# AFS505 Module 1 Final Exam

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## Question 1

What are the basic R data structures? What are the differences between them? In what context would you use one versus the other

The different basic data types in R are Characters, Numeric (both real and decimal), Integers, Logical (TRUE/FALSE), and complex.

Common basic data structures are vectors, Matrices, Lists, and Data Frames.

A vector is a sequence of data elements that must all be the same type. Vectors are technically one dimensional arrays. I would use a vector to store a simple collection of related data. Lists are a kind of vector technically, but most of the time I work with atomic vectors where the data type is restricted. One of the ways I often use vectors is to populate the data for matrices since the mat() function takes a data vector, character string, or list as its arguments. If you attempt to mix different data types in a vector r will coerce the data into the same type.

A matrix is a collection of data elements arranged into a two dimensional rectangular layout with both rows and columns. In this way they are a natural extension of atomic vectors with the additional quality of having dimensionality. Vectors and Matrices require that the data you include be the same data type. A vector is what you should use if your data is one dimensional. You would use a matrix if your data was multidimensional since a matrix has both columns and rows to organize data into.

A data frame is a list of vectors of equal length. it can handle data of different types if you have a mixture of string, logical, and numerical variables. If you have data with different types you should use a list or a data frame instead of a vector/matrix. A data frame is basically a data table. I use data frames to organize and interpret data sets with different variables. The function to fit linear models takes data frames as potential arguments so data frames are good for regression analysis. I often default to data frames for cleaning my data by assigning subsets of old data frames to new data frames however this practice can be cluttered and waste valuable memory. Data frames have attributes like rownames() which is helpful for enhancing the readability of your code. The read family of functions (read.csv or read.table for example) output data frames which is perhaps one of the most important uses of the data frame structure.

A list is the most generic data type. It is basically a vector of objects which can be any type of r object even data structures. A list can even have other lists as its elements. Lists are incredibly useful for working with certain functions in the apply family like sapply and lappy which can take lists or vectors as arguments. The base apply function takes a matrix as an input. Beyond "apply" the way functions in R work is very compatible with lists since functions return single objects a series of function results can be banded together in a list for a function to return.

## Question 2

You are provided a folder with three location (county) names, each of which has subfolders for one or two crops, which in turn has a data file.

I think it's easiest to combine steps a and b for question 2

#### Part a and b

a iterate through the folders to deal all the files and merge them into a single data frame. You can use a "loop" to iterate or for efficiency check out the list.files() function

b add four additional columns to the merged dataframe corresponding to the county name, crop name, latitude and longitude of the data. You must get this information from the directroy structure you are looping through or the strings returned by the call to list.files()

```
### use list files function to get string vectors of the filenames
### we will use fn as shorthand for filenames
fnlocations <- list.files(path = "C:/Users/otten/Desktop/AFS Repository/Will-O-AFS505/CropModelResults"
print(fnlocations)
## [1] "Okanogan"
                     "WallaWalla" "Yakima"
fncrops <- list.files(paste0("C:\\Users\\otten\\Desktop\\AFS Repository\\Will-O-AFS505\\CropModelResult</pre>
print(fncrops)
## [1] "Corn grain"
                      "Winter Wheat"
fncords <- list.files(paste0("C:\\Users\\otten\\Desktop\\AFS Repository\\Will-O-AFS505\\CropModelResult</pre>
print(fncords)
## [1] "48.15625N119.71875W" "48.96875N119.65625W"
### Create an initial data frame to rbind the rest of the data frames to when we read them
### In this step we must also create the new columns appropriately using information from the list.file
Q2maindf <- read.csv(paste0("C:\\Users\\otten\\Desktop\\AFS Repository\\Will-O-AFS505\\CropModelResults
lllist <- strsplit(fncords[1],"N")</pre>
latlong <- lllist[[1]]</pre>
latlong[1] <- pasteO(latlong[1],"N")</pre>
latlong
## [1] "48.15625N" "119.71875W"
Q2maindf$countyname <- fnlocations[1]
Q2maindf$cropname <- fncrops[1]
Q2maindf$latitude <- latlong[1]</pre>
Q2maindf$longitude <- latlong[2]
### All of this is to make the code general so as to not hard code the known length of the file lists
loclength <- length(fnlocations)</pre>
croplength <- length(fncrops)</pre>
cordlength <- length(fncords)</pre>
### Create a for loop to cycle through each location
### within that loop create a for loop to cycle through each crop
### within that loop create a for loop to cycle through each lat/long
### The loop should read a csv file and store it as a data frame. The loop should then create new colum
for (a in 1:loclength) {
  for(b in 1:croplength){
    for (c in 1:cordlength) {
       fncords <- list.files(paste0("C:\\Users\\otten\\Desktop\\AFS Repository\\Will-0-AFS505\\CropMode</pre>
       tempdf <- read.csv(paste0("C:\\Users\\otten\\Desktop\\AFS Repository\\Will-O-AFS505\\CropModelRe</pre>
```

```
### I like to print a piece of my loop just so I can see it is working while I run the code... T
       print(fncords[c])
       lllist <- strsplit(fncords[c],"N")</pre>
       latlong <- lllist[[1]]</pre>
       latlong[1] <- paste0(latlong[1],"N")</pre>
      tempdf$countyname <- fnlocations[a]</pre>
      tempdf$cropname <- fncrops[b]</pre>
      tempdf$latitude <- latlong[1]</pre>
      tempdf$longitude <- latlong[2]</pre>
      Q2maindf <- rbind(Q2maindf,tempdf)</pre>
    }
  }
}
## [1] "48.15625N119.71875W"
## [1] "48.96875N119.65625W"
## [1] "48.15625N119.71875W"
## [1] "48.53125N119.59375W"
## [1] "46.03125N118.21875W"
## [1] "46.28125N118.65625W"
## [1] "46.03125N118.21875W"
## [1] "46.03125N118.40625W"
## [1] "46.15625N119.96875W"
## [1] "46.21875N119.34375W"
## [1] "46.15625N119.34375W"
## [1] "46.21875N119.34375W"
### remove the duplicate values since one file was read twice
Q2maindf <- Q2maindf[!duplicated(Q2maindf),]</pre>
```

