Quick Summary of Covariate Data for Lab 05

Here, we only have 2 covariates so summarizing things is fairly simple. There is more one could do but the key is to understand the values that you have, consider whether there might be data errors, and to evaluate relationships between the covariates.

```
# SwampSquirrel data summary
# read data
dat=read.table("http://www.montana.edu/rotella/502/SwampSquirrels.inp",
     header=FALSE, skip=2,
     colClasses=c("character","integer","numeric","numeric","character"))
# remove column of semi-colons
dat=dat[,-5]
# assign column names
colnames(dat) <- c("ch","freq","birth_date","tail_length")</pre>
# summarize the birthday and tail length covariates
summary(dat[,3:4])
##
     birth_date
                 tail_length
## Min. :189 Min. :227
## 1st Qu.:204
                1st Qu.:236
## Median :207
                 Median :243
## Mean :207
                 Mean :240
## 3rd Qu.:212
                 3rd Qu.:245
  Max. :225
##
                Max. :246
# examine correlation between the 2 covariates
cor(dat[,3:4])
##
              birth_date tail_length
## birth_date
              1.000
                               0.347
                 0.347
## tail_length
                               1.000
```

```
par(mfrow = c(2, 1))
hist(dat$birth_date)
abline(v = mean(dat$birth_date), col = "red", lwd = 2)
hist(dat$tail_length)
abline(v = mean(dat$tail_length), col = "blue", lwd = 2)
```

Histogram of dat\$birth_date



Histogram of dat\$tail_length



par(mfrow = c(1, 1))

with(data = dat, plot(birth_date, tail_length))



You often want a table of summary statistics that includes specific statistics of interest that might not be included in the resuls of R's 'summary' function. If you do, you can write a function customized to your liking. Here's an example.

```
# write a function called 'stats'
# provide names for the stats you want to report
stat.names=c("mean","sd","min","max",".025q",".5q",".975")
# record the number of stats you want to report
num.stats=length(stat.names)
stats=function(x){
      y=as.matrix(x)
       cols=dim(y)[2]
       b=matrix(NA,cols,num.stats)
       for (i in 1:cols){
         a=y[,i]
         b[i,]=t(rbind(mean(a),sd(a),min(a),max(a),
                t(t(quantile(a, probs = c(0.025, 0.50, 0.975))))))
      }
row.names(b)=names(x)
colnames(b)=stat.names
     b
}
stats(dat[3:4])
##
                mean
                     sd min max .025q .5q .975
## birth_date 207.2 7.189 189 225
                                     189 207
                                              221
## tail_length 239.9 5.379 227 246
                                     227 243 245
```