



Part 1

Question 1

Two methods were used to determine the level of E. Coli. The aim was to evaluate the agreement between HEC (in \log_{10} CFU/ml) and a more complex test HGMF (in \log_{10} CFU/ml). Portions of beef were contaminated with E. Coli and then each piece was tested by both methods. Data is contained in the file `ecoli.txt`

HEC	HGMF
0.50	0.42
0.06	0.20
0.20	0.42
0.61	0.33
0.20	0.42
0.56	0.64
-0.82	-0.82
0.67	1.06
1.02	1.21
1.20	1.25
0.93	0.83
2.27	2.37
2.02	2.21
2.32	2.44
2.14	2.28
2.09	2.69
2.30	2.43
-0.10	1.07

- Inspect the scatterplot of the data.
- Fit a model with HEC as the response
- Check if the model is appropriate and comment on the required assumptions for the data.
- Identify the outlier, which was found to be as a result of incorrect procedures.
- Make the necessary change to the data (remove the outlier) and refit the model. Then record your conclusions and comment on assumptions. Also obtain confidence and prediction intervals for a HGMF value of 1.

Hint: After loading the data (e.g. `ecoli = read.table("ecoli.txt", header = TRUE)`), the model can be fit with the code `mylm = lm(HEC ~ HGMF, data = ecoli)`.

Warning: Be careful when copying and pasting code between PDF and the R window.
The tilde character on PDF font is different to the ~ font character in R

Question 2

- Construct 95% confidence interval for the slope.
- Using this confidence interval test the hypothesis that the slope is 1.

Question 3 (to be done by hand)

Suppose `cleanEcoli` is the R object that contains the data without the outlier.

```
x = cleanEcoli$HGMF
xbar = mean(x)
n = length(x)
Sxx = (n - 1) * var(x)
c(n, xbar, Sxx)
```

```
# [1] 17.000000  1.198824 16.961176
```

Using the above summary statistics and previous regression output, answer the questions below:

- Calculate the confidence and prediction intervals for a HGMF value of 1. Compare these to the values obtained from R.
- Explain why the prediction interval is wider.
- If we had a single measurement of HGMF which is the appropriate interval for the variability in an estimate of HEC.

Question 4

Compute the Analysis of Variance for your clean regression object using the `anova` command. Show that the relationship between the t statistic from the table of coefficients and the F -test is as expected.

Hint: Suppose you typed `mycleanlm = lm(HEC ~ HGMF, data = cleanEcoli)` and `cleanEcoli` is the `ecoli` data with the outlier removed. Your clean linear regression analysis was named `mycleanlm`, to compute the simple linear regression then you can compute the ANOVA by typing `anova(mycleanlm)` to see the ANOVA table for that regression analysis.

Question 5:

Why is it not appropriate to rearrange the regression equation to give HGMF as a response? The model definition may be helpful.

Part 2

As you may have already known, **RMarkdown** is a file format that allows for the usage of both the R language, a language for statistics programming, and the **Tex** language, a language for mathematical typesetting.

Question 1

Complete the installations if you haven't done it before. Specifically, to install the **RMarkdown** components, you could type in the R console

```
install.packages("rmarkdown")
```

- For Mac users, you may be asked to install Xcode (a rather big installation). We only need a small piece of it called the `command-line tools`. Run the following line: `xcode-select --install` in the `Terminal` to continue. You should be able to find the `Terminal` tab next to the `Console` tab in `RStudio`.

Further, to incorporate the `Tex` language, you could type in the `R` console

```
install.packages("tinytex")  
tinytex::install_tinytex()
```

Question 2

`RMarkdown` can turn a combination of math formula, `R` output, and text into a `HTML` webpage. To have a try,

- a) Create a new sample `RMarkdown` file using `File > New File > RMarkdown` on the `RStudio` menu; for the `Default Output Format` option, choose `HTML`.
- b) Click on the new tab that just popped up, which by default has a name of `Untitled1`. This tab include a `RMarkdown` file. Knit this `RMarkdown` file by one of the ways below. After knitting, in the pop-up box, name the file `Test1` and put it to your working directory.
 - clicking on the `Knit` button right below the tabs
 - following the path of `File > Knit Document` on the `RStudio` menu
 - use a keyboard shortcut: `Ctrl+Shift+K` for Window & Linux and `Shift+Command+K` for Mac.

(When knitting, `RStudio` will predominantly take the current `RMarkdown` file as input and execute the code in a **new, separate R session** and therefore largely unaffected from whatever have been previously run in the `R` console, such as reading in a dataset or setting up a working directory.)

- c) Now there should be two files with the name of `Test1` in your working directory. The extension of one file should be `html`. What's the extension of the second file? Try to open this second file with `RStudio`.

Question 3

The `tinytex` package we have installed allows `RMarkdown` to export a collection of math formula, `R` output, and text into a `PDF` file. To see how,

- a) Create a new sample `RMarkdown` file using `File > New File > RMarkdown` on the `RStudio` menu; for the `Default Output Format` option, choose `PDF`.
- b) Click on the new tab that just popped up. What is there after the `output:` on the third line of this `RMarkdown` file? Is this the same as in the third line of the `RMarkdown` file in the previous question?
- c) Knit this `RMarkdown` file as in the previous question. After knitting, in the pop-up box, name the file `Test2` and put it to your working directory. Then try to find the file `Test2.pdf` in your working directory.