CSC 423 Homework 3

Akhil Kumar Ramasagaram April 16, 2016

Deep space survey of quasars

- a) Hypothesize a first-order model for equivalent width, y, as a function of the first four variables. $y = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \beta_3 x_3 + \beta_4 x_4$
- b) Give the least squares prediction equation.

```
load("rdata/QUASAR.Rdata")
lm_model <- lm(RFEWIDTH ~ REDSHIFT + LINEFLUX + LUMINOSITY + AB1450, data = QUASAR)
summary(lm_model)

##
## Call:
## lm(formula = RFEWIDTH ~ REDSHIFT + LINEFLUX + LUMINOSITY + AB1450,</pre>
```

```
##
       data = QUASAR)
##
## Residuals:
##
      Min
                1Q Median
                                ЗQ
                                       Max
## -19.757 -9.039
                   -2.250
                             1.756
                                   48.628
##
## Coefficients:
##
                Estimate Std. Error t value Pr(>|t|)
## (Intercept) 21087.951 18553.161
                                     1.137
                                              0.2691
## REDSHIFT
                 108.451
                             88.740
                                     1.222
                                              0.2359
## LINEFLUX
                 557.910
                            315.990
                                     1.766
                                              0.0927 .
                -340.166
                            320.763 -1.060
## LUMINOSITY
                                              0.3016
## AB1450
                  85.681
                              6.273 13.658 1.34e-11 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 15.42 on 20 degrees of freedom
## Multiple R-squared: 0.9118, Adjusted R-squared: 0.8942
## F-statistic: 51.72 on 4 and 20 DF, p-value: 2.867e-10
```

```
lm_coeff <- summary(lm_model)$coefficients[,1]
lm_coeff</pre>
```

```
## (Intercept) REDSHIFT LINEFLUX LUMINOSITY AB1450
## 21087.95124 108.45084 557.90980 -340.16553 85.68102
```

The least square prediction equation is RFEWIDTH = 21087.95 + 108.45 * REDSHIFT + 557.90 * LINEFLUX - 340.165 * LUMINOSITY + 85.68 * AB1450

c) Interpret the β estimates in the model. β_0 is the intercept on y axis which is the value when all our predictors are zero. β_1 is our first predictor

which is REDSHIFT, the same can be said for β_2 , β_3 , β_4 for LINEFLUX, LUMINOSITY & AB1450. For every 1 unit increase in REDSHIFT the RFEWIDTHincreases by 108.45 value. The same can be interpreted for other β values. A positive/negative represents increase of decrease for unit increase.

- d) Test to determine whether redshift (x1) is a useful linear predictor of equivalent width (y), using $\alpha = .05$.
 - At $\alpha = 0.05$, the p-value of redsgift is 0.23, which is more than α , we can conclude that this is not a useful linear predictor.
- e) Locate R^2 and R_a^2 from the output. Interpret these values. Which statistic is the preferred measure of model fit? Explain?
 - The R^2 and R_a^2 from the text book output is 0.912 & 0.894 which can be located in the second table under model summary. The R-Square is more related to individual features performance and adjusted R-square is the performance of all the variable combined. Since we user multiplt variable in our model we should use adjusted R-square here.
- f) Locate the global F-value? we can locate the F-value in the third table(ANOVA) which is 51.720.

Removing oil from a water/oil mix.

```
load("rdata/WATEROIL.Rdata")
lm_model <- lm(VOLTAGE ~ VOLUME + SALINITY + SURFAC, data = WATEROIL)
predict(lm_model, newdata = data.frame('VOLUME' = 80, 'SALINITY' = 1, 'SURFAC' =2), interval = 'predicti</pre>
```

```
## fit lwr upr
## 1 -0.09795082 -1.233442 1.03754
```

the above prediction interval is an estimate of an interval in which our future observations will fall, with a 95% probability, given what has already been observed.

Arsenic in groundwater.

