

### Chapter 1

- $Y_i = \beta_0 + \beta_1 X_i + \varepsilon_i$  where  $\varepsilon_i \sim$  independent  $N(0, \sigma^2)$
- $\hat{Y}_i = b_0 + b_1 X_i$  where  $b_1 = \frac{\sum_{i=1}^n (X_i - \bar{X})(Y_i - \bar{Y})}{\sum_{i=1}^n (X_i - \bar{X})^2} = \frac{\sum_{i=1}^n X_i Y_i - \frac{1}{n} \left( \sum_{i=1}^n X_i \right) \left( \sum_{i=1}^n Y_i \right)}{\left( \sum_{i=1}^n X_i^2 \right) - \frac{1}{n} \left( \sum_{i=1}^n X_i \right)^2}$  and  $b_0 = \bar{Y} - b_1 \bar{X}$
- $e_i = Y_i - \hat{Y}_i$
- $SSE = \sum_{i=1}^n (Y_i - \hat{Y}_i)^2$
- $E(W_1 + c) = E(W_1) + c$
- $E(cW_1) = c \cdot E(W_1)$
- $E(bW_1 + c) = b \cdot E(W_1) + c$
- $E(W_1 + W_2) = E(W_1) + E(W_2)$
- $E(W_1 \cdot W_2) \neq E(W_1) \cdot E(W_2)$  – equality occurs when  $W_1$  and  $W_2$  are independent
- $MSE = \frac{SSE}{n-2}$

### Chapter 2

- $b_1 \sim N(\beta_1, \sigma^2 / \sum (X_i - \bar{X})^2)$
- $\text{Var}(\sum a_i Y_i) = \sum a_i^2 \text{Var}(Y_i)$
- $\hat{\text{Var}}(b_1) = MSE / \sum_{i=1}^n (X_i - \bar{X})^2$
- $t^* = (b_1 - \beta_1) / \sqrt{\hat{\text{Var}}(b_1)}$  and use a  $t(n-2)$  distribution
- $b_1 \pm t(1 - \alpha / 2; n - 2) \cdot \sqrt{\hat{\text{Var}}(b_1)}$
- $b_0 \sim N\left(\beta_0, \sigma^2 \left( \frac{1}{n} + \frac{\bar{X}^2}{\sum (X_i - \bar{X})^2} \right)\right)$
- $\hat{Y}_h \pm t(1 - \alpha / 2; n - 2) \sqrt{\hat{\text{Var}}(\hat{Y}_h)}$  where  $\hat{\text{Var}}(\hat{Y}_h) = MSE \left( \frac{1}{n} + \frac{(X_h - \bar{X})^2}{\sum (X_i - \bar{X})^2} \right)$
- $\hat{Y}_h \pm t(1 - \alpha / 2; n - 2) \sqrt{\hat{\text{Var}}(Y_{h(\text{new})} - \hat{Y}_h)}$  where  $\hat{\text{Var}}(Y_{h(\text{new})} - \hat{Y}_h) = MSE \left( 1 + \frac{1}{n} + \frac{(X_h - \bar{X})^2}{\sum (X_i - \bar{X})^2} \right)$
- $\hat{Y}_h \pm t(1 - \alpha / 2; n - 2) \sqrt{\hat{\text{Var}}(\bar{Y}_{h(\text{new})} - \hat{Y}_h)}$  for  $\bar{Y}_{h(\text{new})} = \frac{1}{m} \sum_{j=1}^m Y_{j,h(\text{new})}$  where  

$$\hat{\text{Var}}(\bar{Y}_{h(\text{new})} - \hat{Y}_h) = MSE \left( \frac{1}{m} + \frac{1}{n} + \frac{(X_h - \bar{X})^2}{\sum (X_i - \bar{X})^2} \right)$$
- $SSTO = \sum_{i=1}^n (Y_i - \bar{Y})^2$

- $SSR = \sum_{i=1}^n (\hat{Y}_i - \bar{Y})^2$
- $SSTO = SSE + SSR$
- $MSR = SSR/1 = SSR$  when there is  $\beta_0$  and  $\beta_1$  in the regression model
- $F^* = \frac{MSR}{MSE}$  and use a  $F(1, n-2)$  distribution
- $R^2 = \frac{SSR}{SSTO} = \frac{SSTO - SSE}{SSTO}$

