# Lab 4: Heart of the (Tiny) Tiger

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### Part I

1. Loading the data from the MASS package.

```
library(MASS)

## Warning: package 'MASS' was built under R version 3.1.3

data(cats)
summary(cats)
```

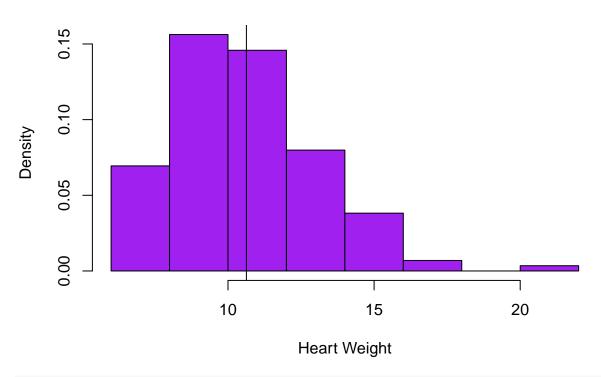
```
##
    Sex
                 Bwt
                                  Hwt
    F:47
                   :2.000
                                    : 6.30
##
           Min.
                             Min.
   M:97
##
           1st Qu.:2.300
                             1st Qu.: 8.95
           Median :2.700
                             Median :10.10
##
##
           Mean
                   :2.724
                             Mean
                                    :10.63
                             3rd Qu.:12.12
##
           3rd Qu.:3.025
##
           Max.
                   :3.900
                                    :20.50
                             Max.
```

The summary shows that the there were 47 female and 97 male cats whose weight and heart weights were collected. The minimum body weight out of all the cats was 2kg and the maximum was 3.9kg with an average body weight of 2.724kg. The minimum heart weight was 6.30g and the maximum was 30.50g with an average heart weight of 10.63g. There are other statistics present in the summary such as the first and third quartile as well as the median.

2.

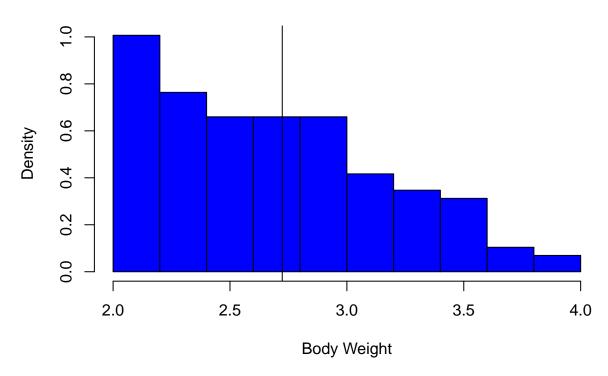
```
hist(cats$Hwt, xlab="Heart Weight",col="purple",probability=TRUE)
abline(v=mean(cats$Hwt))
```

## Histogram of cats\$Hwt



hist(cats\$Bwt, xlab="Body Weight",col="blue",probability=TRUE)
abline(v=mean(cats\$Bwt))

# Histogram of cats\$Bwt



3. Function param calculates the a and s

$$a = \frac{a^2 s^2}{as^2} = \frac{\mu^2}{\sigma^2}$$
$$s = \frac{as^2}{as} = \frac{\sigma^2}{\mu}$$

and assuming that  $\mu$  and  $\sigma$  come from a random sample from a gamma distribution.

```
param <- function(mu,var){
  a=(mu^2)/(var)
  s=(var)/mu
  return(c('a'=a,'s'=s))
}</pre>
```

4.

```
hmean <-mean(cats$Hwt)
hsd <-sd(cats$Hwt)
hvar <-var(cats$Hwt)
cat("The a and s are:", param(hmean,hvar),"respectively.")</pre>
```

## The a and s are: 19.06531 0.5575862 respectively.

5. The function *cat.stats()* computes the mean, variance, a (shape) and s(scale) of the input vector of values

```
cat.stats <- function(vec)
{
    m<-mean(vec)
    var <- var(vec)
    a<-(m^2)/var
    s<-var/m
    return(c("Mean"=m,"Variance"=var,"a"=a,"s"=s))
}</pre>
```

#### Part II

6.

```
males<-which(cats$Sex=="M")
cat("The body weight parameters for male cats are:",cat.stats(cats$Bwt[males]))</pre>
```

## The body weight parameters for male cats are: 2.9 0.2185417 38.48236 0.0753592

```
cat("The heart weight paramters for male cats are:", cat.stats(cats$Hwt[males]))
```

