

# Chapter 2 R Notebook

This is an R Markdown Notebook. When you execute code within the notebook, the results appear beneath the code.

Try executing this chunk by clicking the *Run* button within the chunk or by placing your cursor inside it and pressing *Ctrl+Shift+Enter*.

Add a new chunk by clicking the *Insert Chunk* button on the toolbar or by pressing *Ctrl+Alt+I*.

When you save the notebook, an HTML file containing the code and output will be saved alongside it (click the *Preview* button or press *Ctrl+Shift+K* to preview the HTML file).

The preview shows you a rendered HTML copy of the contents of the editor. Consequently, unlike *Knit*, *Preview* does not run any R code chunks. Instead, the output of the chunk when it was last run in the editor is displayed.

## Chapter 2

This is an R Notebook with the code from Machine Learning with R, Lantz.

### Blog post about Projects and Notebooks

Prime Hings for Running A Data Project in R

### Good website for learning about R.

Quick-R

## Chapter 2: Managing and Understanding Data

### Libraries

```
library(here)
```

```
## here() starts at /home/esuess/classes/2017-2018/03 - Spring2018/Stat6620/Projects/Chap02
```

```
library(lattice)  
library(corrgram)  
library(gmodels)
```

```
here::here()
```

```
## [1] "/home/esuess/classes/2017-2018/03 - Spring2018/Stat6620/Projects/Chap02"
```

### R data structures

#### Vectors

create vectors of data for three medical patients

```
subject_name <- c("John Doe", "Jane Doe", "Steve Graves")
temperature <- c(98.1, 98.6, 101.4)
flu_status <- c(FALSE, FALSE, TRUE)
```

access the second element in body temperature vector

```
temperature[2]
```

```
## [1] 98.6
```

examples of accessing items in vector

include items in the range 2 to 3

```
temperature[2:3]
```

```
## [1] 98.6 101.4
```

exclude item 2 using the minus sign

```
temperature[-2]
```

```
## [1] 98.1 101.4
```

use a vector to indicate whether to include item

```
temperature[c(TRUE, TRUE, FALSE)]
```

```
## [1] 98.1 98.6
```

## Factors

add gender factor

```
gender <- factor(c("MALE", "FEMALE", "MALE"))
gender
```

```
## [1] MALE FEMALE MALE
```

```
## Levels: FEMALE MALE
```

add blood type factor

```
blood <- factor(c("O", "AB", "A"),
               levels = c("A", "B", "AB", "O"))
blood
```

```
## [1] O AB A
```

```
## Levels: A B AB O
```

add ordered factor

```
symptoms <- factor(c("SEVERE", "MILD", "MODERATE"),
                  levels = c("MILD", "MODERATE", "SEVERE"),
                  ordered = TRUE)
symptoms
```

```
## [1] SEVERE MILD MODERATE
```

```
## Levels: MILD < MODERATE < SEVERE
```

check for symptoms greater than moderate

```
symptoms > "MODERATE"
```

```
## [1] TRUE FALSE FALSE
```

## Lists

display information for a patient

```
subject_name[1]
```

```
## [1] "John Doe"
```

```
temperature[1]
```

```
## [1] 98.1
```

```
flu_status[1]
```

```
## [1] FALSE
```

```
gender[1]
```

```
## [1] MALE
```

```
## Levels: FEMALE MALE
```

```
blood[1]
```

```
## [1] O
```

```
## Levels: A B AB O
```

```
symptoms[1]
```

```
## [1] SEVERE
```

```
## Levels: MILD < MODERATE < SEVERE
```

create list for a patient and display the patient

```
subject1 <- list(fullname = subject_name[1],  
                temperature = temperature[1],  
                flu_status = flu_status[1],  
                gender = gender[1],  
                blood = blood[1],  
                symptoms = symptoms[1])
```

```
subject1
```

```
## $fullname
```

```
## [1] "John Doe"
```

```
##
```

```
## $temperature
```

```
## [1] 98.1
```

```
##
```

```
## $flu_status
```

```
## [1] FALSE
```

```
##
```

```
## $gender
```

```
## [1] MALE
```

```
## Levels: FEMALE MALE
```

```
##
```

```
## $blood
```

```
## [1] 0
## Levels: A B AB 0
##
## $symptoms
## [1] SEVERE
## Levels: MILD < MODERATE < SEVERE

methods for accessing a list
get a single list value by position (returns a sub-list)
```

```
subject1[2]
```

```
## $temperature
## [1] 98.1

get a single list value by position (returns a numeric vector)
```

```
subject1[[2]]
```

```
## [1] 98.1

get a single list value by name
```

```
subject1$temperature
```

```
## [1] 98.1

get several list items by specifying a vector of names
```

```
subject1[c("temperature", "flu_status")]
```

```
## $temperature
## [1] 98.1
##
## $flu_status
## [1] FALSE
```

access a list like a vector get values 2 and 3

```
subject1[2:3]
```

```
## $temperature
## [1] 98.1
##
## $flu_status
## [1] FALSE
```

## Data frames

create a data frame from medical patient data and display the data frame

```
pt_data <- data.frame(subject_name, temperature, flu_status, gender,
                      blood, symptoms, stringsAsFactors = FALSE)
pt_data
```

```
##  subject_name temperature flu_status gender blood symptoms
## 1   John Doe         98.1      FALSE MALE     0   SEVERE
## 2   Jane Doe         98.6      FALSE FEMALE  AB    MILD
## 3  Steve Graves     101.4      TRUE  MALE     A   MODERATE
```

accessing a data frame

get a single column

```
pt_data$subject_name
```

```
## [1] "John Doe"      "Jane Doe"      "Steve Graves"
```

get several columns by specifying a vector of names

```
pt_data[c("temperature", "flu_status")]
```

```
##   temperature flu_status
## 1         98.1     FALSE
## 2         98.6     FALSE
## 3        101.4      TRUE
```

this is the same as above, extracting temperature and flu\_status

```
pt_data[2:3]
```

```
##   temperature flu_status
## 1         98.1     FALSE
## 2         98.6     FALSE
## 3        101.4      TRUE
```

accessing by row and column

```
pt_data[1, 2]
```

```
## [1] 98.1
```

accessing several rows and several columns using vectors

```
pt_data[c(1, 3), c(2, 4)]
```

```
##   temperature gender
## 1         98.1  MALE
## 3        101.4  MALE
```