LATITUDE

-1279.76

```
load("rdata/ASWELLS.Rdata")
lm_model <- lm(ARSENIC ~ (LATITUDE + LONGITUDE) * DEPTHFT, data = ASWELLS)</pre>
summary(lm model)
##
## Call:
## lm(formula = ARSENIC ~ (LATITUDE + LONGITUDE) * DEPTHFT, data = ASWELLS)
##
## Residuals:
##
       Min
                1Q
                    Median
                                 3Q
                                        Max
## -175.75
           -65.04
                    -23.02
                              29.82
                                     480.01
## Coefficients:
##
                      Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                      10845.07
                                 67720.06
                                             0.160
                                                     0.8729
```

0.2252

1053.11 -1.215

```
## LONGITUDE
                     217.40
                                814.50 0.267
                                                0.7897
## DEPTHFT
                   -1549.22
                                985.58 -1.572
                                                0.1170
## LATITUDE:DEPTHFT
                     -11.00
                                11.86 -0.927
                                                0.3547
                                                0.0755 .
## LONGITUDE:DEPTHFT
                                 11.20
                                        1.783
                      19.98
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 103.1 on 321 degrees of freedom
    (1 observation deleted due to missingness)
## Multiple R-squared: 0.1372, Adjusted R-squared: 0.1238
## F-statistic: 10.21 on 5 and 321 DF, p-value: 4.306e-09
```

The least square equation for the above model can be written as ARSEINC = 10845.07 - 1279.76 * LATITUDE + 217.40 * LONGITUDE - 1549.22 * DEPTH - 11 * (LATITUDE : DEPTHFT) + 19.98 * (LONGITUDE : DEPTHFT)

With alpha = 0.05 the intereaction term does not have an effect on the arsenic level. The p-value for the LATITUDE:DEPTHFT term is 0.35 with t = -0.93 and for LONGITUDE:DEPTHFT term the p-value is 0.0755 with t = 1.78.

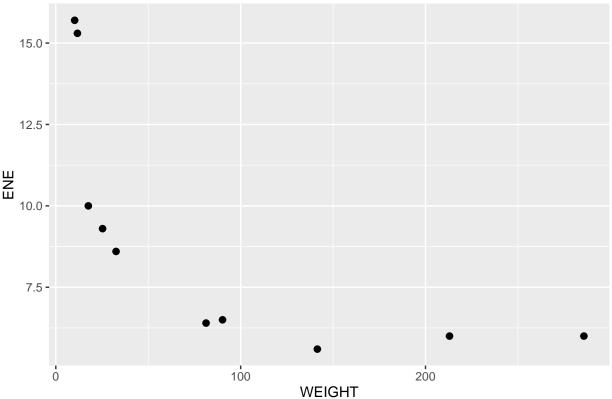
Carp diet study.

```
library(ggplot2)
```

Warning: package 'ggplot2' was built under R version 3.2.4

```
load("rdata/CARP.Rdata")
ggplot(CARP, aes(x = WEIGHT, y = ENE)) + geom_point(size = 2) +
    ggtitle("Scatterplot between Weight & Ene")
```

Scatterplot between Weight & Ene



Yes, there is a clear pattern in the above scatterplot. From the output, we can see that the p-value for the β_2 is 0.031. Our H_o : $\beta_2 = 0$ is rejected.

Homework assistance for accounting students.

(Intercept)

FULL

2.4333

-0.4833

0.4941

0.7813 -0.619

```
library(stringr)
library(nnet)
load("rdata/ACCHW.Rdata")
ACCHW$ASSIST <- as.character(ACCHW$ASSIST)
ACCHW$ASSIST <- str_trim(ACCHW$ASSIST)
ACCHW <- cbind(ACCHW, data.frame(class.ind(ACCHW$ASSIST)))
lm_model <- lm(IMPROVE ~ FULL + CHECK, ACCHW)</pre>
summary(lm_model)
##
## Call:
## lm(formula = IMPROVE ~ FULL + CHECK, data = ACCHW)
##
## Residuals:
##
      Min
              1Q Median
                             3Q
                                   Max
## -5.433 -2.433 0.050 1.567 6.567
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
```

4.925 5.2e-06 ***

0.538

```
## CHECK    0.2867    0.7329    0.391    0.697
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.706 on 72 degrees of freedom
## Multiple R-squared: 0.01244, Adjusted R-squared: -0.01499
## F-statistic: 0.4535 on 2 and 72 DF, p-value: 0.6372
```

We can use categorical features in a regression by converting into dummy variable. In the above case the class ind function from the neural net package creates a dummy variable. The equation can be writtern as Improve = 2.433 - 0.48 * Full + 0.28 * Check. Note that the model fails the overall F-test (p-value = 0.6372).