### Chapter 3

- $e_i^* = \frac{e_i}{\sqrt{MSE}}$
- $Y_i^{(\lambda)} = \beta_0 + \beta_1 X_i + \varepsilon_i$  where  $Y^{(\lambda)} = \begin{cases} Y^\lambda & \lambda \neq 0 \\ \log_e(Y) & \lambda = 0 \end{cases}$
- $d_{i1} = |e_{i1} - \tilde{e}_1|$  and  $d_{i2} = |e_{i2} - \tilde{e}_2|$ , median residual values are  $\tilde{e}_1$  and  $\tilde{e}_2$ ,  $t_i^* = \frac{\bar{d}_1 - \bar{d}_2}{s \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}}$  and use a  $t(n-2)$  distribution,  $s^2 = \frac{\sum (d_{i1} - \bar{d}_1)^2 + \sum (d_{i2} - \bar{d}_2)^2}{n-2}$
- $\log(\sigma_i^2) = \gamma_0 + \gamma_1 X_i$ ,  $X_{BP}^2 = \frac{SSR^*/2}{(SSE/n)^2}$  and use a  $\chi^2$  distribution with 1 degree of freedom

### Chapter 4

- $b_0 \pm t(1 - \alpha / 4; n - 2) * \sqrt{\hat{Var}(b_0)}$ ,  $b_1 \pm t(1 - \alpha / 4; n - 2) * \sqrt{\hat{Var}(b_1)}$
- $\hat{Y}_h \pm t\left(1 - \frac{\alpha}{2g}; n - 2\right) \sqrt{\hat{Var}(\hat{Y}_h)}$
- $\hat{Y}_h \pm t\left(1 - \frac{\alpha}{2g}; n - 2\right) \sqrt{\hat{Var}(\hat{Y}_{h(new)})}$
- $\hat{X}_{h(new)} = \frac{Y_{h(new)} - b_0}{b_1}$
- $\hat{X}_{h(new)} \pm t(1 - \alpha / 2, n - 2) \sqrt{\hat{Var}(\hat{X}_{h(new)})}$  where  $\hat{Var}(\hat{X}_{h(new)}) = \frac{MSE}{b_1^2} \left[ 1 + \frac{1}{n} + \frac{(\hat{X}_{h(new)} - \bar{X})^2}{\sum (X_i - \bar{X})^2} \right]$

### Chapter 5

- $Cov(Z_1, Z_2) = E[(Z_1 - \mu_1)(Z_2 - \mu_2)]$
- $Corr(Z_1, Z_2) = \frac{Cov(Z_1, Z_2)}{\sqrt{Var(Z_1)}\sqrt{Var(Z_2)}}$  and  $Cov(\mathbf{Z}) = \begin{bmatrix} Var(Z_1) & Cov(Z_1, Z_2) \\ Cov(Z_1, Z_2) & Var(Z_2) \end{bmatrix}$
- $\mathbf{Y} = E(\mathbf{Y}) + \varepsilon$  where  $E(\mathbf{Y}) = \mathbf{X}\beta$