Leave a row or column blank to extract all rows or columns

```
# column 1, all rows
```

```
pt_data[, 1]
```

```
## [1] "John Doe"      "Jane Doe"      "Steve Graves"
```

```
# row 1, all columns
```

```
pt_data[1, ]
```

```
##   subject_name temperature flu_status gender blood symptoms
## 1   John Doe         98.1     FALSE  MALE    0   SEVERE
```

```
# all rows and all columns
```

```
pt_data[ , ]
```

```
##   subject_name temperature flu_status gender blood symptoms
## 1   John Doe         98.1     FALSE  MALE    0   SEVERE
## 2   Jane Doe         98.6     FALSE FEMALE  AB    MILD
## 3 Steve Graves      101.4      TRUE   MALE    A MODERATE
```

the following are equivalent

```
pt_data[c(1, 3), c("temperature", "gender")]
```

```
## temperature gender  
## 1 98.1 MALE  
## 3 101.4 MALE
```

```
pt_data[-2, c(-1, -3, -5, -6)]
```

```
## temperature gender  
## 1 98.1 MALE  
## 3 101.4 MALE
```

## Matrixes

create a 2x2 matrix

```
m <- matrix(c(1, 2, 3, 4), nrow = 2)
```

```
m
```

```
##      [,1] [,2]  
## [1,] 1   3  
## [2,] 2   4
```

equivalent to the above

```
m <- matrix(c(1, 2, 3, 4), ncol = 2)
```

```
m
```

```
##      [,1] [,2]  
## [1,] 1   3  
## [2,] 2   4
```

create a 2x3 matrix

```
m <- matrix(c(1, 2, 3, 4, 5, 6), nrow = 2)
```

```
m
```

```
##      [,1] [,2] [,3]  
## [1,] 1   3   5  
## [2,] 2   4   6
```

create a 3x2 matrix

```
m <- matrix(c(1, 2, 3, 4, 5, 6), ncol = 2)
```

```
m
```

```
##      [,1] [,2]  
## [1,] 1   4  
## [2,] 2   5  
## [3,] 3   6
```

extract values from matrixes

```
m[1, 1]
```

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

```
m[3, 2]
```

```
## [1] 6
```

extract rows

```
m[1, ]
```

```
## [1] 1 4
```

extract columns

```
m[, 1]
```

```
## [1] 1 2 3
```

## Managing data with R

saving, loading, and removing R data structures

show all data structures in memory

```
ls()
```

```
## [1] "blood"          "flu_status"    "gender"        "m"  
## [5] "pt_data"         "subject_name" "subject1"      "symptoms"  
## [9] "temperature"
```

remove the m and subject1 objects

```
rm(m, subject1)
```

```
ls()
```

```
## [1] "blood"          "flu_status"    "gender"        "pt_data"  
## [5] "subject_name"  "symptoms"      "temperature"
```

```
rm(list=ls())
```

## Exploring and understanding data

data exploration example using used car data

```
usedcars <- read.csv("usedcars.csv", stringsAsFactors = FALSE)
```

get structure of used car data

```
str(usedcars)
```

```
## 'data.frame': 150 obs. of 6 variables:  
## $ year : int 2011 2011 2011 2011 2012 2010 2011 2010 2011 2010 ...  
## $ model : chr "SEL" "SEL" "SEL" "SEL" ...  
## $ price : int 21992 20995 19995 17809 17500 17495 17000 16995 16995 16995 ...  
## $ mileage : int 7413 10926 7351 11613 8367 25125 27393 21026 32655 36116 ...  
## $ color : chr "Yellow" "Gray" "Silver" "Gray" ...  
## $ transmission: chr "AUTO" "AUTO" "AUTO" "AUTO" ...
```

## Exploring numeric variables

summarize numeric variables

```
summary(usedcars$year)
```

```
##   Min. 1st Qu.  Median    Mean 3rd Qu.  Max.
##   2000   2008   2009     2009   2010   2012
```

```
summary(usedcars[c("price", "mileage")])
```

```
##      price      mileage
## Min.   : 3800   Min.   : 4867
## 1st Qu.:10995   1st Qu.: 27200
## Median :13592   Median : 36385
## Mean   :12962   Mean   : 44261
## 3rd Qu.:14904   3rd Qu.: 55124
## Max.   :21992   Max.   :151479
```

calculate the mean income

```
(36000 + 44000 + 56000) / 3
```

```
## [1] 45333.33
```

```
mean(c(36000, 44000, 56000))
```

```
## [1] 45333.33
```

the median income

```
median(c(36000, 44000, 56000))
```

```
## [1] 44000
```

the min/max of used car prices

```
range(usedcars$price)
```

```
## [1] 3800 21992
```

the difference of the range

```
diff(range(usedcars$price))
```

```
## [1] 18192
```

IQR for used car prices

```
IQR(usedcars$price)
```

```
## [1] 3909.5
```

use quantile to calculate five-number summary

```
quantile(usedcars$price)
```

```
##      0%      25%      50%      75%     100%
## 3800.0 10995.0 13591.5 14904.5 21992.0
```

the 99th percentile

```
quantile(usedcars$price, probs = c(0.01, 0.99))
```

```
##      1%      99%
## 5428.69 20505.00
```

quintiles

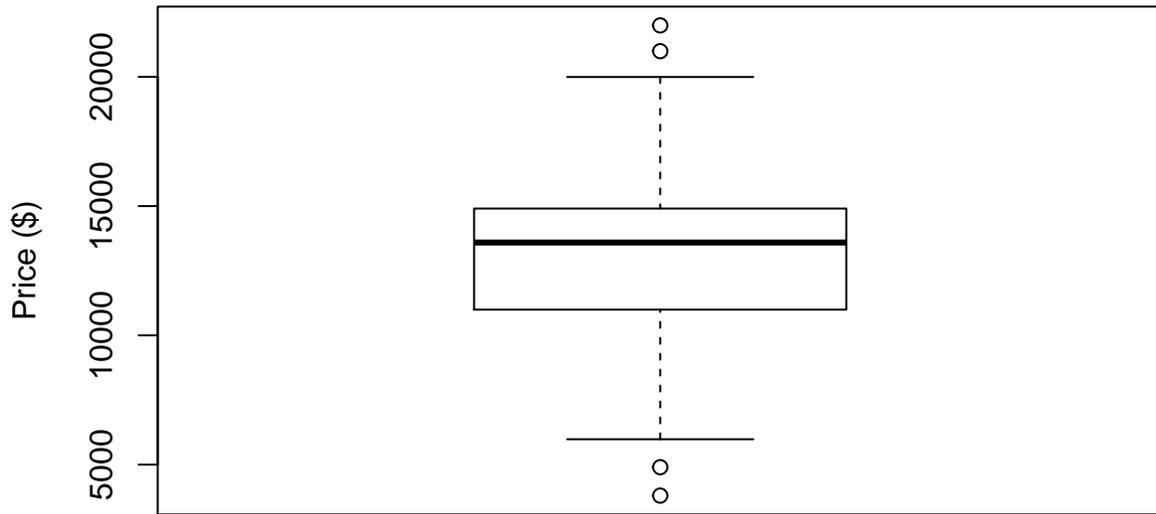
```
quantile(usedcars$price, seq(from = 0, to = 1, by = 0.20))
```

```
##      0%      20%      40%      60%      80%     100%
## 3800.0 10759.4 12993.8 13992.0 14999.0 21992.0
```

