# [STAT 4400] HW-1

Michael Ghattas

1/17/2022

#### Question-1

$$n = 3$$

$$\overline{y} = \frac{1}{n} \sum_{i=1}^{n} y_i = \frac{1+2+3}{3} = \frac{6}{3} = 2$$

$$\overline{x} = \frac{1}{n} \sum_{i=1}^{n} x_i = \frac{2+0+4}{3} = \frac{6}{3} = 2$$

(a)

$$\hat{\beta}_{0} = \bar{y} - \hat{\beta}_{1}\bar{x}$$

$$\hat{\beta}_{1} = \frac{\sum_{i=1}^{n} (y_{i} - \bar{y})(x_{i} - \bar{x})}{\sum_{i=1}^{n} (x_{i} - \bar{x})^{2}}$$

$$\hat{\sigma}^2 = \frac{1}{n-2} \sum_{i=1}^n (y_i - \hat{\beta}_0 - \hat{\beta}_1 x_i)^2$$

(b)

$$\hat{\beta}_1 = \frac{\sum_{i=1}^3 (y_i - 2)(x_i - 2)}{\sum_{i=1}^3 (x_i - 2)^2} = \frac{(-1 + 0 + 1)(0 - 2 + 2)}{(0)^2 + (-2)^2 + (2)^2} = \frac{(0)(0)}{4 + 4} = \frac{0}{8} = 0$$

$$\hat{\beta}_0 = 2 - (0)\bar{x} = 2 - 0 = 2$$

$$\hat{\sigma}^2 = \frac{1}{3-2} \sum_{i=1}^{3} (y_i - 2 - (0)x_i)^2 = \frac{1}{1} \sum_{i=1}^{3} (y_i - 2)^2 = 1[(-1)^2 + (0)^2 + (1)^2] = 1 + 0 + 1 = 2$$

$$\hat{\epsilon}_i = y_i - \hat{y}_i = y_i - \hat{\beta}_0 - \hat{\beta}_1 x_i = y_i - 2 - 0 = y_i - 2$$

$$SSE = \sum_{i=1}^{3} \hat{\epsilon}_i^2 = (1-2)^2 + (2-2)^2 + (3-2)^2 = (-1)^2 + (0)^2 + (1)^2 = 1 + 0 + 1 = 2$$

$$\hat{y}_{i} = \hat{\beta}_{0} - \hat{\beta}_{1}x_{i} + \hat{\epsilon}_{i} = 2 - 0 + \hat{\epsilon}_{i} = 2 + \hat{\epsilon}_{i}$$

$$SSR = \sum_{i=0}^{3} (\hat{y}_{i} - 2)^{2} = ((2 + (1 - 2)) - 2)^{2} + ((2 + (2 - 2)) - 2)^{2} + ((2 + (3 - 2)) - 2)^{2} = ((2 - 1) - 2)^{2} + ((2 - 0) - 2)^{2} + ((2 + 1) - 2)^{2} = (1 - 2)^{2} + (2 - 2)^{2} + (3 - 2)^{2} = (-1)^{2} + (0)^{2} + (1)^{2} = 1 + 0 + 1 = 2$$

$$SST = \sum_{i=1}^{3} (y_i - 2)^2 = (1 - 2)^2 + (2 - 2)^2 + (3 - 2)^2 = (-1)^2 + (0)^2 + (1)^2 = 1 + 0 + 1 = 2$$

$$R^2 = 1 - \frac{SSE}{SST} = \frac{2}{2} = 1$$

(d)

$$H_0: \beta_1 = 0$$

$$H_1:\beta_1\neq 0$$

$$\beta_1 * = 0$$

$$\hat{\beta}_1 \sim N(0, \tau)$$

$$\hat{\tau} = \sqrt{\frac{\sigma^2}{\sum_{i=1}^3 (x_i - \bar{x})^2}} = \sqrt{\frac{2}{\sum_{i=1}^3 (x_i - 2)^2}} = \sqrt{\frac{2}{(1 - 2)^2 + (2 - 2)^2 + (3 - 2)^2}} = \sqrt{\frac{2}{(-1)^2 + (0)^2 + (1)^2}} = \sqrt{\frac{2}{1 + 0 + 1}} = \sqrt{\frac{2}{2}} = \sqrt{1 = 1}$$

