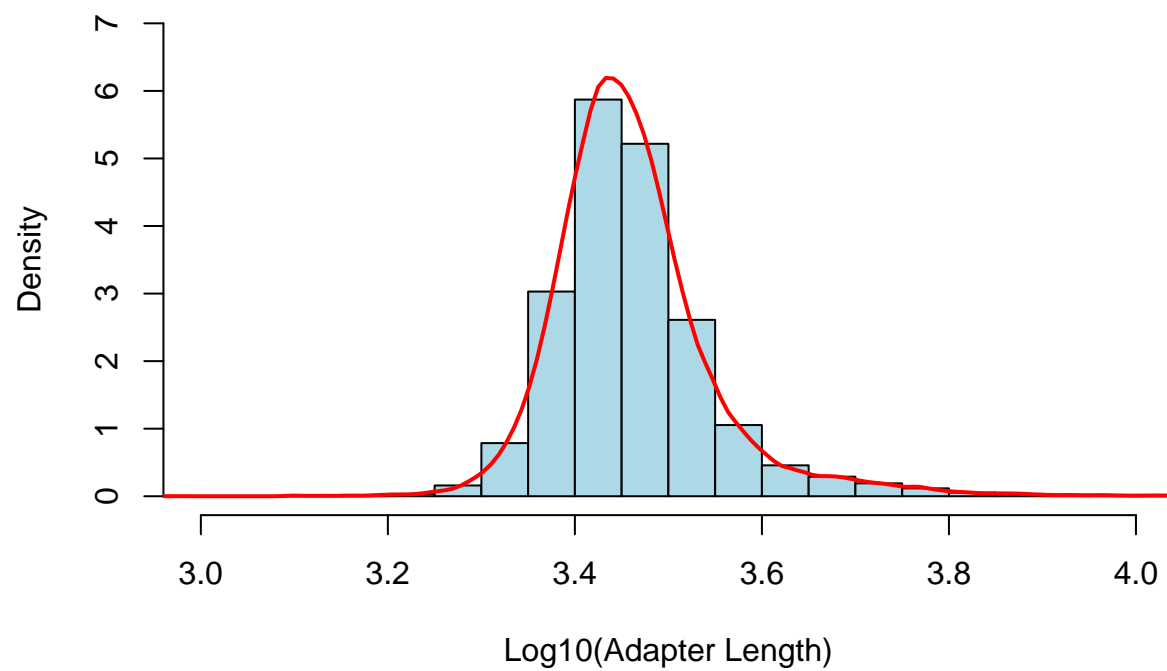
**Density Plot of log-transformed adapter lengths – a60\_unmod**



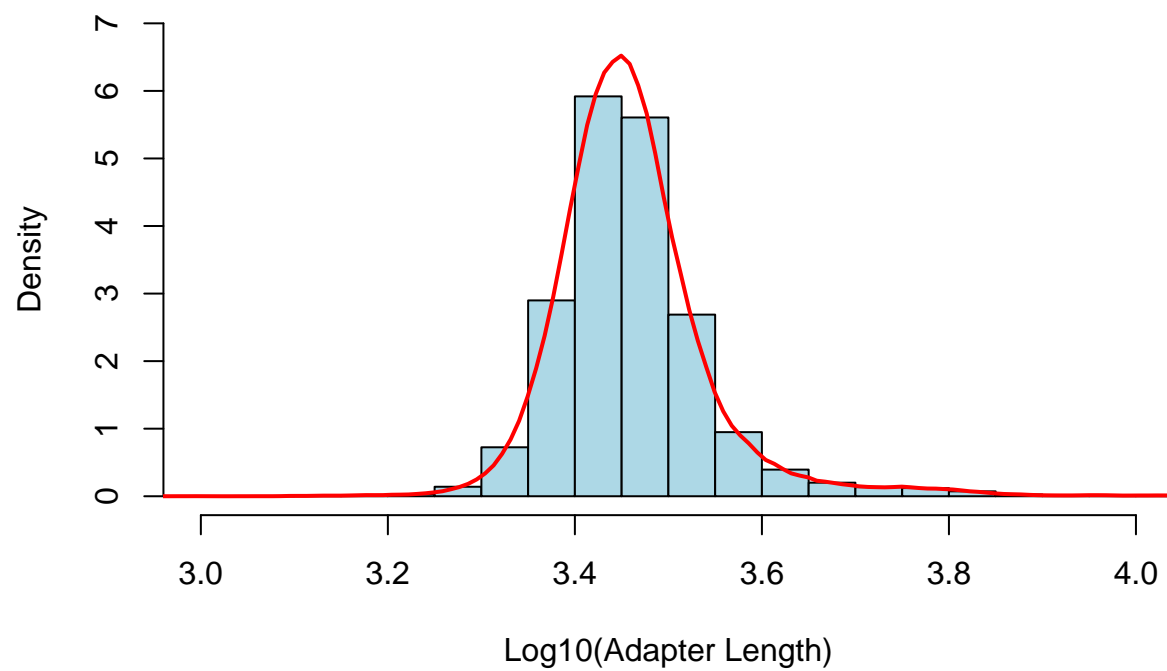
**Density Plot of log-transformed adapter lengths – a120\_1mod**



**Density Plot of log-transformed adapter lengths – a120\_2mod**



**Density Plot of log-transformed adapter lengths – a120\_4mod**



**Density Plot of log-transformed adapter lengths – a120\_unmod**