boxplot of used car prices and mileage

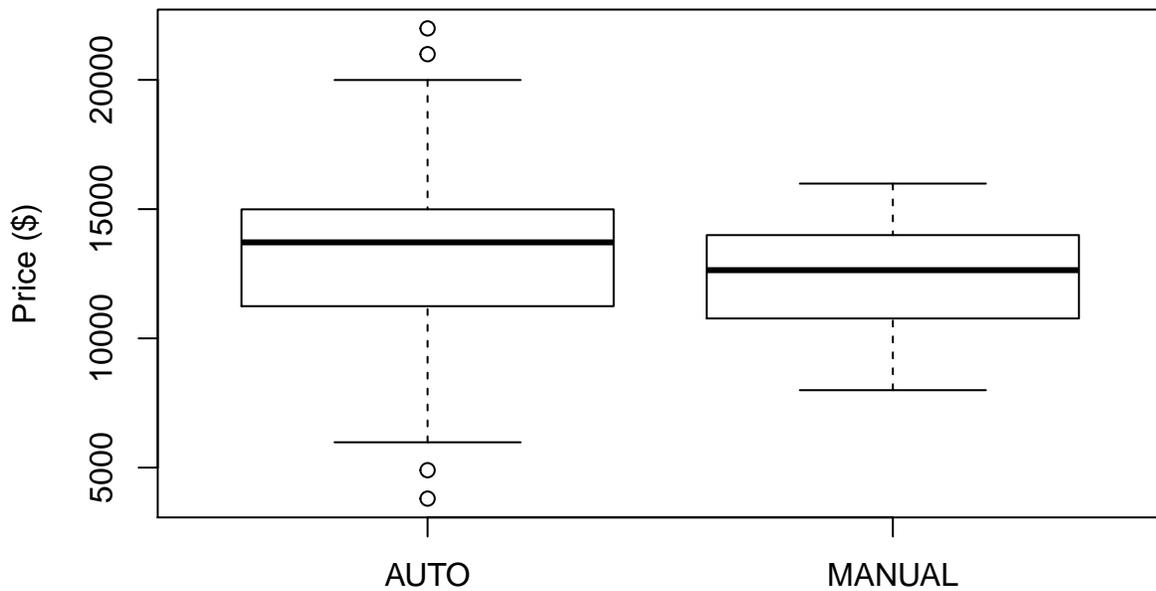
```
boxplot(usedcars$price, main="Boxplot of Used Car Prices",
        ylab="Price ($)")
```

### Boxplot of Used Car Prices



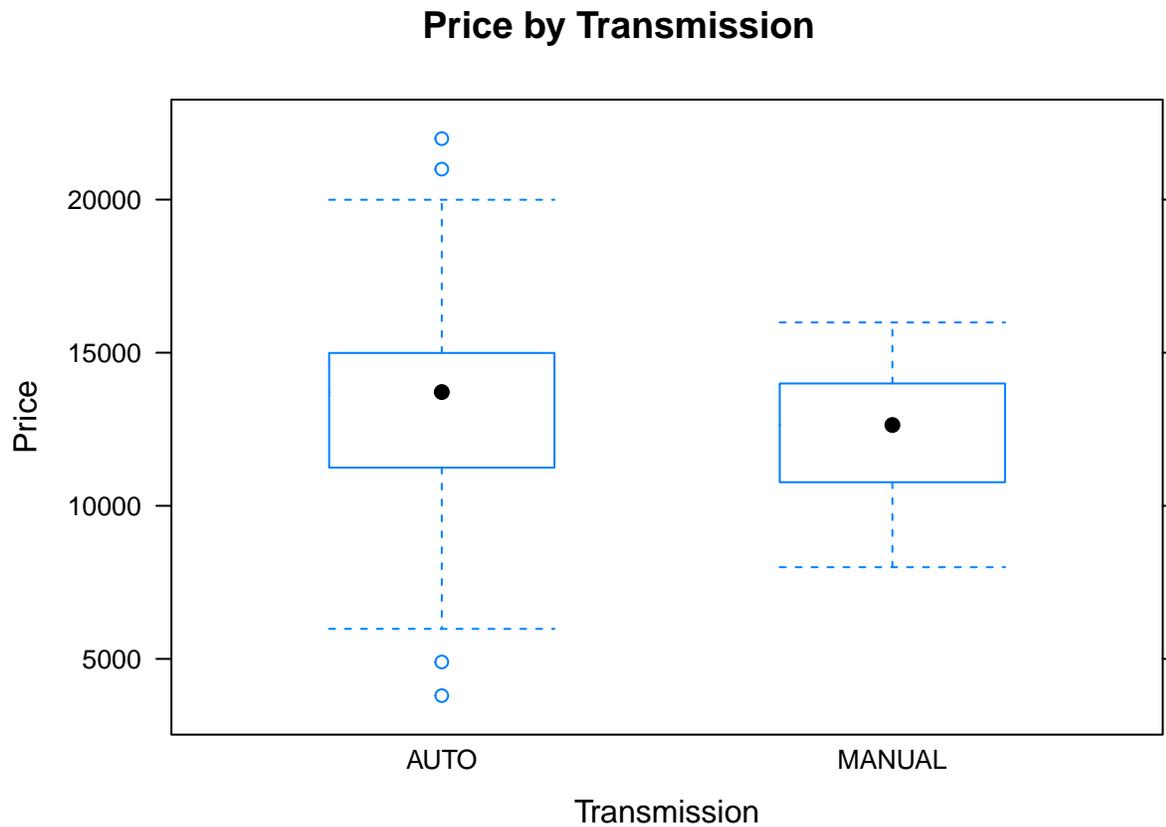
```
boxplot(usedcars$price ~ usedcars$transmission, main="Boxplot of Used Car Prices by Transmission",
        ylab="Price ($)")
```

