STAT 302: Statistical Computing

Winter 2024

Lecture 7: Midterm Review

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This note is designed to give a brief review on lecture materials [Zhang, 2023] and highlight those important concepts/results in STAT 302 before the midterm exam. The review is by no means comprehensive and in order to excel at the midterm exam, a student is expected to master those R fundamentals in the course instead of simply memorizing the key functions or commands in R.

Disclaimer: Parts of this note are modified from the midterm review of STAT 133 in Fall 2016 at UC Berkeley by Professor Deborah Nolan.

1 R Basics and Syntax

- Binary arithmetic operators: +, -, *, /, %% (for mod), %/% (for integer division), and ^ (exponentiation), etc.
- Comparison operators: >, <, <=, >=, and !=. The returning value of a comparison statement is TRUE or FALSE.
- Logical operators: &, |, and !. A logical operator takes one or more comparison statements and return TRUE or FALSE.

 \star Notes: Pay attention to the order/priority of operations as well as the difference between simple and compound expressions.

Data Types in R:

- Basic types in R: numeric (and integer), character, logical, factor, etc.
- Factor has a level (string) and is stored internally as an integer.
- Special values include NA (not available), NaN (not a number), NULL (empty vector), and Inf (infinity).

We can use the built-in function class() to determine the data type of an R object.

 \star Note: The function typeof() determines the (R internal) type or storage mode of any R object. Its returning value may not be the same as class().

2 Data Structures in R

Vector calculations:

- Vector is an ordered container of primitive elements of the same data type.
- R is vectorized so that we can perform operations on vectors.
- Remember the recycling rule in R when the vectors are of different lengths:

> x = rep(c(1,2), times = c(2, 4))
> y = 1:3
> x^y
[1] 1 1 8 2 4 8

Matrices and arrays: A matrix or an array is a rectangular collections of primitive elements of the same data type. They are stored in R as a vector with shape information.

Lists: A list is an ordered container of objects, e.g., mixtures of vectors, data frames, lists, functions.

Data frames: A data frame is an ordered container of vectors, where the vectors are of the same length and possibly different data types.

Practice Problem: Describe two important differences between a data frame and a matrix in R.

Answer: Here are some acceptable answers.

- A data frame is essentially a list of vectors of the same length, whereas a matrix is essentially a vector with shape information.
- Data frames can have columns/vectors that are different types, whereas all values in a matrix must be the same primitive element.
- Data frames can be indexed with \$.
- Data frames require variable names.

Subsetting: We use the family.txt data as an example.

```
> head(family)
firstName sex age height weight
                                        bmi overWt
                 77
                          70
                                 175 25.16239
                                                 TRUE
1
        Tom
               m
2
       Maya
               f
                  33
                          64
                                 124 21.50106
                                                FALSE
3
        Joe
               m
                 79
                          73
                                 185 24.45884
                                                FALSE
4
                  47
                          67
                                 156 24.48414
                                                FALSE
     Robert
               m
5
        Sue
               f
                  27
                          61
                                  98 18.51492
                                                FALSE
6
        Liz
               f
                  33
                          68
                                 190 28.94981
                                                 TRUE
> sapply(family, class)
firstName
                   sex
                                           height
                                                        weight
                                                                         bmi
                                                                                   overWt
                                 age
"character" "character"
                            "integer"
                                          "integer"
                                                       "integer"
                                                                    "numeric"
                                                                                  "logical"
```

• Position – using the indices of elements, rows, columns that we want.

> family[,1] [1] "Tom" "Maya" "Robert" "Sue" "Joe" "Liz" "Jon" "Sally" "Tim" [10] "Tom" "Ann" "Dan" "Art" "Zoe" > family[c(1, 3, 6),] firstName sex age height weight bmi overWt 1 Tom 77 70 175 25.16239 TRUE m 3 Joe 79 73 185 24.45884 FALSE m 6 Liz f 33 68 190 28.94981 TRUE

• Exclusion – using the indices of elements to exclude.

