STAT W4201 001, Homework 5

Brian Weinstein (bmw2148)

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Code is attached here and also posted at https://github.com/BrianWeinstein/advanced-data-analysis. Where relevant, code snippets and output are are included in-line.

Problem 1: Ramsey 7.18

(a) Find the standard error of prediction for the prediction of pH at 5 hours after slaughter. The calculations in Display 7.12 give us $\hat{\beta}_0 = 6.9836$, $\hat{\beta}_1 = -0.7257$, $\hat{\sigma} = 0.08226$, n = 10, $\overline{X} = 1.190$, $s_X^2 = 0.6344$.

Therefore, the standard error of prediction for the pH at 5 hours is

SE [Pred{
$$Y|X_0 = \log(5) = 1.609438$$
}] = $\hat{\sigma}\sqrt{1 + \frac{1}{n} + \frac{(X_0 - \overline{X})^2}{(n-1)s_X^2}}$
= $(0.08226)\sqrt{1 + \frac{1}{10} + \frac{(1.609438 - 1.190)^2}{9 \cdot 0.6344}}$
= 0.0875

(b) Construct a 95% prediction interval at 5 hours after slaughter. The prediction of pH level at 5 hours is,

$$Pred\{Y|X_0 = \log(5) = 1.609438\} = \beta_0 + \beta_1 \cdot \log(5)$$
$$= 6.9836 - 0.7257 \cdot 1.609438$$
$$= 5.8156.$$

A 95% prediction confidence interval at 5 hours is given by

$$5.8156 \pm t_8(0.975) \cdot \text{SE} \left[\text{Pred} \{ Y | X_0 = \log(5) \} \right]$$
$$5.8156 \pm 2.3060 \cdot 0.0875$$
$$5.8156 \pm 0.2017$$
$$\Rightarrow [5.6139, 6.0173].$$

Problem 2: Ramsey 7.24

(a) With a statistical computer package and the data in the file ex0724, obtain the least squares fits to the four simple regressions, individually, to confirm the estimates and standard errors presented in Display 7.17.

Confirming the estimates and standard errors from Display 7.17:

i. Denmark

ii. The Netherlands

iii. Canada

iv. United States

(b) Obtain the t-statistic for the test that the slopes of the regressions are zero, for each of the four countries. Is there evidence that the proportion of male births is truly declining?

The t-statistics and associated two-sided p-values are computed and shown in the output of part (a). For the Year variable:

i. Denmark

The t-statistic is -2.0726, with a one-sided p-value of 0.0221. The data provides moderate, but not convincing evidence, that the proportion of male births is truly declining in Denmark.

ii. The Netherlands

The t-statistic is -5.7102, with a one-sided p-value of 0.000000482. The data provides overwhelming evidence that the proportion of male births is truly declining in the Netherlands.

iii. Canada

The t-statistic is -4.0167, with a one-sided p-value of 0.000369. The data provides convincing evidence that the proportion of male births is truly declining in Canada.

iv. United States

The t-statistic is -5.7792, with a one-sided p-value of 0.00000720. The data provides overwhelming evidence that the proportion of male births is truly declining in the United States.

(c) Explain why the United States can have the largest of the four t-statistics (in absolute value) even though its slope is only the third largest (in absolute value).

In the hypothesis that the slopes of the regressions are zero, the t-statistic is defined as Estimate/SE(Estimate). So even though the slope is only the third largest (in absolute

value), the standard error on the estimate is small enough to make the Estimate/SE(Estimate) ratio largest for the USA.

(d) Explain why the standard error of the estimated slope is smaller for the United States than for Canada, even though the sample size is the same.

The standard error of the estimated slope is given by

$$SE(\hat{\beta}_1) = \hat{\sigma} \sqrt{\frac{1}{(n-1)s_X^2}},$$

where s_X^2 is the sample variance of the X's.

For the US and Canada, n and s_X^2 are the same (it's the same sample size, using the same subset of years), but $\hat{\sigma}$ is different.

$$\hat{\sigma} = \sqrt{\frac{\text{RSS}_j}{\text{Degrees of freedom}}} = \sqrt{\frac{\text{RSS}_j}{n-2}} = \sqrt{\frac{\text{RSS}_j}{19}},$$

where RSS_j is the sum of squared residuals for group j.