### Boxplot of Used Car Prices by Transmission



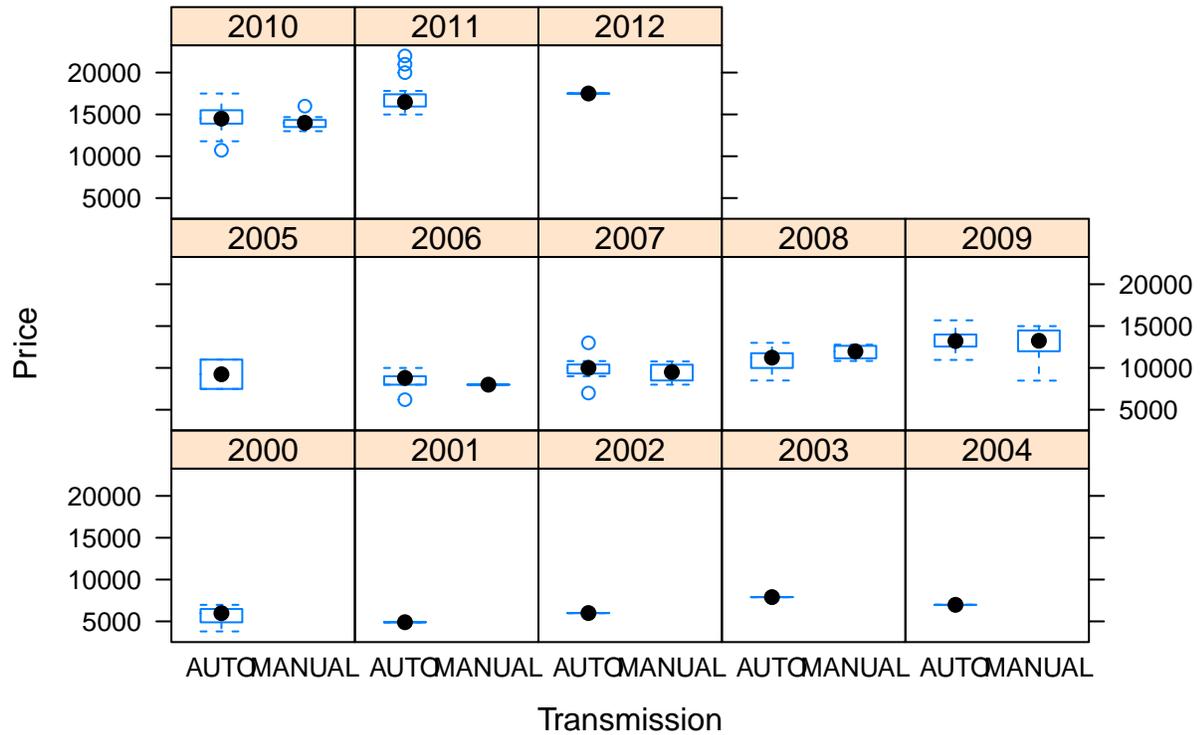
using the lattice package

```
lattice::bwplot(usedcars$price~usedcars$transmission,  
  ylab="Price", xlab="Transmission",  
  main="Price by Transmission")
```



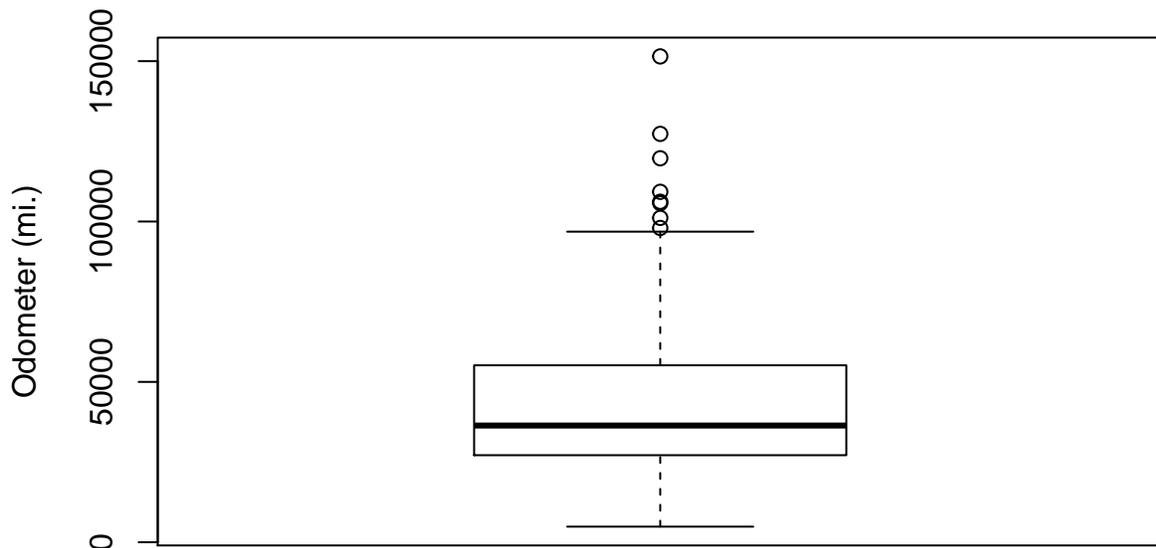
```
usedcars$year <- as.character(usedcars$year)  
  
lattice::bwplot(usedcars$price~usedcars$transmission|usedcars$year,  
  ylab="Price", xlab="Transmission",  
  main="Price by Transmission and Year", layout=(c(5,3)))
```