- Let  $\mathbf{W} = \mathbf{A}\mathbf{Y}$  where  $\mathbf{Y}$  is a random vector and  $\mathbf{A}$  is a matrix of constants. Then  $E(\mathbf{A}) = \mathbf{A}$ ,  $E(\mathbf{W}) = \mathbf{A}E(\mathbf{Y})$ , and  $\text{Cov}(\mathbf{W}) = \mathbf{A}\text{Cov}(\mathbf{Y})\mathbf{A}'$
- $\mathbf{b} = (\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'\mathbf{Y}$
- $\hat{\mathbf{Y}} = \mathbf{X}\mathbf{b} = \mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'\mathbf{Y} = \mathbf{H}\mathbf{Y}$  where  $\mathbf{H} = \mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'$
- $\mathbf{e} = \mathbf{Y}(\mathbf{I} - \mathbf{H})$
- $\text{SSTO} = \mathbf{Y}'\mathbf{Y} - \frac{1}{n}\mathbf{Y}'\mathbf{J}\mathbf{Y}$ ,  $\text{SSE} = \mathbf{e}'\mathbf{e}$ ,  $\text{SSR} = \mathbf{b}'\mathbf{X}'\mathbf{Y} - \frac{1}{n}\mathbf{Y}'\mathbf{J}\mathbf{Y}$
- $\text{Cov}(\mathbf{b}) = \sigma^2(\mathbf{X}'\mathbf{X})^{-1}$
- $\widehat{\text{Var}}(\hat{Y}_h) = \text{MSE}(\mathbf{X}_h'(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}_h)$
- $\widehat{\text{Var}}(Y_{h(\text{new})} - \hat{Y}_h) = \text{MSE}(1 + \mathbf{X}_h'(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}_h)$

### Chapter 6

- $t^* = \frac{b_k - \beta_k}{\sqrt{\widehat{\text{Var}}(b_k)}}$  and use a  $t(n-p)$  distribution
- $\text{MSR} = \text{SSR}/(p-1)$
- $\text{MSE} = \text{SSE}/(n-p)$
- $F^* = \text{MSR}/\text{MSE}$  and use a  $F(p-1, n-p)$  distribution
- $R_a^2 = 1 - \frac{n-1}{n-p}(1-R^2)$

### Chapter 7

- $\text{SSR}(X_2|X_1) = \text{SSE}(X_1) - \text{SSE}(X_1, X_2)$
- $F^* = \frac{(\text{SSE}(X_1, \dots, X_g) - \text{SSE}(X_1, \dots, X_g, X_{g+1}, \dots, X_{p-1})) / (p-1-g)}{\text{SSE}(X_1, \dots, X_g, X_{g+1}, \dots, X_{p-1}) / (n-p)} = \frac{\text{MSR}(X_{g+1}, \dots, X_{p-1} | X_1, \dots, X_g)}{\text{MSE}(X_1, \dots, X_g, X_{g+1}, \dots, X_{p-1})}$  and use a  $F(p-1-g, n-p)$  distribution
- $R_{Y_3|12}^2 = \frac{\text{SSR}(X_3 | X_1, X_2)}{\text{SSE}(X_1, X_2)} = \frac{\text{SSE}(X_1, X_2) - \text{SSE}(X_1, X_2, X_3)}{\text{SSE}(X_1, X_2)}$

### Chapter 8

- Second order model for two predictor variables:  $E(Y_i) = \beta_0 + \beta_1 Z_{i1} + \beta_2 Z_{i2} + \beta_3 Z_{i1} Z_{i2} + \beta_4 Z_{i1}^2 + \beta_5 Z_{i2}^2$

### Chapter 9

- $C_p = \frac{\text{SSE}(X_1, \dots, X_{p-1})}{\text{MSE}(X_1, \dots, X_{p-1})} - (n - 2p)$
- $\text{AIC}_p = n \cdot \log(\text{SSE}_p) - n \cdot \log(n) + 2p$
- $\text{SBC}_p = n \cdot \log(\text{SSE}_p) - n \cdot \log(n) + \log(n) \cdot p$
- $\text{PRESS}_p = \sum_{i=1}^n (Y_i - \hat{Y}_{i(i)})^2 = \sum_{i=1}^n \frac{(Y_i - \hat{Y}_i)^2}{(1 - h_i)^2}$

