

Lab 3

AnyaConti28661255

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1. First, the R package MASS is loaded using `library()`, and then the data called `cats` is loaded using `data()`. This data is then summarized using `summary()`. The first section entitled `sex` shows how many females and how many males there are, then the second section called `Bwt` shows summary statistics for the body weight, and the third section called `Hwt` shows the summary statistics for heart weight (`hwt`). There is no cross-analysis between males vs females for these categories however.

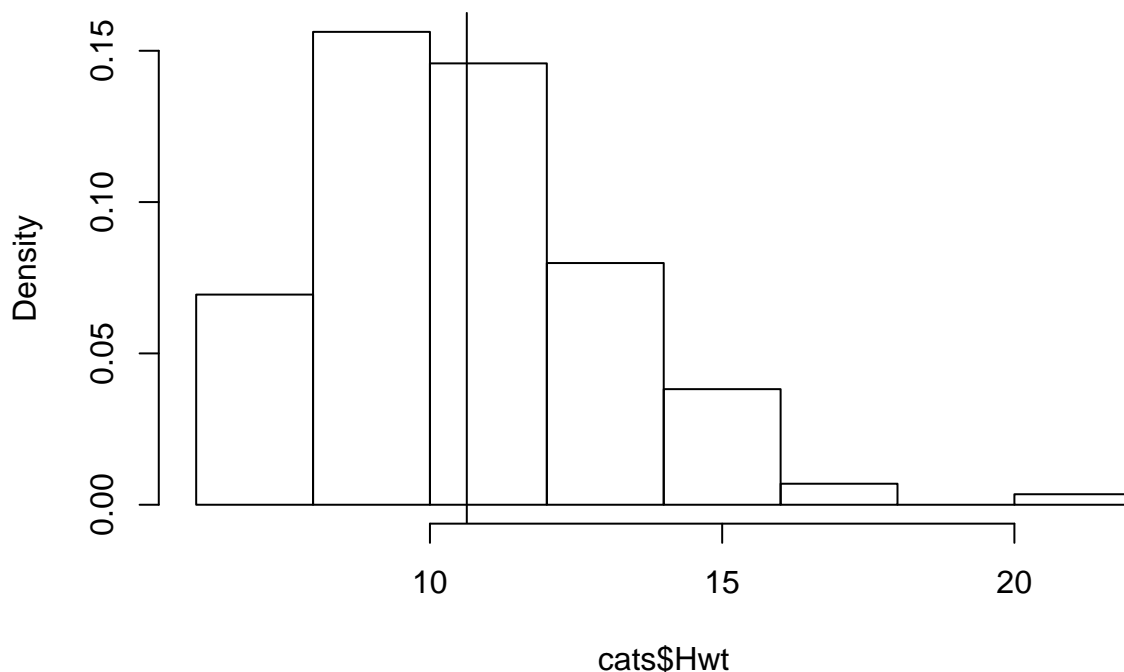
```
library(MASS)
data(cats)
summary(cats)
```

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

2. A histogram of the heart weights is then plotted using the `probability=TRUE` so that the total area under them is equal to one, so it can be a probability function. Then `abline()` was used to add a line. The `v` means vertical, and it is set to be at the mean of the heart weights of `cats`.

```
hist(x=cats$Hwt, probability = TRUE)
abline(v=mean(cats$Hwt))
```

Histogram of cats\$Hwt



3. Writes a function called `gammafunction` which takes two inputs (mean and variance) and calculates `a` and `s` from them, then creates a list called `output` that has both new values inside it, and returns that output.

```
gammafunction <- function(gamma.mean, gamma.var){  
  a = (gamma.mean ^ 2) / (gamma.var)  
  s = (gamma.var) / (gamma.mean)  
  output <- c("a"=a, "s"=s)  
  return(output)  
}
```

4. Calculates the mean of the heart weights of cats which is 10.631. Then calculates the standard deviation of the heart weights of cats which is 2.435. Then calculates the variance of the heart weights of cats which is 5.927. Then uses the gamma function from question 3 with the cat heart weight mean and variance as the inputs to calculate `a` and `s` for the data set. The calculated value for `a` is 19.065 and the calculated value for `s` is 0.558.

```
mean(cats$Hwt)
```

```
## [1] 10.63056
```

```
sd(cats$Hwt)
```

```
## [1] 2.434636
```

```
var(cats$Hwt)
```

```
## [1] 5.927451
```

```
gammafunction(mean(cats$Hwt), var(cats$Hwt))
```

```
##           a           s  
## 19.0653121  0.5575862
```

5. Creates a function called `cat.stats` with 1 input called `data`. The function then does the following. It calculates the mean of the data vector input, and assigns it to a new variable called `cats.mean`. It calculates the variance of the data vector input and assigns it to a new variable called `cats.var`. It calculates an estimated `a` value for the data vector from the mean and the variance, and assigns it to a new variable called `cats.a`. It calculates an estimated `s` value for the data vector from the mean and the variance, and assigns it to a new variable called `cats.s`. It creates a list called `output`, in which it stores all of these values, and then returns the output. An example of it running with the cats heart weights as the input data vector is shown below

```
cat.stats <- function(data){  
  cats.mean = mean(data)  
  cats.var = var(data)  
  cats.a = (cats.mean ^ 2) / (cats.var)  
  cats.s = (cats.var) / (cats.mean)  
  output <- c("mean"=cats.mean, "var"=cats.var, "a"=cats.a, "s"=cats.s)  
  return(output)  
}
```

```
cat.stats(cats$Hwt)
```

```
##      mean      var      a      s  
## 10.630556  5.927451 19.065312 0.5575862
```

