CS-E-106: Data Modeling

Assignment 3

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Solution 1:

```
reg_loop <- function(df, x_cols, y_str) {</pre>
  r2 list = c()
  lm_fits = list({})
  for(i in 1:length(x_cols)){
   x_str = x_cols[i]
   formula = as.formula(paste(y_str,"~", x_str))
   lm = lm(formula, data=df)
   r2_list[[i]] = summary(lm)$r.squared
  r2_df = data.frame(cbind(x_cols, r2_list))
  ordered_df = r2_df[order(r2_list, decreasing = TRUE),]
  print(paste("Variable with maximum R-squared:",x_cols[which(r2_list == max(r2_list))]))
  print(paste("R-squared value:",r2_list[which(r2_list == max(r2_list))]))
  return(ordered_df)
}
cdi = read.csv("cdi.csv")
colnames(cdi)
   [1] "Identification.number"
## [2] "County"
## [3] "State"
## [4] "Land.area"
## [5] "Total.population"
##
  [6] "Percent.of.population.aged.18.34"
## [7] "Percent.of.population.65.or.older"
## [8] "Number.of.active.physicians"
## [9] "Number.of.hospital.beds"
## [10] "Total.serious.crimes"
## [11] "Percent.high.school.graduates"
## [12] "Percent.bachelor.s.degrees"
## [13] "Percent.below.poverty.level"
## [14] "Percent.unemployment"
## [15] "Per.capita.income"
## [16] "Total.personal.income"
## [17] "Geographic.region"
exc = c("Identification.number", "Number.of.active.physicians")
x_cols = setdiff(colnames(cdi), exc)
y_str = "Number.of.active.physicians"
r2_df = reg_loop(df=cdi, x_cols = x_cols, y_str=y_str)
## [1] "Variable with maximum R-squared: County"
## [1] "R-squared value: 0.921236638865801"
```

Thus, county accounts for maximum variability in the number of active physicians. The remainder variables and their respective R^2 is given below in descending order of importance.

```
r2_df
                                                      r2_list
##
                                  x_{cols}
## 1
                                            0.921236638865801
                                  County
## 7
                Number.of.hospital.beds
                                            0.903382565497334
## 14
                  Total.personal.income
                                            0.898913655463206
## 4
                       Total.population
                                            0.884067412249688
## 8
                   Total.serious.crimes
                                            0.673153752663095
## 13
                      Per.capita.income
                                           0.0999411008221881
## 2
                                   State
                                            0.063458287522148
## 10
             Percent.bachelor.s.degrees
                                           0.0560578858594275
## 5
       Percent.of.population.aged.18.34
                                           0.0143279081163583
## 3
                               Land.area 0.00609565213240784
            Percent.below.poverty.level
                                         0.00411345912581384
## 11
## 12
                   Percent.unemployment 0.00255187801271758
## 15
                      Geographic.region 0.000607428795977754
          Percent.high.school.graduates 1.8046222736129e-05
     Percent.of.population.65.or.older 9.78832264836169e-06
## 6
Solution 2:
confint_regions <- function(df) {</pre>
  regions = levels(factor(df$Geographic.region))
  lm_fits = list({})
  for(i in regions){
    formula = as.formula(paste("Number.of.active.physicians","~", "Percent.bachelor.s.degrees"))
    lm = lm(formula, data=df[df$Geographic.region==i,])
    lm_fits[[i]] = lm
    print(paste("Region:",i))
    print(confint(lm, level=0.9)[2,])
    print(summary(lm)$coefficients)
    cat("\n")
  }
}
confint_regions(df=cdi)
## [1] "Region: 1"
##
        5 %
                95 %
## 37.52342 99.62067
##
                                 Estimate Std. Error
                                                         t value
                                                                     Pr(>|t|)
## (Intercept)
                               -399.33942 429.03669 -0.9307815 0.3541860656
## Percent.bachelor.s.degrees
                                 68.57205
                                            18.70308 3.6663503 0.0003944817
##
##
  [1] "Region: 2"
        5 %
##
                95 %
## 15.34141 88.23268
##
                                 Estimate Std. Error
                                                         t value
                                                                   Pr(>|t|)
## (Intercept)
                               -163.67212 463.91810 -0.3528039 0.72493602
## Percent.bachelor.s.degrees
                                51.78704
                                            21.96372 2.3578445 0.02021586
##
## [1] "Region: 3"
```

```
##
        5 %
                95 %
## 21.38184 58.19190
##
                               Estimate Std. Error
                                                        t value
                              -16.12031 251.04144 -0.06421374 0.9488855299
## (Intercept)
## Percent.bachelor.s.degrees 39.78687
                                          11.12036 3.57784135 0.0004669194
##
## [1] "Region: 4"
                    95 %
##
          5 %
##
   -2.927066 149.839187
##
                                Estimate Std. Error
                                                        t value Pr(>|t|)
## (Intercept)
                              -265.54556 1063.69869 -0.2496436 0.8035455
## Percent.bachelor.s.degrees
                                73.45606
                                            45.86403 1.6016049 0.1134469
Thus, the slopes for the regression lines for differen regions vary from one another.
Solution 3:
(a)
gpa = read.csv("GPA.csv")
lm_gpa = lm(GPA~ACT, data=gpa)
summary(lm_gpa)
##
## Call:
## lm(formula = GPA ~ ACT, data = gpa)
##
## Residuals:
##
        Min
                  1Q
                      Median
                                     30
                                             Max
## -2.74004 -0.33827 0.04062 0.44064 1.22737
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 2.11405
                           0.32089
                                      6.588 1.3e-09 ***
## ACT
                0.03883
                           0.01277
                                      3.040 0.00292 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.6231 on 118 degrees of freedom
## Multiple R-squared: 0.07262,
                                    Adjusted R-squared: 0.06476
## F-statistic: 9.24 on 1 and 118 DF, p-value: 0.002917
anova(lm_gpa)
## Analysis of Variance Table
## Response: GPA
##
              Df Sum Sq Mean Sq F value
                                          Pr(>F)
               1 3.588 3.5878 9.2402 0.002917 **
## ACT
## Residuals 118 45.818 0.3883
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
(b)
MSR = \sum_{i=1}^{n} (\hat{Y}_i - \bar{Y})^2
```