## Chapter 10

- $r_i = \frac{e_i}{\sqrt{\widehat{\text{Var}}(e_i)}} = \frac{e_i}{\sqrt{\text{MSE}(1-h_{ii})}}$  and  $r_i \sim t(n-p)$
- $d_i = Y_i - \hat{Y}_{i(i)}$
- $t_i = \frac{d_i}{\sqrt{\text{MSE}_{(i)}(1 + \mathbf{X}_i'(\mathbf{X}_{(i)}\mathbf{X}_{(i)})^{-1}\mathbf{X}_i)}} = e_i \left[ \frac{n-p-1}{\text{SSE}(1-h_{ii}) - e_i^2} \right]^{1/2}$  and  $t_i \sim t(n-p-1)$
- Hat matrix diagonal values criteria:  $h_{ii} > 2p/n$ ;  $h_{ii} > 0.5$  indicates very high,  $0.2 < h_{ii} \leq 0.5$  indicate moderate
- $(\text{DFFITS})_i = \frac{\hat{Y}_i - \hat{Y}_{i(i)}}{\sqrt{\text{MSE}_{(i)}h_{ii}}} = t_i \left( \frac{h_{ii}}{1-h_{ii}} \right)^{1/2}$  with  $|(\text{DFFITS})_i| > 1$  for “small to medium” sized data sets and  $|(\text{DFFITS})_i| > 2\sqrt{p/n}$  for “large” data sets as a criteria
- $D_i = \frac{\sum_{j=1}^n (\hat{Y}_j - \hat{Y}_{j(i)})^2}{p\text{MSE}} = \frac{e_i^2}{p\text{MSE}} \left[ \frac{h_{ii}}{(1-h_{ii})^2} \right]$  and  $D_i > F(0.50, p, n-p)$  as a criteria
- $(\text{DFBETAS})_{k(i)} = \frac{b_k - b_{k(i)}}{\sqrt{\text{MSE}_{(i)}c_{kk}}}$  with  $|(\text{DFBETAS})_{k(i)}| > 1$  for “small to medium” sized data sets and  $|(\text{DFBETAS})_{k(i)}| > 2/\sqrt{n}$  for “large” data sets as a criteria
- $(\text{VIF})_k = (1-R_k^2)^{-1}$  with  $(\text{VIF})_k > 10$  as a criteria

## Chapter 11

- $\mathbf{b}_w = (\mathbf{X}'\mathbf{W}\mathbf{X})^{-1}\mathbf{X}'\mathbf{W}\mathbf{Y}$  with  $\mathbf{W} = \begin{bmatrix} w_1 & 0 & \dots & 0 \\ 0 & w_2 & \dots & 0 \\ \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & \dots & w_n \end{bmatrix} = \begin{bmatrix} 1/\sigma_1^2 & 0 & \dots & 0 \\ 0 & 1/\sigma_2^2 & \dots & 0 \\ \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & \dots & 1/\sigma_n^2 \end{bmatrix}$
- $\text{Cov}(\mathbf{b}_w) = (\mathbf{X}'\mathbf{W}\mathbf{X})^{-1}$
- $d_i = \left[ \frac{(X_{i1} - X_{h1})^2}{S_1} + \frac{(X_{i2} - X_{h2})^2}{S_2} \right]^{1/2}$  for two predictor variables
- $w_i = \begin{cases} [1 - (d_i / d_q)^3]^3 & d_i < d_q \\ 0 & d_i \geq d_q \end{cases}$

## Test #1 R part:

Syntax	Description
<code>lm(formula = y ~ x, data = data.frame)</code>	Find the sample regression model (and various other measures) with the response variable y and predictor variable x
<code>summary(mod.fit)</code>	Summarizes information in <i>mod.fit</i>
<code>predict(object = mod.fit, newdata = data.frame, interval = "confidence", se.fit = TRUE, level = 1-alpha)</code>	Predicts the response variable at the predictor variable values in the <i>data.frame</i> using model information in <i>mod.fit</i> . The <i>data.frame</i> needs to have the same predictor variable as the original data set specified in <i>mod.fit</i> . The standard errors produced are $\sqrt{\widehat{\text{Var}}(\hat{Y}_h)}$ . A confidence interval is produced through the interval option. Prediction intervals are produced by changing "confidence" to "prediction". The <i>level</i> option specifies the confidence level.
<code>anova(mod.fit)</code>	Finds the ANOVA table using information in an object called <i>mod.fit</i>
<code>boxcox(object = mod.fit, lambda = seq(from = -2, to = 2, by = 0.01))</code>	Finds $\hat{\lambda}$ for the Box-Cox transformation by looking in a region between -2 and 2
<code>ifelse(test = condition to test, yes = yes value, no = no value)</code>	Returns a vector of values corresponding to the result of the test
<code>leveneTest(y = mod.fit\$residuals, group = group)</code>	Perform Levene's test for residuals in <i>mod.fit</i> that have been split into <i>group</i> ; this function is in the <i>car</i> package
<code>bptest(formula = y ~ x, data = data.frame, studentize = FALSE)</code>	Perform the Breusch-Pagan Test where y is the response variable, x is the predictor variable, and they are stored in a particular <i>data.frame</i> ; this function is in the <i>lmtest</i> package
<code>examine.mod.simple(mod.fit.obj = mod.fit, const.var.test = TRUE, boxcox.find = TRUE)</code>	Runs my diagnostics function that is contained in <i>examine.mod.simple.R</i> and uses the model specified in <i>mod.fit</i> , constant variance tests are run and the Box-Cox transformation is found
<code>confint(object = mod.fit, level = 1 - alpha/g)</code>	Finds Bonferroni joint confidence intervals for g $\beta$ 's in the model specified in <i>mod.fit</i>