## Price by Transmission and Year



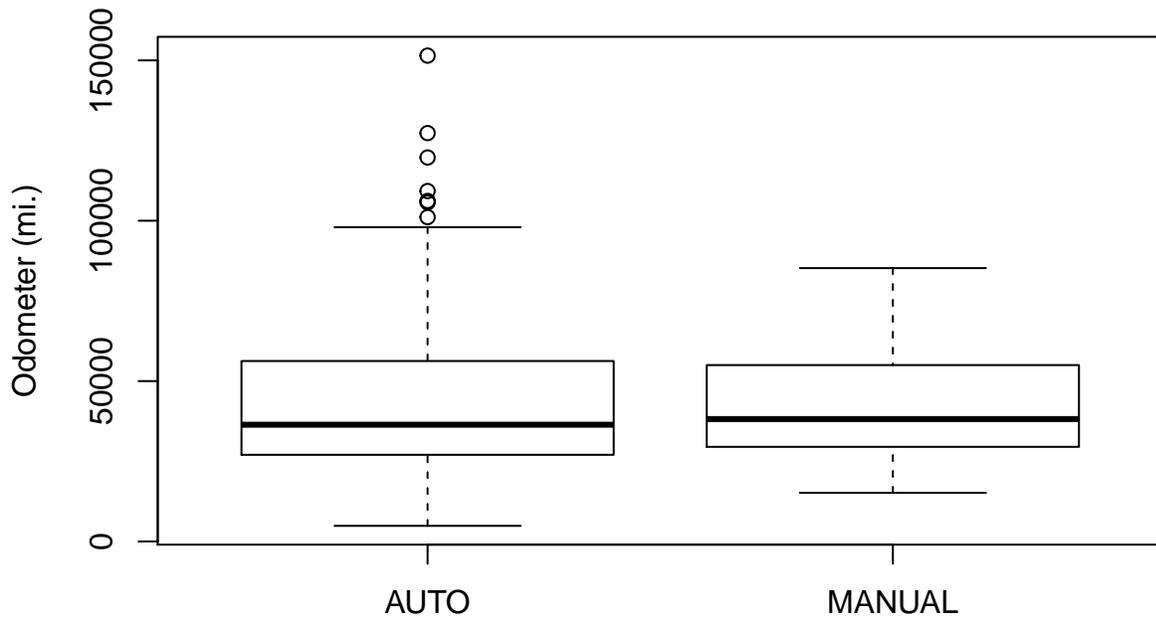
```
boxplot(usedcars$mileage, main="Boxplot of Used Car Mileage",
        ylab="Odometer (mi.)")
```

## Boxplot of Used Car Mileage



```
boxplot(usedcars$mileage ~ usedcars$transmission, main="Boxplot of Used Car Mileage by Transmission", y
```

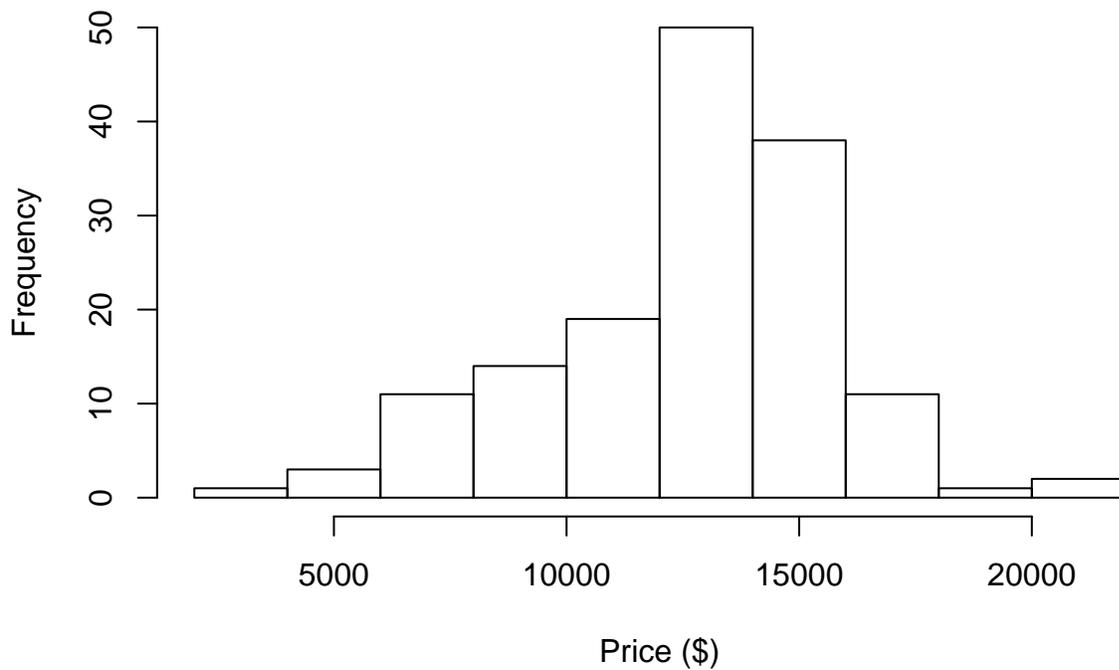
## Boxplot of Used Car Mileage by Transmission



histograms of used car prices and mileage

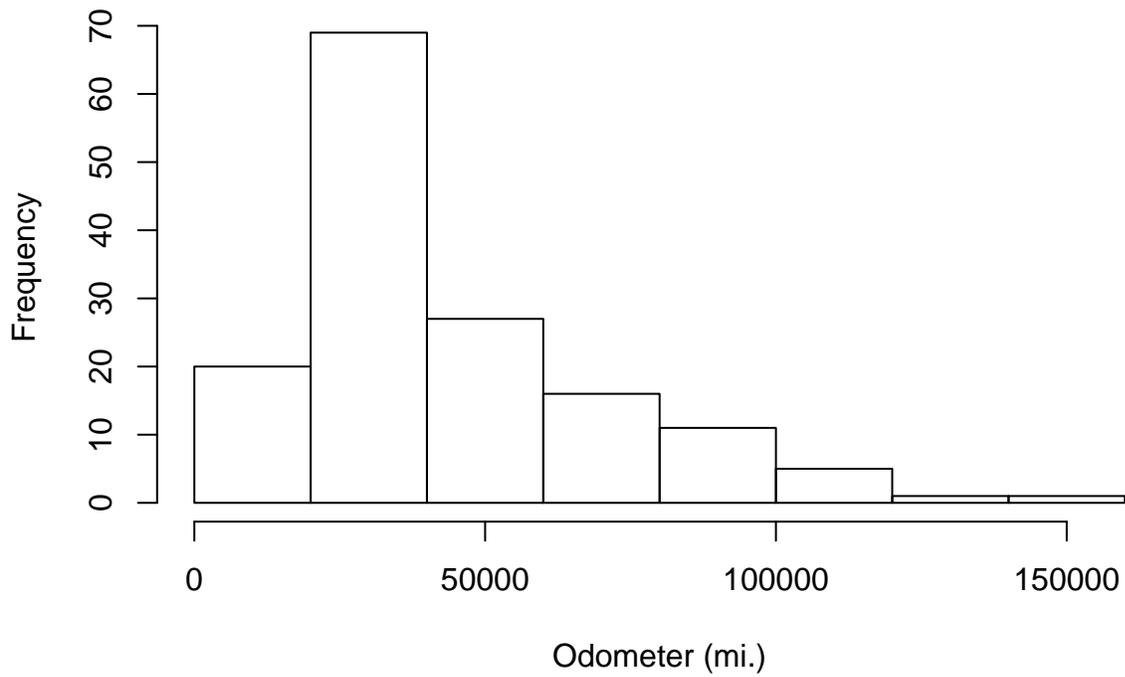
```
hist(usedcars$price, main = "Histogram of Used Car Prices",  
     xlab = "Price ($)")
```

