# Example 4.1

Kevin Cummiskey October 15, 2019

## Example 4.1 Recovering Polyphenols (pg 272)

### Lesson 16 - Simple Linear Regression

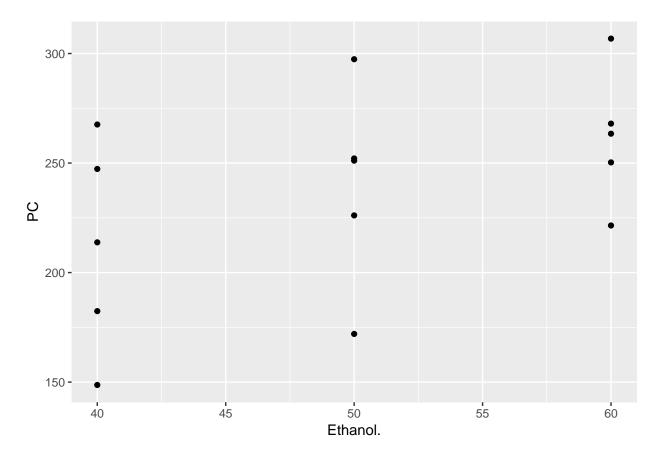
Objectives:

- 1. Describe association between two quantivative variables.
- 2. Interpret least-squares regression models.
- 3. Compare and contrast separate versus linear regression models.

What is the observational unit? explanatory variable? response variable?

What type of study is this?

```
grapes = read.table(file = "http://www.isi-stats.com/isi2/data/Polyphenols.txt", header = T)
grapes %>% ggplot(aes(x = Ethanol., y = PC)) + geom_point()
```



Write a statistical model for a separate mean for each ethanol level.

Fit the model.

```
grapes$Ethanol_cat = factor(grapes$Ethanol.)
contrasts(grapes$Ethanol_cat) = contr.sum
model_anova = lm(PC ~ Ethanol_cat, data = grapes)
summary(model_anova)
##
## Call:
## lm(formula = PC ~ Ethanol_cat, data = grapes)
##
## Residuals:
              1Q Median
##
      Min
                            3Q
                                 Max
## -67.74 -21.60
                  1.84 23.85 57.66
##
## Coefficients:
##
                Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                  237.90
                              10.91 21.805 5.08e-11 ***
                  -25.94
                              15.43 -1.681
                                              0.119
## Ethanol_cat1
## Ethanol_cat2
                              15.43
                                    0.119
                                              0.907
                    1.84
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 42.26 on 12 degrees of freedom
## Multiple R-squared: 0.2268, Adjusted R-squared: 0.09794
```

Instead, let's say we fit the following regression model:

What would you conclude from this model?

$$y_i = \beta_0 + \beta_1 x_i + \epsilon_i$$
$$\epsilon_{ij} \sim \mathcal{N}(0, \sigma^2)$$

where i = 1, ..., 15 is grape,  $x_i$  is the ethanol concentration used on the *i*th grape, and  $y_i$  is the PC of the *i*th grape.

How is this model different that the ANOVA model?

Let's fit a regression model.

```
model_regression = lm(PC ~ Ethanol., data = grapes)
summary(model_regression)
##
## Call:
## lm(formula = PC ~ Ethanol., data = grapes)
## Residuals:
##
     Min
             1Q Median
                           3Q
                                 Max
## -65.90 -21.55
                  0.92 24.31
                               59.50
##
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 112.800
                           65.081
                                    1.733
                                            0.1067
                                            0.0734 .
## Ethanol.
                 2.502
                            1.285
                                    1.948
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 40.62 on 13 degrees of freedom
## Multiple R-squared: 0.2259, Adjusted R-squared: 0.1663
```

What would you conclude from this output? What has changed from the ANOVA model?

#### Lesson 17 - Inference for Simple Linear Regression

Objectives:

- 1. Simulation-based inference for relationship between quantitative variables.
- 2. Theory-based approach for relationship between quantitaive variables.
- 3. Evaluate validity conditions for theory-based tests.

Why do we want to conduct inference?

Write the null and alternative hypothesis to test whether there is an association between ethanol concentration used and PC.

#### Simulation-based approach

Describe how we would conduct a simulation to conduct this test.

Let's conduct a simulation-based test (pg 291).

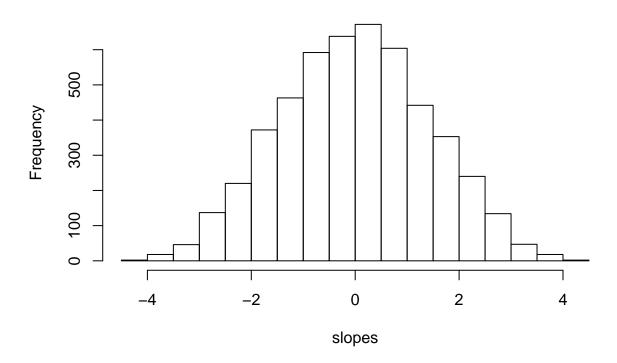
```
m = 5000 #number of iterations
slopes = c() # empty vector
grapes.sim = grapes #copy of data

for(i in 1:m){
    grapes.sim$PC.sim = sample(grapes.sim$PC) #shuffle the response
    model.sim = lm(PC.sim ~ Ethanol., data = grapes.sim) # fit model to shuffled data
    slopes[i] = coef(model.sim)[2] # extract the slope from the model
}
```

Here is a plot of the distribution of the simulated slopes:

hist(slopes)

## **Histogram of slopes**



Here is the p-value:

```
sum(slopes > coef(model_regression)[2])/m
```

## [1] 0.0402

What would we conclude?

#### Theory-based test

The *p*-value is in the linear model object:

```
summary(model_regression)
##
## Call:
## lm(formula = PC ~ Ethanol., data = grapes)
##
## Residuals:
##
     Min
             1Q Median
                           3Q
                                 Max
## -65.90 -21.55 0.92 24.31 59.50
##
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 112.800 65.081 1.733 0.1067
                            1.285 1.948 0.0734 .
## Ethanol.
                 2.502
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 40.62 on 13 degrees of freedom
## Multiple R-squared: 0.2259, Adjusted R-squared: 0.1663
## F-statistic: 3.793 on 1 and 13 DF, p-value: 0.07338
A t confidence interval for the population slope \beta_1 is (pg295):
```

```
b1 = summary(model_regression) $coefficients[2,1]
se_b1 = summary(model_regression)$coefficients[2,2]
tstar = qt(0.975, 13)
upper = b1 + tstar * se_b1
lower = b1 - tstar * se_b1
lower
## [1] -0.2732065
upper
## [1] 5.277207
# or you can just do
confint(model_regression)
##
                     2.5 %
                                97.5 %
## (Intercept) -27.7982917 253.398292
## Ethanol.
                -0.2732065
                              5.277207
```