## Additional general R functions and examples

Syntax	Description
<b>Basic functions</b>	
<code>class(name)</code>	Finds the class of an object
<code>head(data.frame)</code>	Print the first six rows of the <i>data.frame</i>
<code>library(name)</code>	Make a library of functions called <i>name</i> ready to go in R
<code>mean(x = data)</code>	Find the mean of what is inside of <i>data</i>

Syntax	Description
<code>names(mod.fit)</code>	Allows one to see the elements in the list object called <i>mod.fit</i>
<code>set.seed(number)</code>	Set a seed number
<code>sum(x = data)</code>	Find the sum of what is inside of <i>data</i>
<b>Read in data</b>	
<code>read.table(file = "c:\\chris\\datafile.txt", header = TRUE, sep = "")</code>	Read in a text data file with variable names in the first row and spaces separating the variable names and their values. Comma delimited data files can be read in using the <code>sep=","</code> option (or <code>read.csv()</code> instead). If the first row does not contain the variable names, use <code>header = FALSE</code> and <code>col.names = c("var1", ..., "var.c")</code> to name the variables yourself.
<code>write.table(x = set1, file = "C:\\out_file.csv", quote = FALSE, row.names = FALSE, sep=",")</code>	Save data in a data frame to a file on the hard drive. The data was in the data frame <i>set1</i> and it will be written as a comma delimited file named <i>out_file.csv</i> .
<b>Plotting</b>	
<code>abline(h = 0)</code>	Plots a horizontal line at 0
<code>axis(side = 1, at = seq(from = 0, to = 4.5, by = 0.5), labels = true)</code>	Plot x-axis labels at 0 to 4.5 by 0.5.
<code>boxplot(x = x)</code>	Construct a box plot for data in a <i>vector x</i>
<code>curve(expr = x^2, xlim = c(-1, 1))</code>	Plots $f(x) = x^2$ for $-1 < x < 1$ ; the <code>add = TRUE</code> option can be used to add the curve to the current open plot
<code>hist(x = x)</code>	Construct a histogram for data in a <i>vector x</i>
<code>identify(x = x, y = y)</code>	This function allows you to interact with a plot to identify the observations on a y vs. x plot
<code>legend(locator(1), legend = vector of names, lty = vector of line types, col = vector of colors, bty = "n")</code>	Puts a legend on a plot where the insertion place is specified interactively with a left mouse click; no box is drawn around the legend
<code>plot(x = x, y = y)</code>	Plots <i>y</i> on the y-axis and <i>x</i> on the x-axis
<code>points(x = x, y = y, pch = 1, col = "red", cex = 2)</code>	Plots <i>y</i> on the y-axis and <i>x</i> on the x-axis with a plotting character of 1 (open circles), color of red, and symbol size that is twice as large as the default
<code>segments(x0 = x0, y0 = y0, x1 = x1, y1 = y1, lty=2)</code>	Draws a line segment between the points ( <i>x</i> <sub>0</sub> , <i>y</i> <sub>0</sub> ) and ( <i>x</i> <sub>1</sub> , <i>y</i> <sub>1</sub> ) using line type #2 (dashed line)
<code>stripchart(x = x, method = "jitter", vertical = TRUE)</code>	Construct a dot plot for data in a <i>vector x</i> with the points jittered; the points are plotted in a vertical manner
<code>win.graph(width = 6, height = 6, pointsize = 10)</code>	Opens a new graphics window that is 6"×6" with font size of 10
<b>Probability distributions</b>	