## Histogram of Used Car Prices



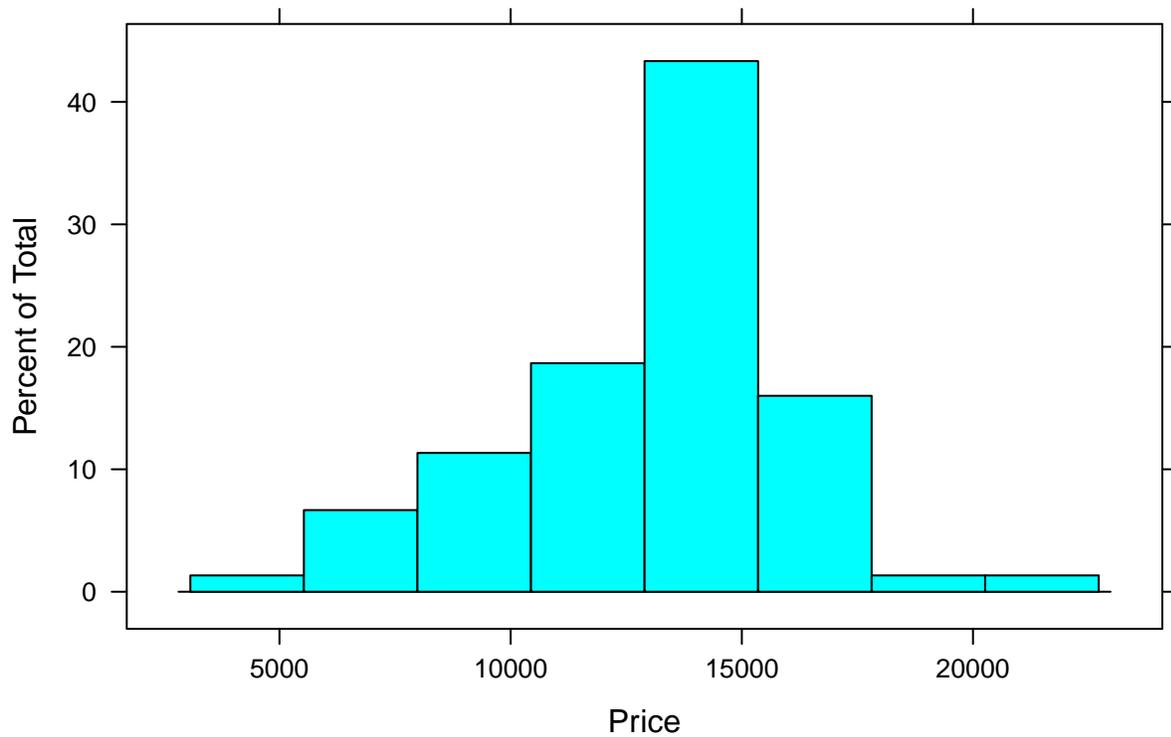
```
hist(usedcars$mileage, main = "Histogram of Used Car Mileage",  
     xlab = "Odometer (mi.)")
```

## Histogram of Used Car Mileage



```
lattice::histogram(~ usedcars$price,  
  xlab="Price",  
  main="Distribution of Price")
```

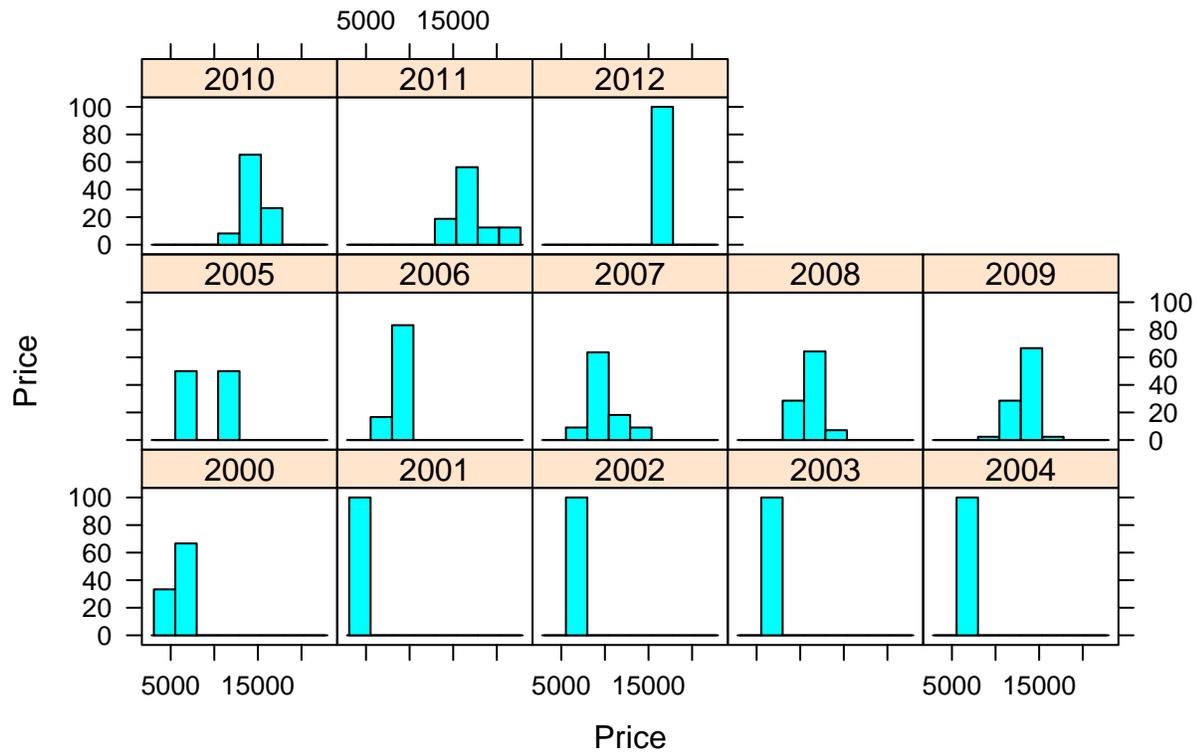
## Distribution of Price



```
usedcars$year <- as.character(usedcars$year)

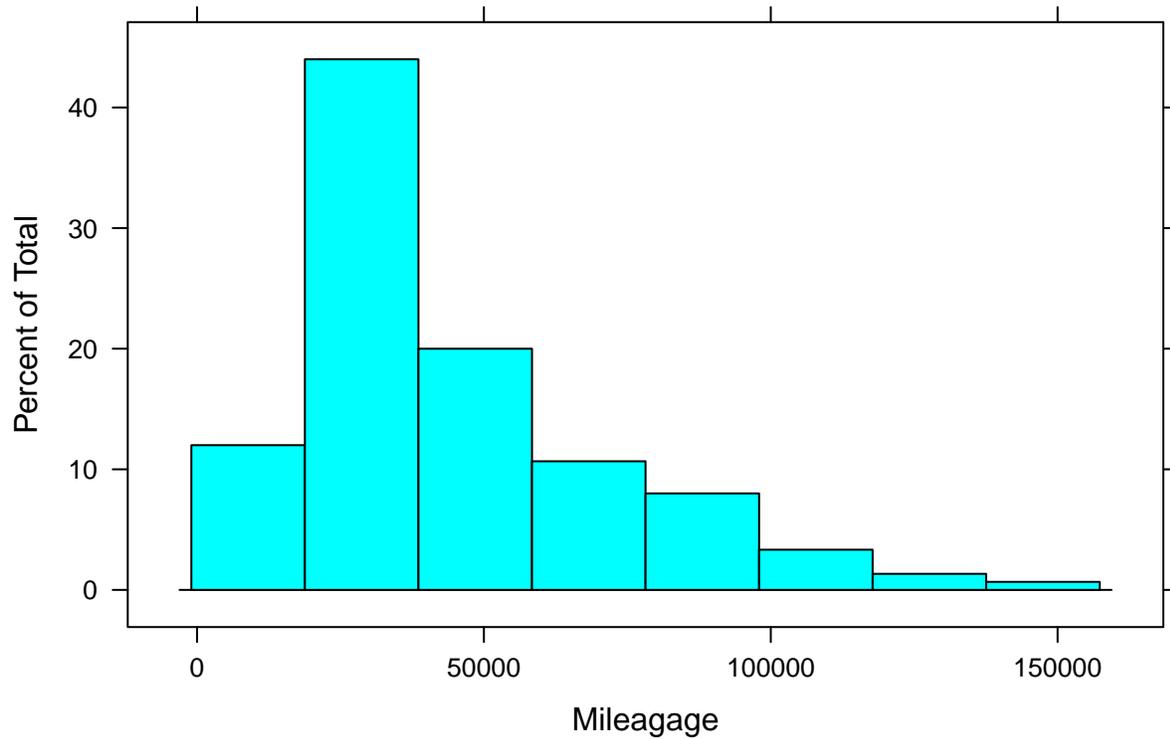
lattice::histogram(~ usedcars$price | usedcars$year,
  ylab="Price", xlab="Price",
  main="Distribution of Price by Year", layout=(c(5,3)))
```