6. First the `cat.stats` function is run with the cat heart weight data, but only for the entries where the sex is female. The results of this were saved as `FHwt`. This is then repeated, but only for the entries where the sex is male, and the results were saved as `MHwt`.

```

FHwt <- cat.stats(cats$Hwt[cats$Sex=="F"])
FHwt

##      mean      var      a      s
## 9.2021277 1.8432562 45.9399792 0.2003076

MHwt <- cat.stats(cats$Hwt[cats$Sex=="M"])
MHwt

##      mean      var      a      s
## 11.3226804 6.4632302 19.8357612 0.5708216

```

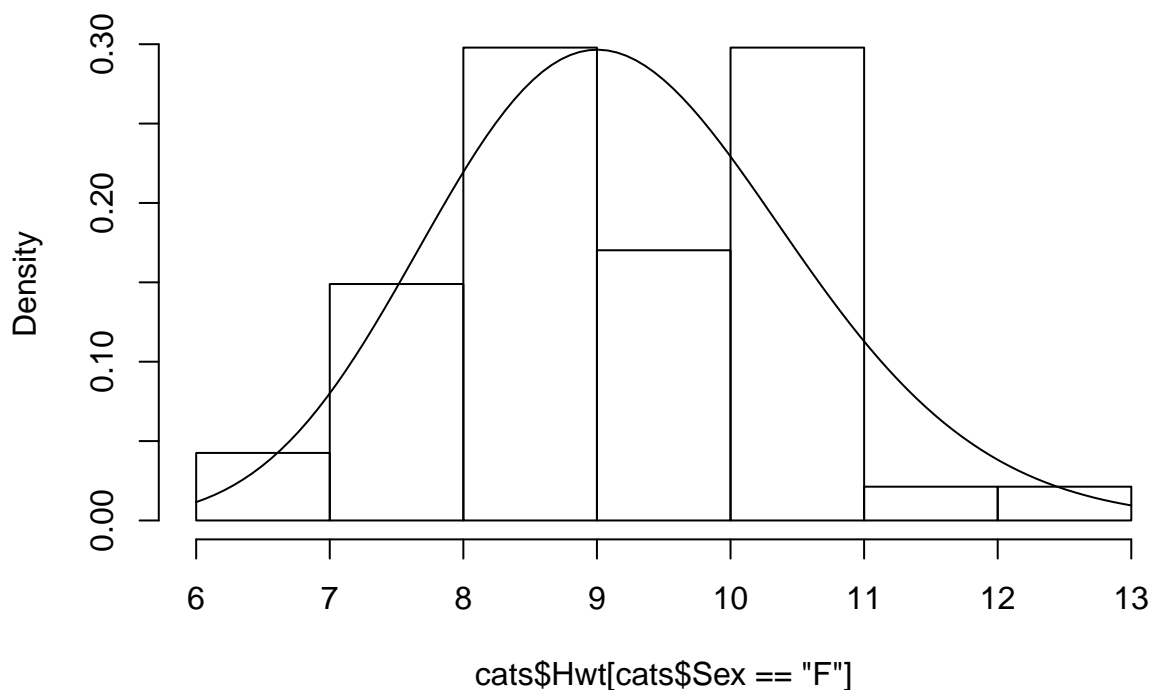
7. A histogram was created using the heart weight data, but only where the Sex is equal to “F” for female. Then the probability option was used, so that it would be a probability distribution function. Then a curve was added to this, where the expression was a gamma function with shape equal to the a value from the Female heart weight calculations from the previous problem (the 3rd entry of FHwt) and scale equal to the s value from the Female heart weight calculations (the 4th entry of FHwt).

```

hist(x=cats$Hwt[cats$Sex=="F"], probability = TRUE)
curve(dgamma(x, shape = FHwt[[3]], scale = FHwt[[4]]), add=TRUE)

```

Histogram of cats\$Hwt[cats\$Sex == "F"]



8. A histogram was created using the heart weight data, but only where the Sex is equal to “M” for male. Then the probability option was used, so that it would be a probability distribution function. (The y axis was adjust to have a slightly larger range so that the curve would fit on the graph.) Then a curve was added to this, where the expression was a gamma function with shape equal to the a value from the Male heart weight calculations from the previous problem (the 3rd entry of MHwt) and scale equal to the s value from the Female heart weight calculations (the 4th entry of MHwt).

```

hist(x=cats$Hwt[cats$Sex=="M"], probability = TRUE, ylim = c(0,0.16))
curve(dgamma(x, shape = MHwt[[3]], scale = MHwt[[4]]), add=TRUE)

```

Histogram of cats\$Hwt[cats\$Sex == "M"]