<b>Syntax</b>	<b>Description</b>
<code>pchisq(q = y, df = degrees of freedom)</code>	Finds $P(Y \leq y)$ for Y that has a chi-square distribution with a given degrees of freedom
<code>qchisq(p = 1-<math>\alpha</math>, df = degrees of freedom)</code>	Finds $1-\alpha$ quantile from a chi-square distribution for a given degrees of freedom
<code>pf(q = y, df1 = degrees of freedom 1, df2 = degrees of freedom 2)</code>	Finds $P(Y \leq y)$ for Y that has a F-distribution with a numerator (df1) and denominator (df2) degrees of freedom
<code>qf(p = 1-<math>\alpha</math>/2, df = degrees of freedom)</code>	Finds $1-\alpha/2$ quantile from a F-distribution with a numerator (df1) and denominator (df2) degrees of freedom
<code>pnorm(q = y)</code>	Finds $P(Y \leq y)$ for Y that has a standard normal distribution
<code>qnorm(p = 1-<math>\alpha</math>/2)</code>	Finds $1-\alpha/2$ quantile from a standard normal distribution
<code>pt(q = y, df = degrees of freedom)</code>	Finds $P(Y \leq y)$ for Y that has a t-distribution with a given degrees of freedom
<code>qt(p = 1-<math>\alpha</math>/2, df = degrees of freedom)</code>	Finds $1-\alpha/2$ quantile from a t-distribution for a given degrees of freedom
<b>Code examples</b>	
<code>group&lt;-ifelse(test = x &lt; median(x), yes = 1, no = 2) levene.test(y = mod.fit\$residuals, group = group)</code>	Perform Levene's test for residuals in <i>mod.fit</i> that have been split into two groups based on the median of x; note that the <i>levene.test()</i> function is in the car package; this code is similar to what is in HS_college_GPA_ch3.R

**Test #2 R part:**

Syntax	Description
<code>matrix(data = c( ), nrow = r , ncol = c, byrow = TRUE)</code>	Create a matrix of size $r \times c$ with data that is in the <code>c()</code> vector. The data is put into the matrix by rows.
<code>t(A)</code>	Find the transpose of a matrix A
<code>A%*%B</code>	Multiply the matrices A and B
<code>solve(A)</code>	Find the inverse of a matrix A
<code>J&lt;-matrix(data = 1, nrow = n, ncol = n)</code>	Create a $n \times n$ matrix of 1's
<code>as.numeric(object)</code>	Treat an object's contents as numerical only
<code>diag(n)</code>	Create a $n \times n$ identity matrix
<code>vcov(mod.fit)</code>	Calculate the estimated covariance matrix for <b>b</b> using results from the <code>lm()</code> function stored in <code>mod.fit</code>
<code>cbind(1, c( ))</code>	Combine by columns the number 1 and a vector of data given by the <code>c()</code> function. The 1 value is "recycled" to match the same length as what is given by the <code>c()</code> function
<code>scatter3d(x = x, y = y, z = z, fit="linear", bg="white", grid=TRUE)</code>	Create a 3D scatter plot of the data in x, y, and z, and plot the multiple linear regression model using y as the response variable
<code>nba[nba\$last.name == "Jordan"   nba\$last.name == "Stockton",]</code>	Extract rows of a data set called <code>nba</code> that match <code>last.name</code> values of Jordan <b>or</b> Stockton
<code>anova(mod.fit.red, mod.fit.comp, test = "F")</code>	Calculate the partial F-test statistic and p-value where <code>mod.fit.red</code> contains the reduced model results and <code>mod.fit.comp</code> contains the complete model results using separate implementations of the <code>lm()</code> function
<code>lm(formula = y ~ x + I(x^2), data = set1)</code>	This is an example of how to include a quadratic term in a regression model
<code>lm(formula = y ~ I(x-mean(set1\$x)) + I((x-mean(set1\$x))^2), data = set1)</code>	This is an example of how to include a mean adjusted quadratic term in a regression model
<code>library(Rcmdr)</code> <code>library(rgl)</code> <code>rgl.clear("all")</code> <code>rgl.light()</code> <code>rgl.bbox()</code> <code>scatter3d(x = x, y = y, z = z, fit="quadratic", grid=TRUE, bg.col="black")</code>	This is an example of how to construct a 3D scatter plot with a 2 <sup>nd</sup> order sample regression model
<code>lm(formula = y ~ x1 + x2 + x1:x2, data = set1)</code>	This is an example of how to include an interaction term in a regression model
<code>aggregate(formula = y ~ x, data = set1, FUN = mean)</code>	Finds the mean of y for each unique value of x. Other functions could be used to summarize y too by changing the <code>FUN</code> argument value.
<code>contrasts(x)</code>	Allows one to see how R will code a qualitative variable x by using indicator variables
<code>levels(x)</code>	Gives the levels of a qualitative variable x