### Distribution of Price by Year



```
lattice::histogram(~ usedcars$mileage,
  xlab="Mileage",
  main="Distribution of Mileage")
```

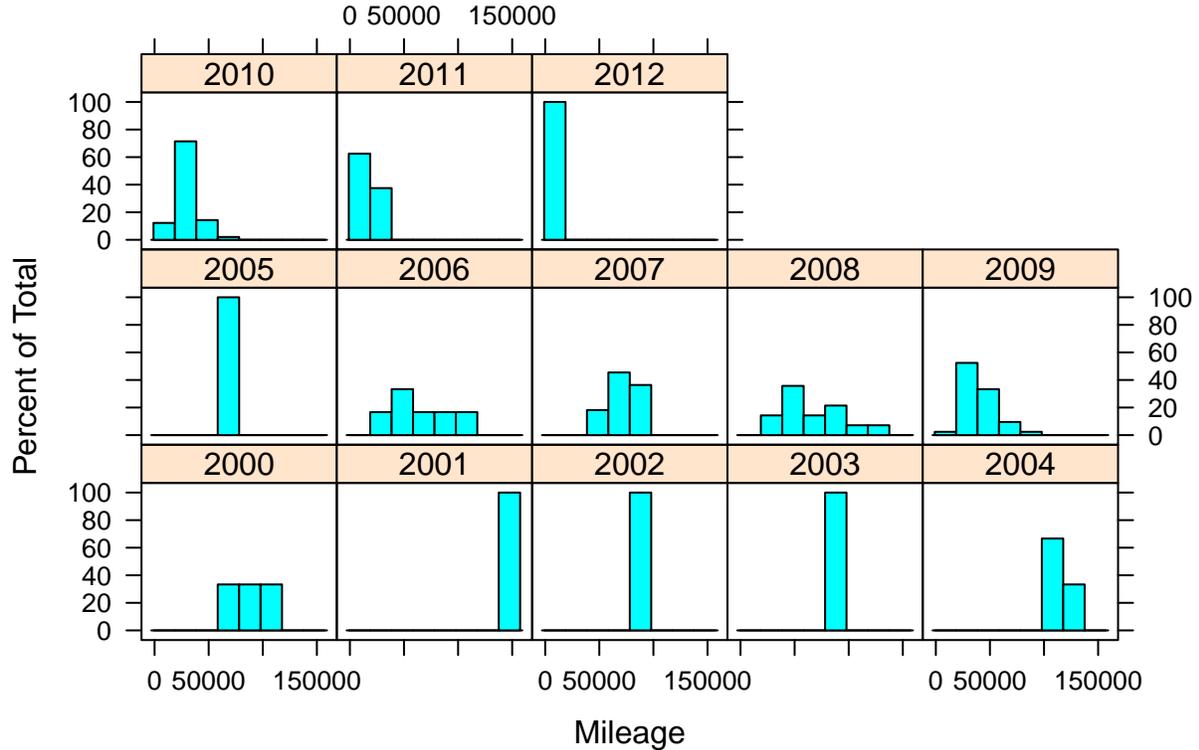
## Distribution of Mileage



```
usedcars$year <- as.character(usedcars$year)

lattice::histogram(~ usedcars$mileage | usedcars$year,
  xlab="Mileage",
  main="Distribution of Mileage by Year", layout=(c(5,3)))
```

## Distribution of Mileage by Year



variance and standard deviation of the used car data

```
var(usedcars$price)
```

```
## [1] 9749892
```

```
sd(usedcars$price)
```

```
## [1] 3122.482
```

```
var(usedcars$mileage)
```

```
## [1] 728033954
```

```
sd(usedcars$mileage)
```

```
## [1] 26982.1
```