$$\hat{t} = \frac{\hat{\beta}_1}{\hat{\tau}}$$

$$p[|t| \ge t^*] = \alpha$$

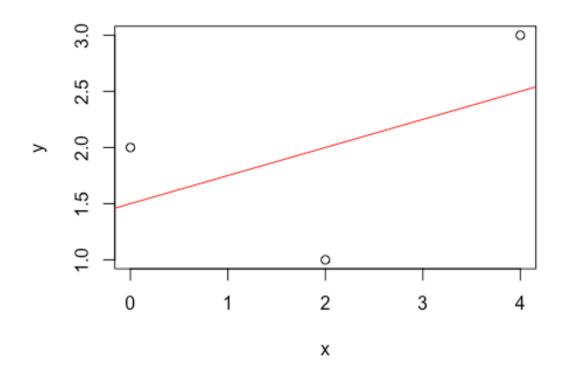
$$y = c(1, 2, 3)$$

$$x = c(2, 0, 4)$$

$$alpha = 0.5$$

$$n = length(x)$$

```
sigma2.hat = 2
std.t = dt(x, (n - 2))
t.star = qt((1 - alpha / 2), df = (n - 2))
y.hat = beta.0.hat + (beta.1.hat * x)
eps.hat = y - y.hat
SSE = sum(eps.hat ** 2)
sigma2.hat = SSE / (n - 2)
tau.hat = sqrt(sigma2.hat / sum((x - mean(x)) ** 2))
t.hat = beta.1.hat / tau.hat
t.star
## [1] 1
t.hat
## [1] 0
print("(|t| < t^*)) Thus we fail to reject the the null-hypothesis")
## [1] "(|t| < t^*) Thus we fail to reject the the null-hypothesis"
(e)
lmod = lm(y \sim x)
plot(x, y)
abline(lmod, col = "red")
```



## **Question-2**

$$0 \le x \le 50 \overline{x} = 35 \sigma_x = 10$$

$$y = a + bx \overline{y} = a + b\overline{x} = a + 35b$$

$$\sigma_y = 15 \overline{y} = 100$$

$$\sigma_y = |b| \sigma_x = 10 |b| 15 = 10 |b| \rightarrow |b| = \frac{15}{10} = 1.5 \rightarrow b = \pm 1.5$$

$$a + 35b = 100 \rightarrow a + 35(\pm 1.5) = 100$$

$$a + 35(-1.5) = 100 \rightarrow a - 52.5 = 100 \rightarrow a = 100 + 52.5 = 152.5$$

$$a + 35(1.5) = 100 \rightarrow a + 52.5 = 100 \rightarrow a = 100 - 52.5 = 47.5$$

```
(a) y = 152.5 - 1.5x (b) y = 47.5 + 1.5x (c)
```

I would recommend the solution from part (a), as it has an increasing function that exhibits a positive correlation.

### **Question-3**

(a)

```
set.seed(123)
var1 = rnorm(1000, mean = 0, sd = 1)
var2 = rnorm(1000, mean = 0, sd = 1)
lmod1 = lm(var1 \sim var2)
summary(lmod1)
##
## Call:
## lm(formula = var1 ~ var2)
##
## Residuals:
##
      Min
              10 Median
                             3Q
                                   Max
## -2.7168 -0.6290 -0.0060 0.6451 3.2383
##
## Coefficients:
             Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 0.01252
                        0.03129
                                 0.400 0.68909
## var2
              ## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

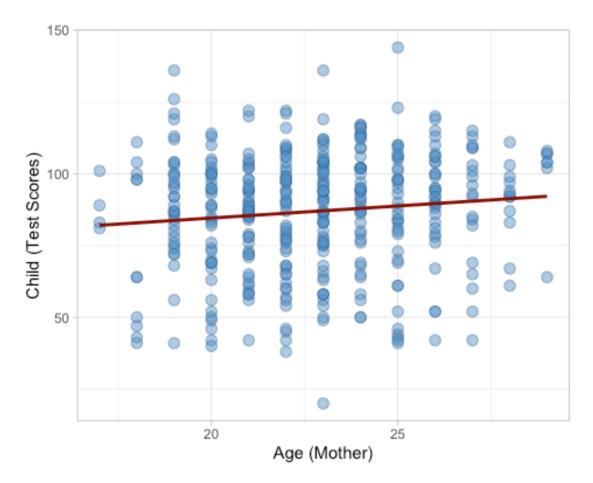
```
##
## Residual standard error: 0.9885 on 998 degrees of freedom
## Multiple R-squared: 0.007479, Adjusted R-squared: 0.006484
## F-statistic: 7.52 on 1 and 998 DF, p-value: 0.006211
print("Yes, based on the p-value generated we can conclude that the slope
coefficient var2 is statistically significant.")
## [1] "Yes, based on the p-value generated we can conclude that the slope
coefficient var2 is statistically significant."
(b)
set.seed(321)
z.scores <- rep (NA, 100)
for (k in 1:100)
 var1 <- rnorm (1000 ,0 ,1)</pre>
 var2 <- rnorm (1000 ,0 ,1)</pre>
 fit <- lm (var2 ~ var1)</pre>
  z.scores[k] <- coef(fit )[2] / summary(fit)$coef[2,"Std. Error"]</pre>
}
alpha = .05
cutoffn = gnorm((1 - alpha) / 2, lower.tail=TRUE)
sum(abs(z.scores) > cutoffn)
## [1] 100
for (k in 1:100)
{
  result = sum(abs(z.scores) < 1.96)
}
result
## [1] 95
```