Syntax	Description
<code>set1\$x&lt;-relevel(set1\$x, ref = "b")</code>	Changes the reference level ("all 0 level") to "b" for the variable x in the data frame <i>set1</i> .
<code>library(package = multcomp) K&lt;-matrix(data = c(0,1,-1), nrow = 1, ncol = 3, byrow = TRUE) compare.means&lt;-glht(model = mod.fit, linfct = K) summary(compare.means, test = adjusted("none"))</code>	This shows how to use the multcomp package to obtain a test for $H_0: \beta_1 - \beta_2 = 0$ vs. $H_a: \beta_1 - \beta_2 \neq 0$ . The test argument of <i>summary()</i> can be changed to <i>adjusted("bonferroni")</i> to produce Bonferroni corrections to p-values when more than one test is performed.
<code>factor(x)</code>	This function creates a qualitative variable from the object x.
<code>fit.stat&lt;-function(model, data) {   mod.fit&lt;-lm(formula = model, data = data)   sum.fit&lt;-summary(mod.fit)   aic.mod&lt;-AIC(object = mod.fit, k = 2)   bic.mod&lt;-AIC(object = mod.fit, k =     log(length(mod.fit\$residuals)))   sse&lt;-anova(mod.fit)\$"Sum Sq"[     length(anova(mod.fit)\$"Sum Sq")]   p&lt;-length(mod.fit\$coefficients)   n&lt;-length(mod.fit\$residuals)   Cp&lt;-sse/MSE.all - (n-2*p)   press&lt;-sum(mod.fit\$residuals^2/     (1 - lm.influence(mod.fit)\$hat)^2)   data.frame(Rsq=sum.fit\$r.squared, AdjRsq =     sum.fit\$adj.r.squared, AIC.stat =     aic.mod, BIC.stat = bic.mod, PRESS =     press, Cp = Cp) } }</code>	Function used to calculate $R^2$ , $R_a^2$ , $AIC_p$ , $SBC_p$ , $PRESS_p$ , and $C_p$ . This function is in the <i>nba_ch9.R</i> program.
<code>text(x = x, y = y, labels = vector of labels)</code>	Put the text specified in labels on a plot at the desired (x, y) location
<code>regsubsets(x = model formula, data = set1,   method = "exhaustive", nbest=4)</code>	Finds the best set of predictor variables for a regression model out of those given in the model formula. The four best models are given for possible sets of 1, 2, ... predictor variables. The function is located in the <i>leaps()</i> package. The <i>summary()</i> and <i>plot()</i> functions can be used with the resulting object to summarize the results. The <i>subsets()</i> function in the <i>car</i> package can summarize the results as well.

