

Homework 2 Instructions

FISH 552 - Kristin Privitera-Johnson

10/06/2021

Part 1 - Programming Assignment

Open a new script in RStudio and create an assignment header using comments.

```
# Name: First Last  
# Homework 1
```

Complete the tasks below. Please label each question and task with comments.

```
##=Question 1a ####  
# Your R code here  
  
##=Question 1b ####  
# Your R code here
```

When your script is complete, save it as **LastName_Homework2.R**, then clear your workspace (Workspace/Clear all) and run through your script again to make sure you don't have any **object not found** errors. Then go to the course website to submit your R script.

Question 1

This question will use the dataset **beaver1**, which is built into R. Run the command **head(beaver1)** to get a sense of what is in the dataset, and go to the help file **?beaver1** to learn more about this data set. Using this dataset:

- Use one function to compute the mean temperature for inside and outside the retreat.
- Recreate the following plot. Note the x-axis, and the green points mark activity outside the retreat.
 - Plot, labels, and title
 - Axis
 - Points
 - Horizontal lines
 - Legend
- Note that the observation at 22:20 is missing. Create a new data frame that includes an additional observation with a temperature of 37.3 and activity outside the retreat at this time.
- How much did the mean temperature during periods of activity outside the retreat change? (i.e. compare **beaver1** and your new data frame)

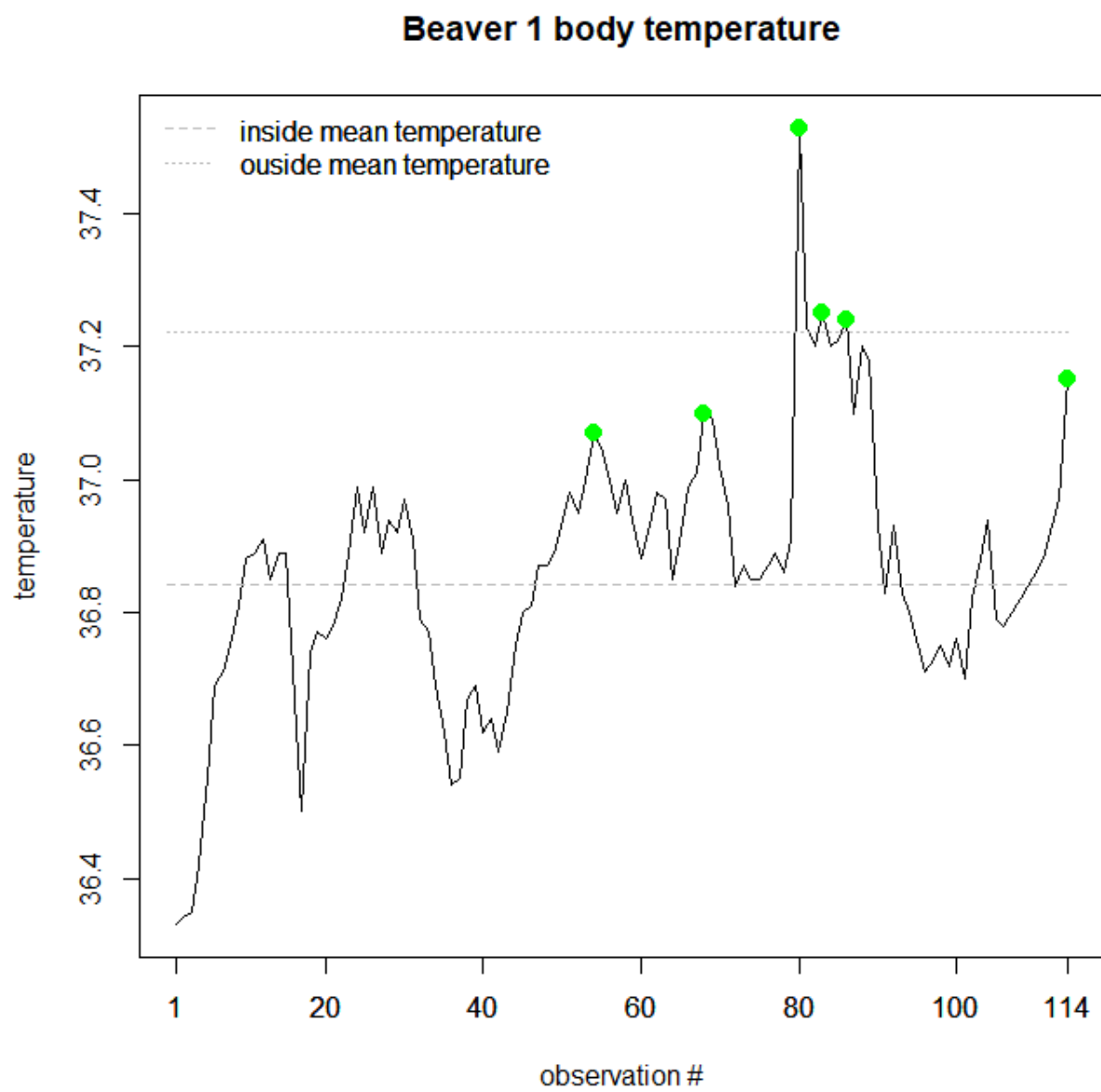


Figure 1: Plot to recreate.

Question 2

This question will use the dataset `crabs`, which is built into R. Type the following command to get access to the dataset (we'll cover what a library is later):

```
library(MASS)
?crabs
head(crabs)
```

Now use the dataset `crabs` to complete the following:

- Use **one function** to compute the mean of frontal lobe size for “blue” and “orange” purple rock crabs. Now compute the standard deviation of frontal lobe size for “blue” and “orange” purple rock crabs.
- Run this line of code: `crabs$sp:crabs$sex`. Explain what's going on here by adding comments to your code above this line of code. Call this new factor `spsex`.
- Change the levels of `spsex` so that they are more informative. So B:F might be called “Blue Female” and so on.
- Use **one function** to compute the mean of frontal lobe size for each sex and color combination of purple rock crab.
- Use **one function** with the vector `spsex` to compute the total number of each sex and color combination of purple rock crab. Call this vector `crab.counts`. Hint: `?table`.
- Create an informative plot using the `crabs` data with appropriate labeling. You may plot part of the data or all of the data, but you are required to create a multipanel plot with at least two dimensions using `layout()` or `par(mfrow)`. Be creative!