```
print("95 estimated slope coefficients are statistically significant at the \alpha = .05 level of significance.") ## [1] "95 estimated slope coefficients are statistically significant at the \alpha = .05 level of significance."
```

### **Question-4**

```
(a)
library(haven)
data <- read dta("/Users/Home/Documents/Michael Ghattas/School/CU Boulder/</pre>
2022/Spring 2022/STAT - 4400/HW/1/child.iq.dta")
head(data)
## # A tibble: 6 × 3
##
      ppvt educ cat momage
     <dh1>
              <dbl> <dbl>
##
                  2
## 1
       120
                         21
## 2
        89
                  1
                         17
## 3
        78
                  2
                         19
## 4
        42
                  1
                         20
## 5
                  4
                         26
       115
## 6
        97
                  1
                         20
lmod = lm(ppvt ~ momage, data = data)
summary(lmod)
##
## Call:
## lm(formula = ppvt ~ momage, data = data)
##
## Residuals:
       Min
##
                10 Median
                                 30
                                        Max
## -67.109 -11.798
                     2.971 14.860 55.210
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
                             8.6880
                                      7.802 5.42e-14 ***
## (Intercept) 67.7827
```

```
## momage
                0.8403
                           0.3786
                                    2.219
                                             0.027 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 20.34 on 398 degrees of freedom
## Multiple R-squared: 0.01223, Adjusted R-squared: 0.009743
## F-statistic: 4.926 on 1 and 398 DF, p-value: 0.02702
library(ggplot2)
ggplot(data, aes(momage, ppvt)) +
geom point(shape = 21, color="steelblue4", fill="steelblue3", size = 3,
alpha=0.5,show.legend = FALSE) +
theme light() + xlab("Age (Mother)") + ylab("Child (Test Scores)") +
geom smooth(method = lm, color="darkred", se=FALSE)
## `geom smooth()` using formula 'y ~ x'
```

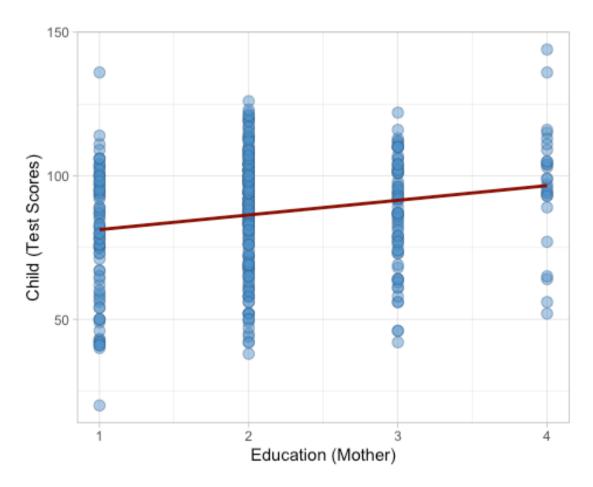


print("Based on our summary and plots, it seems that the mother's age is a significant predictor, though not enough on its own to find a direct correlation. That being said, the plot clearly shows that the majority of the observations with a higher test score belong to the mothers in their late 20's. Thus mothers should give birth in their late 20s.")

## [1] "Based on our summary and plots, it seems that the mother's age is a significant predictor, though not enough on its own to find a direct correlation. That being said, the plot clearly shows that the majority of the observations with a higher test score belong to the mothers in their late 20's. Thus mothers should give birth in their late 20s."