Syntax	Description
<pre>step(object = mod.fit, direction = "forward",       scope=list(lower = model formula, upper = model formula), k = 2)</pre>	<p>Perform forward predictor variable selection between the models listed in the lower and upper list elements. One needs to fit a model first with <i>lm()</i> and use the <i>object</i> option to pass in a corresponding model fit object. The <i>k = 2</i> option corresponds to using the AIC (<i>k = log(sample size)</i>) can be used for SBC or BIC). The <i>direction = "backward"</i> or <i>"both"</i> options can be used to perform backward elimination or stepwise forward, respectively.</p>
<pre>test.var&lt;-function(Ho, Ha, data) {   Ho.mod&lt;-lm(formula = Ho, data = data)   Ha.mod&lt;-lm(formula = Ha, data = data)   anova.fit&lt;-anova(Ho.mod, Ha.mod)   round(anova.fit\$"Pr(&gt;F)"[2], 4) }</pre>	<p>This is a function that can be used to compare a complete and reduced model through the use of a partial F-test. The function is in the nba_ch9.R program.</p>

## Final exam R part:

<b>Syntax</b>	<b>Description</b>
<code>avPlots(model = mod.fit)</code>	Constructs added variable plots corresponding to the information in <i>mod.fit</i> . The function is in the <i>car</i> package.
<code>rstandard(model = mod.fit)</code>	Find the studentized residuals using information in <i>mod.fit</i>
<code>rstudent(model = mod.fit)</code>	Find the studentized deleted residuals using information in <i>mod.fit</i>
<code>outlier.test(model = mod.fit)</code>	Perform a test for outliers using information in <i>mod.fit</i> and with a Bonferroni adjustment. This function is in the <i>car</i> package.
<code>hatvalues(model = mod.fit)</code>	Find the diagonal values of the hat matrix using information in <i>mod.fit</i>
<code>dffits(model = mod.fit)</code>	Find the DFFITS values using information in <i>mod.fit</i>
<code>cooks.distance(model = mod.fit)</code>	Find the Cook's distance values using information in <i>mod.fit</i>
<code>dfbetas(model = mod.fit)</code>	Find the DFBETAS values using information in <i>mod.fit</i>
<code>symbols(x = x, y = y, circles = z, inches = 0.25)</code>	Plot y on the y-axis and x on the x-axis. The plotting point is proportional to z with the largest size being 0.25 inches.
<code>vif(mod.fit)</code>	Find the VIF values using information in <i>mod.fit</i> . This function is in the <i>car</i> package.
<code>examine.mod.multiple.final(mod.fit.obj = mod.fit, first.order = number, const.var.test = TRUE, boxcox.find = TRUE)</code>	Use this function to examine various plots and summary measures to evaluate a model. See the <i>examine.mod.multiple.final.R</i> program file for more information.
<code>runif(n = n, min = 0, max = 1)</code>	Simulate n observations from a uniform distribution between 0 and 1. This was used to split a data set in the <i>NBA_ch10_validation.R</i> program.
<code>quantile(x = x, probs = c(0.2, 0.4, 0.6, 0.8), type = 1)</code>	Find the 0.2, 0.4, 0.6, 0.8 observed quantiles for a vector of data
<code>tapply(X = x, INDEX = groups, FUN = var)</code>	This is an example of how to apply the <code>var()</code> function to a vector of data that has been put into a number of groups. See the <i>weighted_least_squares.R</i> program for an example.
<code>lm(formula = Y ~ X, data = data, weight = 1/w)</code>	Find a regression model using weights in w.
<code>lqs(formula = Y ~ X, data = data, method = "method name")</code>	Find the parameter estimates of a regression model. The method option of "lms" is for least median squares. The method option of "lts" is for least trimmed squares. This function is in the <i>MASS</i> package.

Syntax	Description
<pre>persp(x = x, y = y, z = z, theta = 200, phi = 20,       ticktype = "detailed", expand = 1, shade       =0.5, col = "green3")</pre>	<p>Construct a 3D plot. The <i>theta</i> and <i>phi</i> options control the angle rotation of the plot. The other options can be used as needed to make the plot look nicer.</p>
<pre>loess(formula = Y ~ X, data = data, span = 0.5,       degree = 1)</pre>	<p>Find the first order lowess regression model for a value of <i>q</i> specified in the <i>span</i> option. A higher order model can be found by changing the <i>degree</i> option.</p>