For the US, the RSS is

```
> sum((lmUsa$residuals)^2)
[1] 0.000001290857
```

and for Canada, the RSS is

```
> sum((lmCanada$residuals)^2)
[1] 0.00001120681
```

Since the RSS for the US model is an order of magnitude smaller than the RSS for the Canada model (and since all other parameters in $SE(\hat{\beta}_1)$ are identical) the standard error for the estimated slope is smaller for the US.

(e) Can you think of any reason why the standard deviations about the regression line might be different for the four countries? (Hint: The proportion of males is a kind of average, i.e., the average number of births that are male.)

There's no reason to think that the standard deviations about the regression lines would be the same for the four countries. These are samples from four different populations, and there's no reason to think their sampling distributions will have identical spreads.

Problem 3: Ramsey 7.28

(a) Is the neuron activity different in the stringed musicians and the controls?

A boxplot of the neuronal activity index for the two groups is shown in Figure 1.

The two groups have unequal variances. On the log scale, however, the variances are nearly equal, as shown below and in Figure 2.

```
> # compare group standard deviations on the original and log scales
> brainData %>%
+ group_by(Group) %>%
+ summarize(sd(Activity), sd(LogActivity)) %>%
+ as.data.frame()
    Group sd(Activity) sd(LogActivity)
1 control    2.258318    0.2983304
2 player    5.588928    0.2976929
```

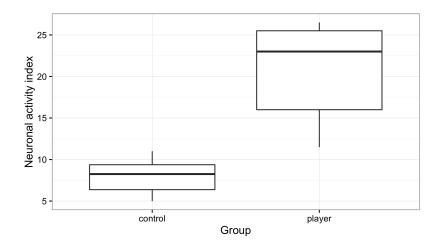


Figure 1: Boxplots of neuronal activity index (D5 dipole strength, in nA-m) for a groups of string players (player) and a group of non-string players (control).

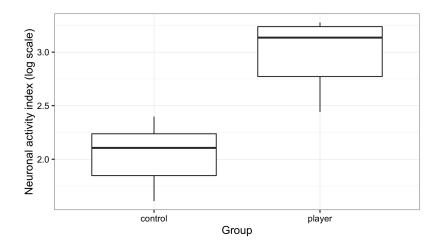


Figure 2: Boxplots of neuronal activity index (D5 dipole strength, in log(nA-m)) for a groups of string players (player) and a group of non-string players (control).

Using a two-sample t-test on the log-transformed neuronal activity index, we find overwhelming evidence that mean log activities are not equal in the two groups on the log scale (two-sided p-value: 0.00004312). Back-transforming the estimate and confidence interval from the log scale to the original scale, it's estimated that the mean neuronal activity index is 2.5731 times higher for the group of string players than the group of non-string players (95% confidence interval: 1.8329 to 3.6123 times).

(b) Is the amount of activity associated with the number of years the individual has been playing the instrument?

This is equivalent to testing if $\hat{\beta}_1 \neq 0$ in the linear regression of $\hat{\mu}\{Y|X\} = \hat{\beta}_0 + \hat{\beta}_1 X$, where Y is the neuronal activity index and X is the number of years an individual has been playing the instrument. The coefficients of the linear regression are shown below, and a plot of Years vs Activity is shown in Figure 3. The residual plot of the fitted model is shown in Figure 4.

The data provides overwhelming evidence that the amount of neuronal activity is associated with the number of years the individual has been playing the instrument. The slope of the regression line is estimated to be $\hat{\beta}_1 = 0.9971$, with a two-sided p-value (for the test that $\beta_1 = 0$) of 6.178×10^{-7} .

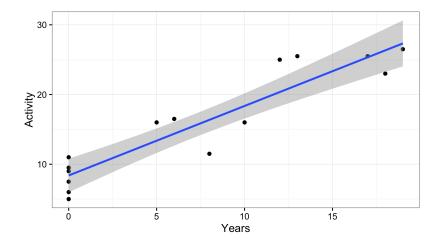


Figure 3: Scatterplot and linear regression of Activity vs Years. A 95% confidence band for the entire regression line is shown in gray.

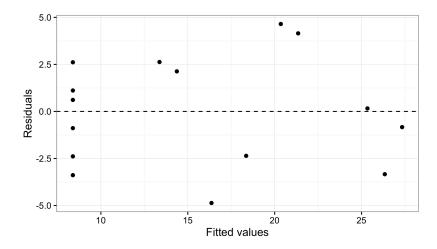


Figure 4: Residual plot for the fitted model from Figure 3.