```
(b)
lmod = lm(ppvt ~ momage + educ_cat, data = data)
summary(lmod)
```

```
##
## Call:
## lm(formula = ppvt ~ momage + educ cat, data = data)
##
## Residuals:
##
       Min
                10 Median
                               30
                                      Max
## -61.763 -13.130 2.495 14.620 55.610
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 69.1554
                           8.5706
                                    8.069 8.51e-15 ***
## momage
                           0.3981
                                    0.862 0.389003
                0.3433
## educ cat
                           1.3165 3.579 0.000388 ***
              4.7114
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 20.05 on 397 degrees of freedom
## Multiple R-squared: 0.04309,
                                  Adjusted R-squared: 0.03827
## F-statistic: 8.939 on 2 and 397 DF, p-value: 0.0001594
library(ggplot2)
ggplot(data, aes(educ cat, ppvt)) +
geom point(shape = 21, color="steelblue4", fill="steelblue3", size = 3,
alpha=0.5, show.legend = FALSE) +
theme_light() + xlab("Education (Mother)") + ylab("Child (Test Scores)") +
geom smooth(method = lm, color="darkred", se=FALSE)
## `geom_smooth()` using formula 'y ~ x'
```



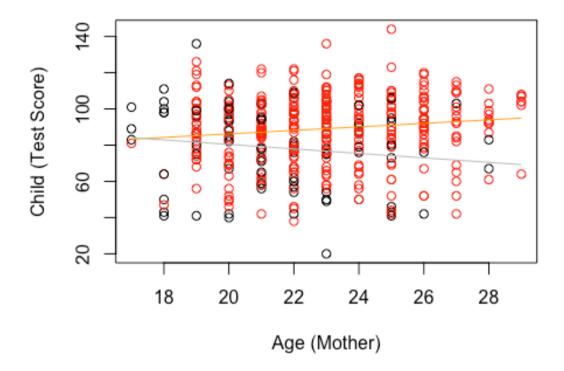
print("Based on our summary and plots, it seems that the mother's education
is a strong and significant predictor. That being said, the plot clearly
shows that the majority of the observations with a higher test score belong
to the mothers having completed a high-school education.")

## [1] "Based on our summary and plots, it seems that the mother's education is a strong and significant predictor. That being said, the plot clearly shows that the majority of the observations with a higher test score belong to the mothers having completed a high-school education."

```
(c)
data$mom.hs <- ifelse(data$educ_cat >= 2, 1, 0)

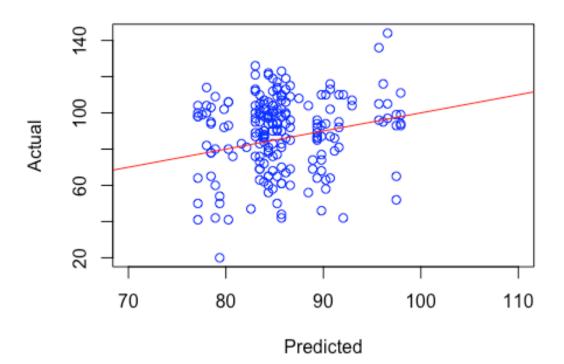
lmod <- lm(ppvt ~ (mom.hs * momage), data = data)
summary(lmod)</pre>
```

```
##
## Call:
## lm(formula = ppvt ~ (mom.hs * momage), data = data)
##
## Residuals:
##
      Min
               10 Median
                               30
                                      Max
## -56.696 -12.407 2.022 14.804 54.343
##
## Coefficients:
                Estimate Std. Error t value Pr(>|t|)
                105.2202
                            17.6454 5.963 5.49e-09 ***
## (Intercept)
## mom.hs
               -38,4088
                            20.2815 -1.894
                                              0.0590 .
## momage
                 -1.2402
                            0.8113 -1.529
                                              0.1271
## mom.hs:momage 2.2097 0.9181 2.407 0.0165 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 19.85 on 396 degrees of freedom
## Multiple R-squared: 0.06417. Adjusted R-squared: 0.05708
## F-statistic: 9.051 on 3 and 396 DF, p-value: 8.276e-06
spread <- ifelse(data$mom.hs == 1, "red", "black")</pre>
plot(data$momage, data$ppvt, xlab = "Age (Mother)", ylab = "Child (Test
Score)", col = spread, pch = 1)
curve(cbind(1, 1, x, 1 * x) %*% coef(lmod), add = TRUE, col = "orange")
# Mother finished hs
curve(cbind(1, 0, x, 0 * x) % coef(lmod), add = TRUE, col = "grey")
# Mother did not finish hs
```



```
(d)
lmod <- lm(ppvt ~ momage + educ_cat, data = data[1:200, ])</pre>
summary(lmod)
##
## Call:
## lm(formula = ppvt ~ momage + educ_cat, data = data[1:200, ])
##
## Residuals:
                    Median
                                 3Q
##
       Min
                 1Q
                                         Max
## -46.358 -12.967
                      2.866
                             14.435
                                      58.428
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
```

