```
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```

```
BirdBones <- read.csv("../data/bird.csv",header = T, sep = ",")
```

Introduction

Research Question

Are the bones of birds significantly distinct per group/functioning of birds, and can we use them to id

Data

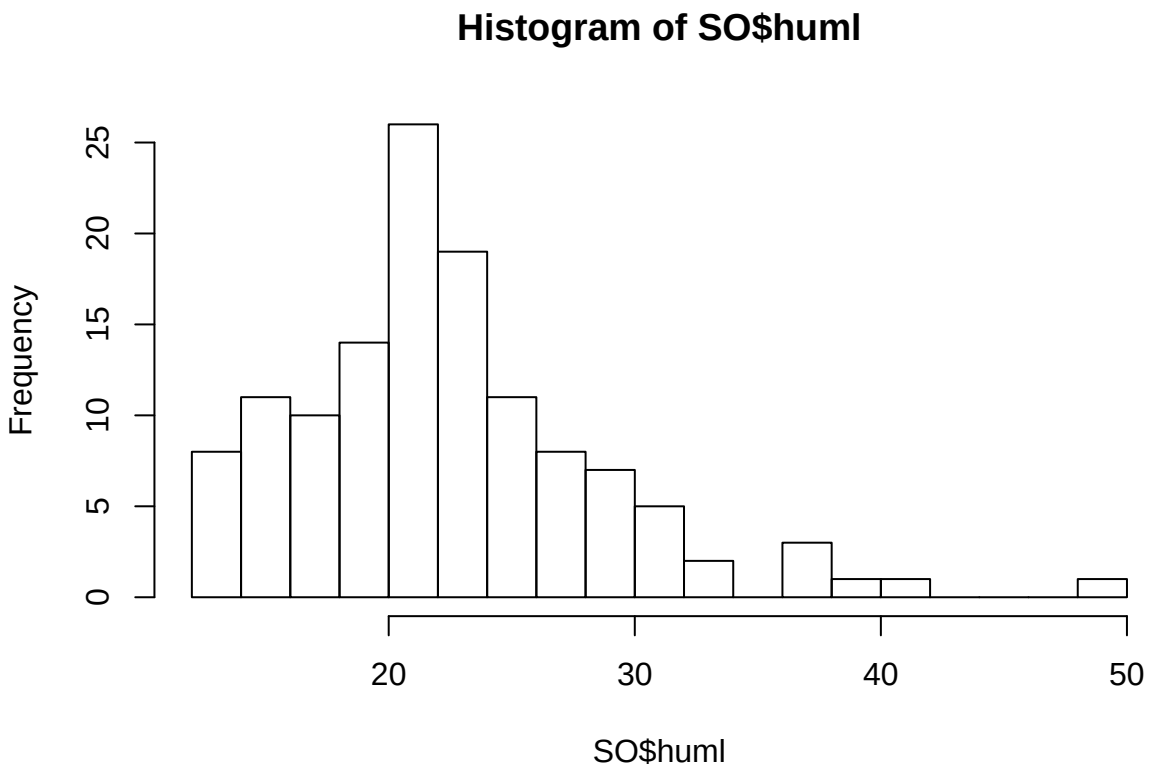
Data recieved from [Birds' Bones and Living Habits](https://www.kaggle.com/zhangjuefei/birds-bones-and-

Bone measurements were measured from a skeleton collection of Natural History Museum of Los Angeles Cou
provided by Dr. D. Liu of beijing Museaum of Natural History

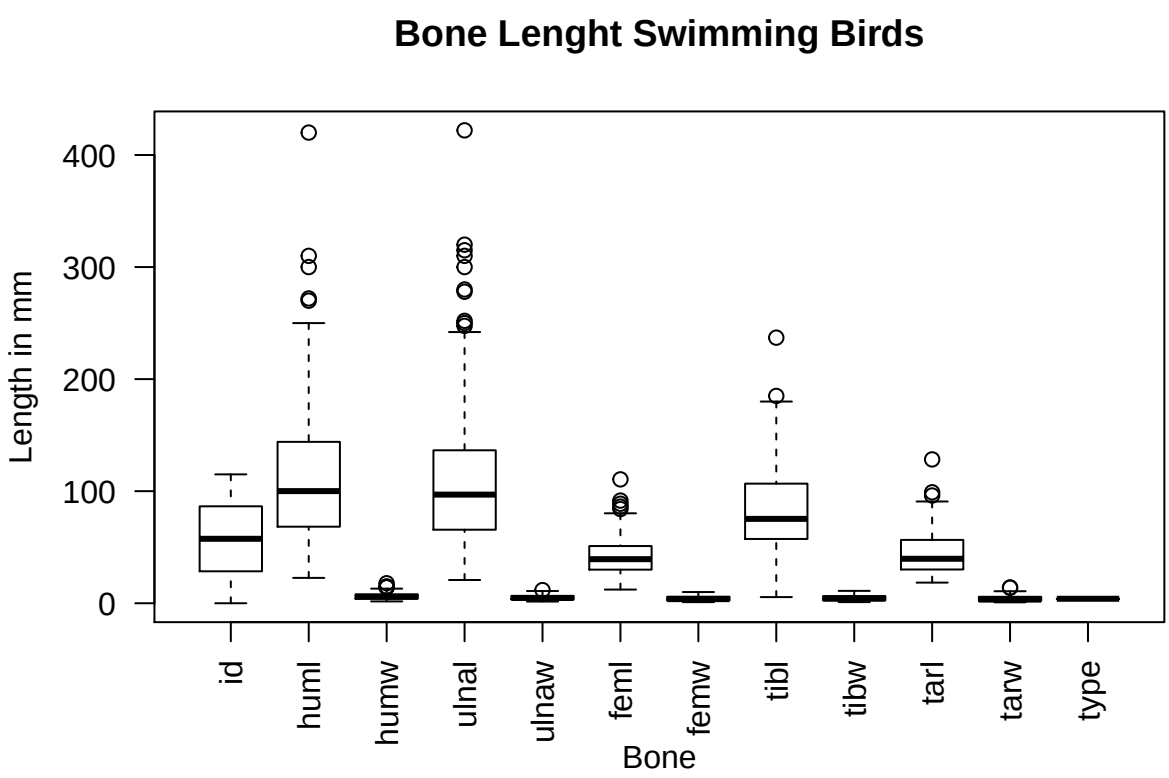
Exploratory Data Analyses

```
# Swimming
SW <- BirdBones[BirdBones$type == "SW",]
# Wading
W <- BirdBones[BirdBones$type == "W",]
# Terrestial
TER <- BirdBones[BirdBones$type == "T",]
# Raptors
R <- BirdBones[BirdBones$type == "R",]
# Scansorial
P <- BirdBones[BirdBones$type == "P",]
# Singing
SO <- BirdBones[BirdBones$type == "SO",]
```

```
source("../scripts/BoneMeans.R")
dataMeans <- BoneMeans(BirdBones)
hist(SO$huml, breaks = 15)
```




```
boxplot(SW, main = "Bone Lenght Swimming Birds",
        xlab = "Bone", ylab = "Length in mm", las=2)
```


Bone Lenght Swimming Birds
Length in mm
0
100
200
300
400
id
huml
humw
ulnal
ulnaw
feml
femw
tibl
tibw
tarl
tarw
type
Bone