MSR measures the effect of the regression line in explaining the total variation in Y_i .

$$MSE = \frac{\sum_{i=1}^{n} (Y_i - \hat{Y}_i)^2}{n-2}$$

MSE measures the mean variation of Y_i around the regression line. Its the average of all the squared distances by which the regression line missed the actual Y_i .

$$E[MSE] = \sigma^2$$

$$E[MSR] = \sigma^2 + \beta_1 \sum_{i=1}^{n} (X_i - \bar{X})^2$$

Thus, MSE and MSR will estimate same quantity when $\beta_1 = 0$ i.e. $Y_i = \bar{Y}$

(c)

Null Hypothesis: $H_0: \beta_1 = 0$; Alternate Hypothesis: $H_1: \beta_1 \neq 0$

Decision Rule:

$$F^* = \frac{MSR}{MSE}$$

- If $F* \leq F(1-\alpha; 1, n-2)$, conclude H_0 ;
- If $F* \geq F(1-\alpha; 1, n-2)$, conclude H_1

```
MSR = 3.5878

MSE = 0.3883

F = MSR/MSE

print(F)
```

[1] 9.239763

```
help(pf)
pf(q=0.01, 1, 118)
```

[1] 0.07948598

Result:

Thus, since $F^* > F(1 - \alpha; 1, n - 2)$, we conclude that $H_1 : \beta_1 \neq 0$ holds.

(d)

The absolute magnitude of the reduction in the variation of Y when X is introduced into the regression model is SST - SSE = SSR = 3.588 (from the ANOVA table above).

The relative measure is given by $\frac{SSR}{SST} = \frac{3.588}{3.588+45.818} = 0.0726$. This measure is also known as the R^2 or the coefficient of determination.

```
R_sq = 3.588/(3.588+45.818)
R_sq
```

[1] 0.07262276

(e)

```
r = sqrt(R_sq)
r
```

[1] 0.2694861

Looking at the summary of the regression model for GPA dataset, β_1 is positive. Hence, r = +0.27.