## The heart weight paramters for male cats are: 11.32268 6.46323 19.83576 0.5708216

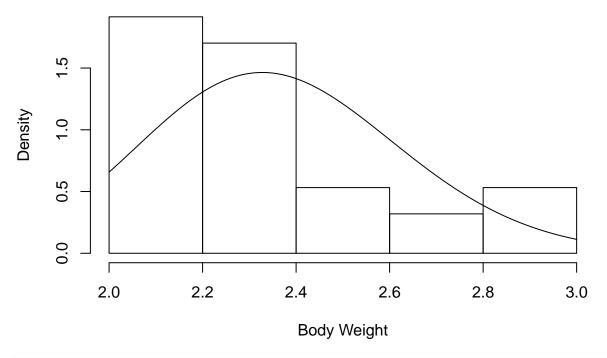
```
females<-which(cats$Sex=="F")
cat("The body weight parameters for female cats are:",cat.stats(cats$Bwt[females]))
## The body weight parameters for female cats are: 2.359574 0.07506938 74.16595 0.0318148
cat("The heart weight parameters for female cats are:", cat.stats(cats$Hw[females]))</pre>
```

## The heart weight paramters for female cats are: 9.202128 1.843256 45.93998 0.2003076

#### 7. Females

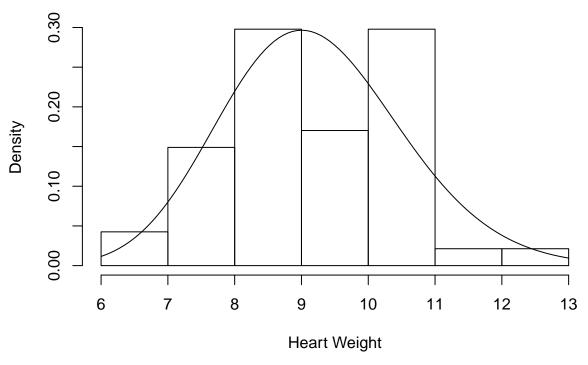
hist(cats\$Bwt[females],main='Female Cats Body Weight',xlab='Body Weight',probability=TRUE)
curve(dgamma(x,shape=cat.stats(cats\$Bwt[females])[3],scale=cat.stats(cats\$Bwt[females])[4]),add=TRUE)

### **Female Cats Body Weight**



hist(cats\$Hwt[females],main='Female Cats Heart Weight',xlab='Heart Weight',probability=TRUE)
curve(dgamma(x,shape=cat.stats(cats\$Hwt[females])[3],scale=cat.stats(cats\$Hwt[females])[4]),add=TRUE)

## **Female Cats Heart Weight**

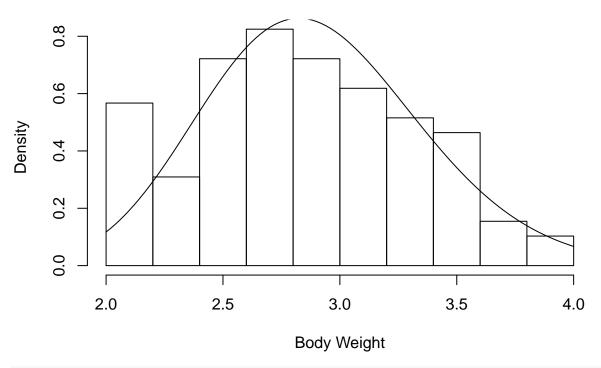


The PDF of the Gamma Distribution distribution seems to be consistent with the empirical probability density of the histogram. However, if we sample more cats, we will get a histogram that looks more like the PDF.

#### 8. Males

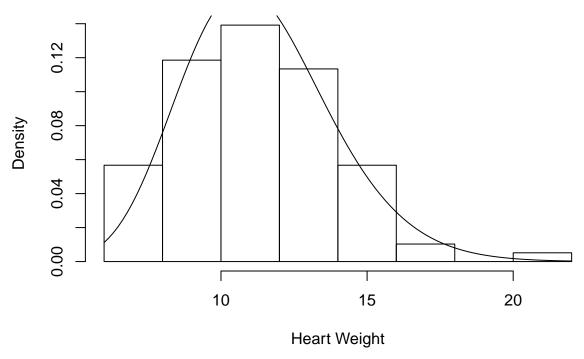
```
# the scales changes by letting porability=TRUE
hist(cats$Bwt[males],main='Male Cats Body Weight',xlab='Body Weight',probability=TRUE)
curve(dgamma(x,shape=cat.stats(cats$Bwt[males])[3],scale=cat.stats(cats$Bwt[males])[4]),add=TRUE)
```

### **Male Cats Body Weight**



hist(cats\$Hwt[males],main='Male Cats Heart Weight',xlab='Heart Weight',probability=TRUE)
curve(dgamma(x,shape=cat.stats(cats\$Hwt[males])[3],scale=cat.stats(cats\$Hwt[males])[4]),add=TRUE)

## **Male Cats Heart Weight**



We can say the same thing about the male cats as we did for female cats. Also, we can say that the distribution is is skewed to the right which means that the right tail of the distribution is heavier.