Problem 4: Ramsey 8.17

(a) Use scatterplots of the raw data, along with trial and error, to determine transformations of Y = Ragwort dry mass and of X = Flea beetle load that will produce an approximate linear relationship.

Scatterplots of the raw data and 3 log transformations of the data are shown in Figure 5. For ease of interpretation, I only examined log-transformed variables.

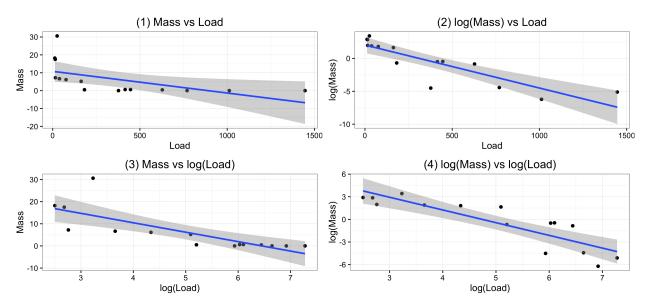


Figure 5: Scatterplots of the raw data and 3 log transformations of the data.

Log-transforming both the response (Mass) and explanatory variables (Load) produces an approximately linear relationship.

(b) Fit a linear regression model on the transformed scale; calculate residuals and fitted values. Using the 1m R function, a linear regression of log(Mass) on log(Load) is shown below.

```
> lmPest <- lm(formula=LogMass~LogLoad, data=pestData)
> summary(lmPest)
lm(formula = LogMass ~ LogLoad, data = pestData)
Residuals:
     Min
               1Q
                    Median
                                 3Q
                                         Max
-2.54217 -1.04130
                   0.06406
                           1.40544
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
               7.988
                          1.383
                                  5.774 0.0000645 ***
(Intercept)
LogLoad
              -1.685
                          0.264
                                -6.383 0.0000241 ***
Signif. codes: 0 *** 0.001 ** 0.01 * 0.05 . 0.1
Residual standard error: 1.646 on 13 degrees of freedom
Multiple R-squared: 0.7581, Adjusted R-squared: 0.7395
F-statistic: 40.74 on 1 and 13 DF, p-value: 0.00002407
```

Residuals and fitted values are shown below.

```
> residuals(lmPest)
          1
                       2
                                    3
                                                4
                                                             5
                                                                          6
                                                                                       7
-0.87164605 -0.60823442 -1.36047140
                                       0.87704266
                                                    0.06405890
                                                                1.13328141
                                                                             2.24600408
                       9
                                   10
                                               11
                                                            12
                                                                         13
 0.09207006
                                      2.01739421 -2.50142865 -1.21096053 -0.84217066
             1.67760559
                         1.82962887
         15
-2.54217409
> fitted(lmPest)
                     2
                                3
                                                        5
         1
                                                           0.6815433 -0.5954242 -0.7812252
3.7730676 3.4704353 3.3373263
                                   2.5439574
                                               1.8320606
                    10
                               11
                                           12
                                                       13
                                                                   14
                                                                              15
-2.1702639 \ -2.2916643 \ -2.8683655 \ -2.0084314 \ -3.2118881 \ -4.2738252 \ -3.6724340
```

(c) Look at the residual plot. Do you want to try other transformations? What do you suggest? The residual plot is shown in Figure 6.

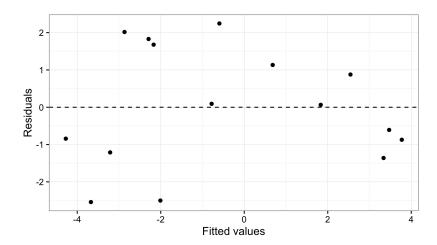


Figure 6: Residual plot for the fitted model from part (b).

There's no horn-shaped pattern present, but it does appear that even the log-transformed data has some nonlinearity — the residuals are negative at the extreme values of fitted values, but positive near the center. This suggests that the addition of a $\log(X)^2$ term may be beneficial. The residual plot does not indicate nonconstant variance or the presence of extreme outliers.

Problem 5: Ramsey 8.20

Problem 6: Ramsey 9.12