(f)

Operationally, \mathbb{R}^2 has more clear interpretation.

- R^2 is the proportion of total variation in Y explained by X. Thus, it is a relative measure of improvement that was made by the introduction of X in the regression model. This can be used in a more direct way compared to r which measures linear association between X and Y.
- R^2 is on a scale of 0 to 1 (1 indicating the highest correlation), whereas, r ranges from -1 to 1 (the extremes indicating highest correlation). Meaning both r=-1 and r=+1 can mean the same level of association between X and Y. Also, the objective of the coefficients of correlation/determination is to measure the overall effectiveness of the model rather than looking at which direction the regression line is going. This is better accomplished by R^2 .

Solution 4:

Response: Y

(a) crime = read.csv("Crime Rate.csv") cor.test(crime\$Y,crime\$X,method="pearson") ## ## Pearson's product-moment correlation ## ## data: crime\$Y and crime\$X ## t = -4.1029, df = 82, p-value = 9.571e-05 ## alternative hypothesis: true correlation is not equal to 0 ## 95 percent confidence interval: -0.5761223 -0.2175580 ## sample estimates: ## cor ## -0.4127033 (b) lm crime = lm(Y~X, data=crime)summary(lm_crime) ## ## Call: ## lm(formula = Y ~ X, data = crime) ## ## Residuals: ## Min 10 Median 3Q Max ## -5278.3 -1757.5 -210.5 1575.3 6803.3 ## ## Coefficients: Estimate Std. Error t value Pr(>|t|) ## (Intercept) 20517.60 6.260 1.67e-08 *** 3277.64 ## X -170.5841.57 -4.103 9.57e-05 *** ## ---## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1 ## ## Residual standard error: 2356 on 82 degrees of freedom ## Multiple R-squared: 0.1703, Adjusted R-squared: 0.1602 ## F-statistic: 16.83 on 1 and 82 DF, p-value: 9.571e-05 anova(lm crime) ## Analysis of Variance Table ##

```
Sum Sq Mean Sq F value
                                                     Pr(>F)
                1 93462942 93462942 16.834 9.571e-05 ***
## Residuals 82 455273165 5552112
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
Null Hypothesis: H_0: \beta_1 = 0; Alternate Hypothesis: H_1: \beta_1 \neq 0
Decision Rule:
F^* = \frac{MSR}{MSE}
   • If F* \leq F(1-\alpha; 1, n-2), conclude H_0;
   • If F* \ge F(1-\alpha; 1, n-2), conclude H_1
MSR = 93462942
MSE = 5552112
F = MSR/MSE
print(F)
## [1] 16.83376
pf(q=0.01, 1, 82)
## [1] 0.07941159
Result:
Thus, since F^* > F(1 - \alpha; 1, n - 2), we conclude that H_1 : \beta_1 \neq 0 holds.
(c)
cor.test(crime$Y,crime$X, method="spearman")
## Warning in cor.test.default(crime$Y, crime$X, method = "spearman"): Cannot
## compute exact p-value with ties
##
##
    Spearman's rank correlation rho
##
## data: crime$Y and crime$X
## S = 140839, p-value = 5.359e-05
## alternative hypothesis: true rho is not equal to 0
## sample estimates:
##
           rho
## -0.4259324
(d)
Null Hypothesis: There is no association between X and Y; Alternate Hypothesis: There is an association
between X and Y
Decision Rule:
t^* = \frac{r_s \sqrt{n-2}}{1-r_s^2}
   • If |t^*| \le t(1 - \alpha/2; n - 2), conclude H_0;
   • If |t^*| \ge t(1 - \alpha/2; n - 2), conclude H_1
r s = -0.4259324
n = nrow(crime)
```

```
t = (r_s*sqrt(n-2)/(1-r_s^2))

t

## [1] -4.711787

pt(0.005, 82)

## [1] 0.5019886
```

Result:

Thus, since $|t^*| \ge t(1 - \alpha/2; n - 2)$, we conclude that H_1 that there is an association between X and Y.