## Exploring numeric variables

one-way tables for the used car data

```
table(usedcars$year)
```

```
##
```

```
## 2000 2001 2002 2003 2004 2005 2006 2007 2008 2009 2010 2011 2012
```

```
## 3 1 1 1 3 2 6 11 14 42 49 16 1
```

```
table(usedcars$model)
```

```
##
```

```
## SE SEL SES
## 78 23 49
table(usedcars$color)

##
## Black Blue Gold Gray Green Red Silver White Yellow
## 35 17 1 16 5 25 32 16 3

compute table proportions

model_table <- table(usedcars$model)
prop.table(model_table)

##
## SE SEL SES
## 0.5200000 0.1533333 0.3266667

round the data

color_table <- table(usedcars$color)
color_pct <- prop.table(color_table) * 100
round(color_pct, digits = 1)

##
## Black Blue Gold Gray Green Red Silver White Yellow
## 23.3 11.3 0.7 10.7 3.3 16.7 21.3 10.7 2.0
```

## Exploring relationships between variables

correlation

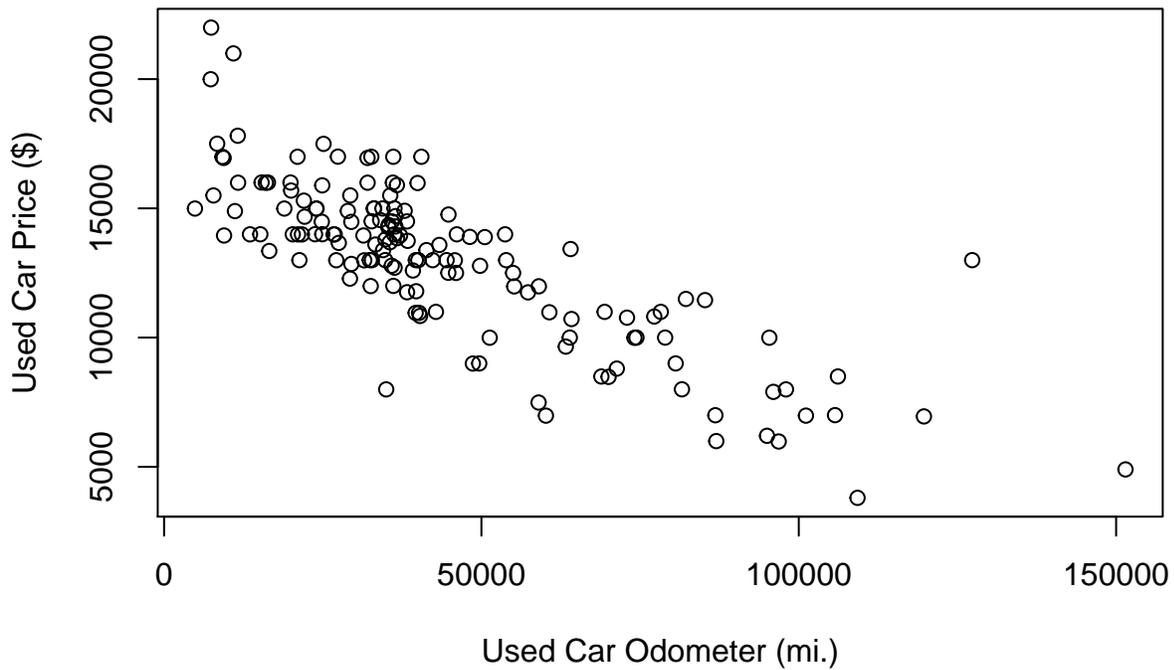
```
cor(x = usedcars$mileage, y = usedcars$price)
```

```
## [1] -0.8061494
```

scatterplot of price vs. mileage

```
plot(x = usedcars$mileage, y = usedcars$price,
     main = "Scatterplot of Price vs. Mileage",
     xlab = "Used Car Odometer (mi.)",
     ylab = "Used Car Price ($)")
```

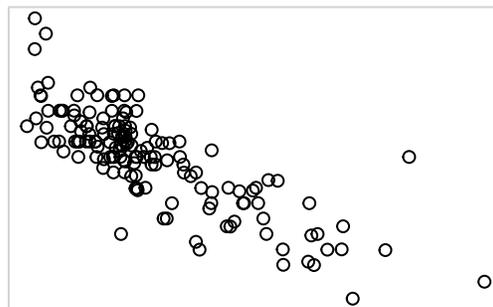
## Scatterplot of Price vs. Mileage



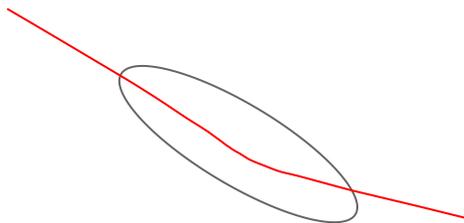
The corrgram package has the corrgram function that is nice for looking at relationships between numeric variable.

```
corrgram::corrgram(usedcars, lower.panel=panel.ellipse,  
  upper.panel=panel.pts)
```

price



mileage



new variable indicating conservative colors

```
usedcars$conservative <-  
  usedcars$color %in% c("Black", "Gray", "Silver", "White")
```

checking our variable

```
table(usedcars$conservative)
```

```
##  
## FALSE TRUE  
##    51    99
```

Crosstab of conservative by model

```
gmodels::CrossTable(x = usedcars$model, y = usedcars$conservative)
```

```
##  
##  
## Cell Contents  
## |-----|  
## |                N |  
## | Chi-square contribution |  
## |      N / Row Total |  
## |      N / Col Total |  
## |      N / Table Total |  
## |-----|  
##  
##  
## Total Observations in Table: 150  
##  
##  
##          | usedcars$conservative  
## usedcars$model |      FALSE |      TRUE | Row Total |  
## -----|-----|-----|-----|  
##          SE |          27 |          51 |          78 |  
##          |          0.009 |          0.004 |          |  
##          |          0.346 |          0.654 |          0.520 |  
##          |          0.529 |          0.515 |          |  
##          |          0.180 |          0.340 |          |  
## -----|-----|-----|-----|  
##          SEL |           7 |          16 |          23 |  
##          |          0.086 |          0.044 |          |  
##          |          0.304 |          0.696 |          0.153 |  
##          |          0.137 |          0.162 |          |  
##          |          0.047 |          0.107 |          |  
## -----|-----|-----|-----|  
##          SES |          17 |          32 |          49 |  
##          |          0.007 |          0.004 |          |  
##          |          0.347 |          0.653 |          0.327 |  
##          |          0.333 |          0.323 |          |  
##          |          0.113 |          0.213 |          |  
## -----|-----|-----|-----|  
## Column Total |          51 |          99 |          150 |  
##          |          0.340 |          0.660 |          |  
## -----|-----|-----|-----|  
##  
##
```