### Part c

c Rename the column irrig to irrigation\_demand and precip to precipitation and export the dataframe as a csy file

```
### Rename the columns
colnames(Q2maindf)[6] = "irrigation_demand"
colnames(Q2maindf)[7] = "precipitation"

### Export the data frame
write.csv(Q2maindf, "C:\\Users\\otten\\Desktop\\AFS Repository\\Will-O-AFS505\\ManagedCropData.csv")
```

### Part d

d summarize the annual irrigation demand by crop name and county name

```
### To summarize by other variables I can convert the character data into factor data
Q2maindf$countyname <- as.factor(Q2maindf$countyname)
Q2maindf$cropname <- as.factor(Q2maindf$cropname)

### summarize irrigation demand by county name
tapply(Q2maindf$irrigation_demand,INDEX = Q2maindf$countyname, summary)

## $Okanogan
## Min. 1st Qu. Median Mean 3rd Qu. Max.</pre>
```

```
##
     272.0
             363.6
                      392.0
                               390.8
                                       416.7
                                                490.5
##
   $WallaWalla
##
##
      Min. 1st Qu.
                     Median
                                Mean 3rd Qu.
                                                 Max.
##
     180.0
             300.0
                      360.0
                               357.0
                                       417.5
                                                520.0
##
## $Yakima
##
      Min. 1st Qu.
                     Median
                                Mean 3rd Qu.
                                                 Max.
##
     390.0
             520.0
                      560.0
                               545.9
                                       580.0
                                                640.0
### summarize irrigation demand by crop name
tapply(Q2maindf$irrigation_demand, INDEX = Q2maindf$cropname, summary)
## $Corn_grain
##
      Min. 1st Qu.
                     Median
                                Mean 3rd Qu.
                                                 Max.
##
     180.0
             360.0
                      413.3
                               422.4
                                       480.0
                                                640.0
##
##
   $Winter_Wheat
##
      Min. 1st Qu.
                     Median
                                Mean 3rd Qu.
                                                 Max.
##
     208.0
             363.9
                      420.1
                               440.3
                                       540.0
                                                640.0
```

#### Part e

e What is the average yield of Winter Wheat in Walla Walla at 46.03125N118.40625W for the year ranges (1981-1990), (1991-2000), and (2001-2019)?

