AMS317 HW5 #2

Lucy Lin

2022-10-06

```
2a)
### ??faraway::seatpos
data = data(seatpos, package = "faraway")
fit1 <- lm(hipcenter ~ Ht, data=seatpos)</pre>
summary(fit1)
##
## Call:
## lm(formula = hipcenter ~ Ht, data = seatpos)
## Residuals:
      Min
               1Q Median
                              3Q
                                       Max
## -99.956 -27.850 5.656 20.883 72.066
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
                           90.6704
## (Intercept) 556.2553
                                    6.135 4.59e-07 ***
              -4.2650
                            0.5351 -7.970 1.83e-09 ***
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
## Residual standard error: 36.37 on 36 degrees of freedom
## Multiple R-squared: 0.6383, Adjusted R-squared: 0.6282
## F-statistic: 63.53 on 1 and 36 DF, p-value: 1.831e-09
Yes, it is significant at alpha = 0.01 (Signif. codes: 0'****)
2b)
fit2 <- lm(hipcenter ~ Weight, data=seatpos)</pre>
summary(fit2)
##
## Call:
## lm(formula = hipcenter ~ Weight, data = seatpos)
##
## Residuals:
##
       Min
                 1Q Median
                                    3Q
                                            Max
```

```
## -109.446 -23.179
                       -2.292
                                20.171 130.567
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
##
## (Intercept)
                1.2422
                           34.0563
                                     0.036
                                              0.971
                            0.2134 -5.002 1.49e-05 ***
## Weight
                -1.0674
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 46.45 on 36 degrees of freedom
## Multiple R-squared:
                        0.41, Adjusted R-squared: 0.3936
## F-statistic: 25.02 on 1 and 36 DF, p-value: 1.493e-05
Yes, it is significant at alpha = 0.01 (Signif. codes: 0^{***})
2c)
fit12 <- lm(hipcenter ~ Ht + Weight, data=seatpos)</pre>
summary(fit12)
##
## Call:
## lm(formula = hipcenter ~ Ht + Weight, data = seatpos)
##
## Residuals:
##
       Min
                1Q Median
                                3Q
                                       Max
## -99.061 -27.970
                    4.611 21.910 70.381
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 589.9006
                          127.4824
                                     4.627 4.92e-05 ***
## Ht
                -4.5697
                            0.9672
                                    -4.725 3.68e-05 ***
                 0.1148
                                     0.380
                                              0.706
## Weight
                            0.3020
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 36.81 on 35 degrees of freedom
## Multiple R-squared: 0.6398, Adjusted R-squared: 0.6192
## F-statistic: 31.08 on 2 and 35 DF, p-value: 1.738e-08
It is no longer significant.
2d)
cor(seatpos$Ht, seatpos$Weight)
```

[1] 0.8285257

Ht is considered to be the "impactful" variable instead since weight and height might be correlated. Height is "credited" as the more impactful variable as a result. Since they are highly correlated, weight does not appear to explain as much of the data.

```
2e) H_0: beta_Ht = 0 H_1: beta_Ht = /= 0
```

```
nested model: Y ~ x_Ht larger model: Y ~ x_Ht + x_Weight  
2f) H_0: beta_Ht = 0 and beta_Weight = 0 H_1: at least one of beta_Ht or beta_Weight =/= 0  
nested model: Y ~ 1 larger model: Y ~ 1 + x_Ht + x_Weight  
2g)
```

anova(fit2,fit12)

```
## Analysis of Variance Table
##
## Model 1: hipcenter ~ Weight
## Model 2: hipcenter ~ Ht + Weight
## Res.Df RSS Df Sum of Sq F Pr(>F)
## 1 36 77664
## 2 35 47420 1 30244 22.322 3.675e-05 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

- i) H_0 : beta_Ht = 0 H_1 : beta_Ht = /= 0
- ii) The F statistic for anova(fit2,fit12) is the square of that t test and have the same p value since they have the same nested and larger models.
- iii) Even though they are testing the same thing, the anova (fit1,fit2) has models: h_0: Y ~ 1 + beta_weight h_1: Y ~ 1 + beta_weight + beta_Ht

and fit1 had models: h_0: Y ~ 1 h_1: Y ~ 1 + beta_Ht

The models are not the same.