```
## (Intercept) 63.6295
                           11.8202
                                     5.383 2.07e-07 ***
## momage
                 0.4473
                            0.5516
                                     0.811 0.41836
## educ cat
                                     2.986 0.00318 **
                 5.4434
                            1.8228
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 19.58 on 197 degrees of freedom
## Multiple R-squared: 0.06199, Adjusted R-squared: 0.05246
## F-statistic: 6.509 on 2 and 197 DF, p-value: 0.001831
pmod <- predict(lmod, data[201:400, ])</pre>
plot(pmod, data$ppvt[201:400], xlim = c(70, 110), xlab = "Predicted",
    ylab = "Actual", col = "blue")
abline(a = 0, b = 1, col = "red")
```



### **Question-5**

(a)

```
library(faraway)
data(prostate)
head(prostate)
##
        lcavol lweight age
                                lbph svi
                                              lcp gleason pgg45
                                                                    lpsa
## 1 -0.5798185 2.7695 50 -1.386294
                                       0 -1.38629
                                                        6
                                                              0 -0.43078
## 2 -0.9942523 3.3196 58 -1.386294
                                       0 -1.38629
                                                        6
                                                              0 -0.16252
## 3 -0.5108256 2.6912 74 -1.386294
                                       0 -1.38629
                                                        7
                                                             20 -0.16252
## 4 -1.2039728 3.2828 58 -1.386294
                                       0 -1.38629
                                                        6
                                                              0 -0.16252
## 5 0.7514161 3.4324 62 -1.386294
                                       0 -1.38629
                                                        6
                                                              0 0.37156
## 6 -1.0498221 3.2288 50 -1.386294
                                       0 -1.38629
                                                              0 0.76547
                                                        6
lmod = lm(log(lpsa) ~ log(lcavol) + ., data = prostate)
## Warning in log(lpsa): NaNs produced
## Warning in log(lcavol): NaNs produced
summary(lmod)
##
## Call:
## lm(formula = log(lpsa) \sim log(lcavol) + ., data = prostate)
##
## Residuals:
       Min
                      Median
##
                 10
                                   30
                                           Max
## -1.42557 -0.12542 0.02419 0.19314 0.51973
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) -0.228166
                          0.665962 -0.343 0.73286
## log(lcavol) -0.073779
                          0.082178
                                   -0.898 0.37221
## lcavol
                                    3.189 0.00209 **
               0.298653
                          0.093642
## lweight
                                    1.543 0.12700
               0.132774
                          0.086027
## age
              -0.008582
                          0.005988 -1.433 0.15596
## lbph
               0.068535
                          0.030029
                                    2.282 0.02535 *
```

```
## svi
               0.247254
                         0.116662
                                    2.119 0.03741 *
## lcp
              -0.049417
                         0.044109 -1.120 0.26619
## gleason
               0.086846
                         0.075419
                                   1.152 0.25323
## pgg45
               0.001981
                         0.002088
                                    0.948 0.34605
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.3323 on 74 degrees of freedom
    (13 observations deleted due to missingness)
## Multiple R-squared: 0.4948, Adjusted R-squared: 0.4334
## F-statistic: 8.053 on 9 and 74 DF, p-value: 3.254e-08
confint(lmod, 'log(lcavol)', level = 0.95)
                   2.5 %
##
                            97.5 %
## log(lcavol) -0.2375227 0.08996445
(b)
lmod = lm(log(lpsa) \sim log(lcavol) + lcavol + lbph + svi, data = prostate)
## Warning in log(lpsa): NaNs produced
## Warning in log(lcavol): NaNs produced
summary(lmod)
##
## Call:
## lm(formula = log(lpsa) \sim log(lcavol) + lcavol + lbph + svi, data =
prostate)
##
## Residuals:
       Min
                     Median
                                  30
                                         Max
                 10
## -1.48196 -0.12939 0.05611 0.22173 0.48638
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
##
                                   2.825 0.005987 **
## (Intercept) 0.35496
                         0.12566
```

```
## lcavol
               0.29829
                          0.08713
                                    3.423 0.000983 ***
## lbph
                                    2.961 0.004043 **
               0.07834
                          0.02645
## svi
                                    2.356 0.020976 *
               0.24564
                          0.10428
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.3384 on 79 degrees of freedom
    (13 observations deleted due to missingness)
## Multiple R-squared: 0.4407, Adjusted R-squared:
## F-statistic: 15.56 on 4 and 79 DF, p-value: 1.98e-09
confint(lmod, 'log(lcavol)', level = 0.95)
##
                   2.5 %
                             97.5 %
## log(lcavol) -0.2350253 0.08509177
```

(c)

The model from part (a) has a slightly better fit, as the R^2 value is slightly higher, indicating a better fitted model.