The instructions don't specify how I need to calculate this information so I am using excel to filter the data quickly and report the average. This is easy because earlier in the exam I exported the data frame as a csv file

For the crop at that location during the years 1981-1990 the average yield is 7660.89502

For the Crop at that location during the years 1991-2000 the average yield is 8086.688721

For the Crop at that location during the years 2001-2019 the average yield is 7720.68303

#### Part f

f which location has the highest yield (average) for the time period (2001-2019) for grain corn

Again because the instructions don't specify I must do this in r I will make the calculations in excel

```
### I will create a subset of my data to figure this out first I am only interested in values related t
library(stringr)

Q2fsub <- subset(Q2maindf ,Q2maindf$cropname == "Corn_grain")

### Secondly I am only interested in the years 2001-2019. It will be easier for me to subset this data

Q2fsub$YYYY.MM.DD.DOY. <- str_trunc(Q2fsub$YYYY.MM.DD.DOY., 4,"right", ellipsis = "")

Q2fsub$YYYY.MM.DD.DOY. <- as.numeric(Q2fsub$YYYY.MM.DD.DOY.)

Q2fsub <- subset(Q2fsub, Q2fsub$YYYY.MM.DD.DOY.) >= 2001)

### I will coerce the location (lat long) into factors so that I can calculate the means for each area

Q2fsub$latitude <- as.factor(Q2fsub$latitude)

Q2fsub$longitude <- as.factor(Q2fsub$longitude)</pre>
```

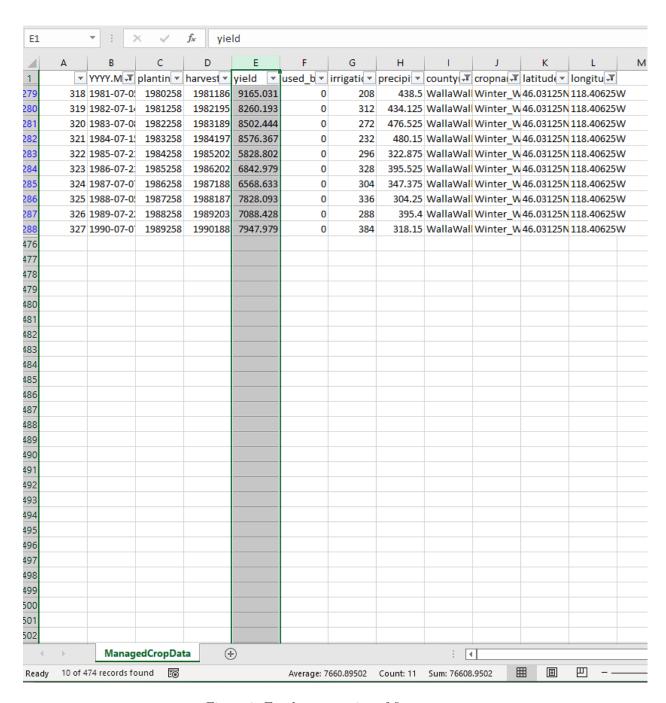


Figure 1: Excel computation of first average

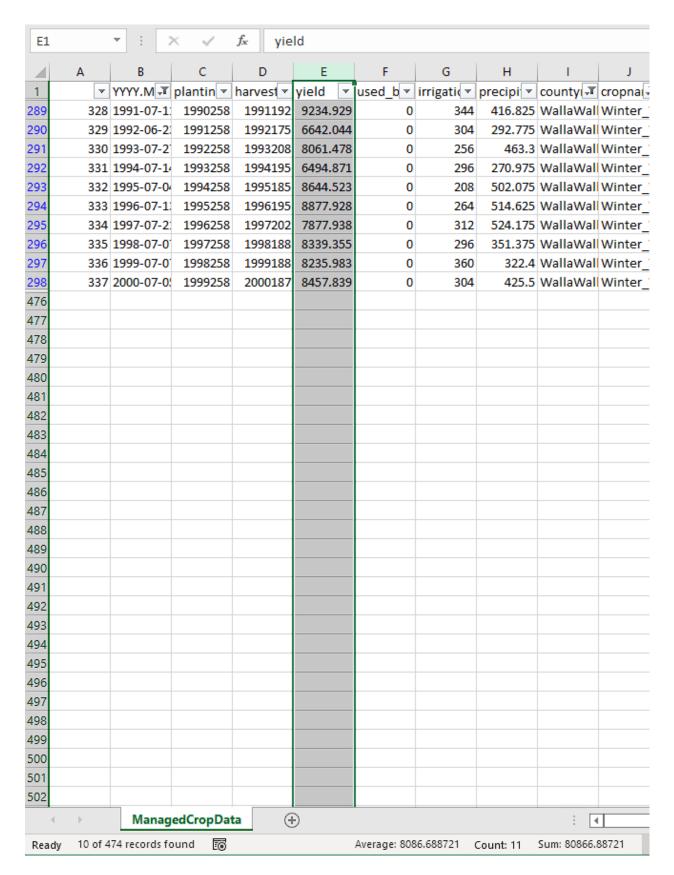


Figure 2: Excel computation of second average

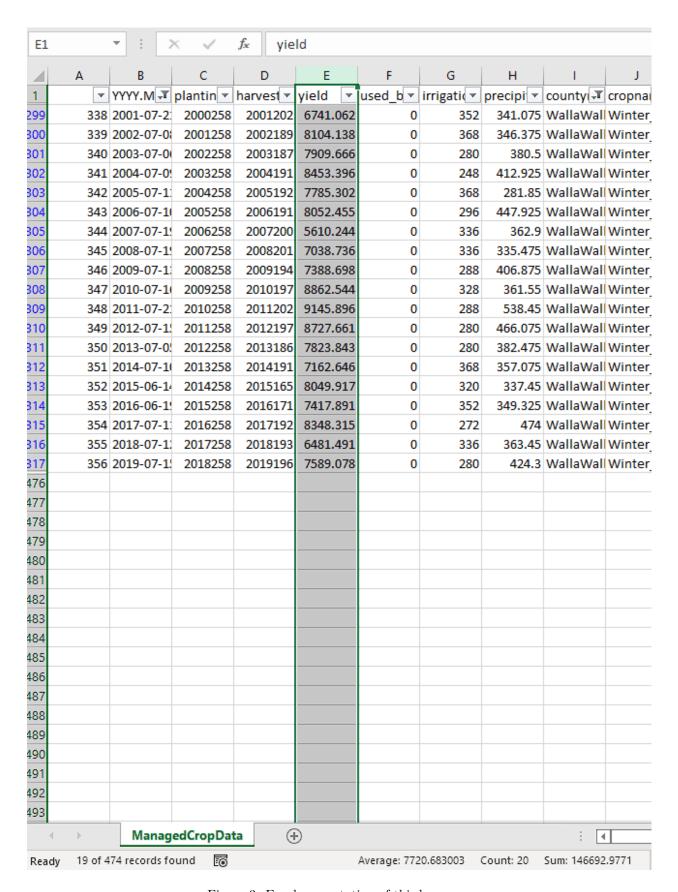


Figure 3: Excel computation of third average

```
### I use the tapply function to compute means
tapply(Q2fsub$yield, INDEX = Q2fsub$latitude, mean)

## 46.03125N 46.15625N 46.21875N 46.28125N 48.15625N 48.96875N

## 9981.876 13536.885 15263.030 12482.990 14092.233 14139.643

tapply(Q2fsub$yield, INDEX = Q2fsub$longitude, mean)
```

```
## 118.21875W 118.65625W 119.34375W 119.65625W 119.71875W 119.96875W ## 9981.876 12482.990 15263.030 14139.643 14092.233 13536.885
```

It is apparent that 46.21875N119.34375W has the highest average yield for grain corn in the period 2001-20019 with an average of 15263.030

#### Github link

Repository link: https://github.com/Will-Ottenheimer/Will-O-AFS505.git

Rmarkdown Permalink: https://github.com/Will-Ottenheimer/Will-O-AFS505/blob/a240a28cfffae1349c2ad 571714fa9d61144dac9/AFS505%20Final%20Exam.Rmd

Rmarkdown PDF Permalink: https://github.com/Will-Ottenheimer/Will-O-AFS505/blob/a240a28cfffae13 49c2ad571714fa9d61144dac9/AFS505-Final-Exam.pdf

 $CSV\ File\ Permalink:\ https://github.com/Will-Ottenheimer/Will-O-AFS505/blob/4f6e32785add3316e0c4f0\ 13e6d832c7e1244ae9/ManagedCropData.csv$ 

### Question 3

Was the data provided to you well described? If not, what information was missing? Comment on what kind of meta data (description about the data) should be included as a best practice while sharing data sets.

The data was poorly described. To begin the unit of measurement is missing for nearly all the variables. I don't know if yield is in bushels or grams, or truckloads so the meta data file should explain the measurement unit for all the numeric data. The Used\_biomass is not labeled as a dummy variable (if it is it should be). The meta data should express what format the planting and harvesting date are in.

The best meta data includes the date the data was created and the date it was last modified. Metadata should also include the name of the author of the data set and their contact info. Additionally, quality metadata should have tags and categories and titles and descriptions for all the data. Databases in particular should include information about the data types, columns, constraints, and relationships between the tabular data.

### Question 4

Create an R Markdown file with different tabs for each of the six parts of question 2. In a seventh tab add the github link which has your Rscript/R markdownfile and the csv file generated from 2(c)

Because I did steps a and b of question 2 in one step there will only be 6 tabs.