<pre>> family</pre>	[-(<mark>2</mark> :1	L2),]				
firstName	e sex	age	height	weight		bmi ove:	rWt
1	Tom	m	77	70	175	25.16239	TRUE
13	Art	m	46	66	150	24.26126	FALSE
14	Zoe	f	48	62	125	22.91060	FALSE

• Logical – using a logical vector of the same length as the vector/rows/columns being subsetted. It will keep the elements corresponding to TRUE.

```
> family[(family$overWt) & (family$age > 60), ]
firstName sex age height weight bmi overWt
1 Tom m 77 70 175 25.16239 TRUE
7 Jon m 67 68 185 28.18797 TRUE
```

• Name – using a character vector of names of elements/rows/columns to keep.

> family[, "age"] [1] 77 33 79 47 27 33 67 52 59 27 55 24 46 48

 \star Note: Subsetting a list or data frame via a double-square bracket returns the element or column within the list or data frame, but not a list or data frame of one element.

```
> class(family[["age"]])
[1] "integer"
> class(family["age"])
[1] "data.frame"
```

Practice Problem: Write down what the value of x will contain after each line of R code, if the commands are executed sequentially.

```
> x = seq(0, 8, length = 5)
> x
[1] 0 2 4 6 8
> x[x<4] = NA
> x
[1] NA NA 4 6 8
> x[5] = 10
> x
[1] NA NA 4 6 10
> x[] = 0
> x
[1] 0 0 0 0 0
> x = 12
> x
[1] 12
```

Practice Problem: Data on 37 parents of babies born at Kaiser Hospital in the 1960s is available in a data frame called **parents**. The variables age, ed, ht, and wt are the mother's age, education level, height and weight. The variables that start with the letter d are corresponding variables for the fathers.

>	> head(parents)								
	age	e ed	ht	wt d	lage	e ded	dht	dwt marital	inc
1	27	College	62	100	31	College	65	110 Married	[2500, 5000)
2	33	College	64	135	38	College	70	148 Married	[7000, 8000)
3	28	High School	64	115	32	Some High School	l NA	NA Married	[5000, 6000)
4	36	College	69	190	43	Some College	68	197 Married	[12500, 15000)
5	23	College	67	125	24	College	NA	NA Married	[2500, 5000)
6	25	High School	62	93	28	High School	64	130 Married	[7000, 8000)

1. Write an R expression to find the subset of parents where the mother is over 40.

```
parents[parents$age > 40, ]
```

2. Write an R expression using an apply function to return the class of each variable in the data frame.

```
sapply(parents, class)
```

3. Write one R expression using an apply function to return the number of NAs in each variable (recall that there is a function called *is.na(*) that returns a logical indicating the presence of NAs).

```
sapply(parents, function(x){
   sum(is.na(x))
})
```

Practice Problem: Here is a list in R:

```
> x
$a
[1] 0.03895442 0.77658866 0.83532332
$b
[,1] [,2]
[1,] 1 4
[2,] 2 5
[3,] 3 6
```

Write one line of R code to extract the first row of the matrix.

```
x$b[1, ]
```

x[["b"]][1,]

* Notes: You should know how to use dim(), head(), class(), length() functions with the data structures in R. In addition, you should also be able to use sapply(), lapply(), apply(), tapply(), and mapply() functions with these structures.

Practice Problem: Suppose we have a matrix m, and we have just executed the following:

> dim(m) [1] 500 3

> head(m)							
	[,1]	[,2]	[,3]				
[<mark>1</mark> ,]	0.9231213	0.2142767	2.544545				
[2,]	0.5669828	0.6981698	1.336078				
[<mark>3</mark> ,]	0.7077386	0.1447157	1.750822				
[4,]	0.6723225	0.6040853	0.042359				
[5,]	0.4925304	-0.4389639	1.504622				
[6,]	0.8370886	0.3683489	1.456362				

We need to create a vector containing the sum of the squared entries in each row of m. Use an anonymous function and the apply() function to do this:

```
apply(m, 1, function(x) {
   sum(x^2)
})
```

3 Control Flow and Basic Iterations

Conditional evaluation: if and if-else statements.

```
if (condition to evaluate to TRUE or FALSE) {
  code block to be run if above is TRUE
}
and
if (condition to evaluate to TRUE or FALSE) {
  code block to be run if above is TRUE
} else {
  code block to be run if above is FALSE
}
```

Iterations: Looping with for and while statements.

```
> x = c(10, 2, 6, 5)
> for (i in 1:3) {
    print(x[i])
  }
[1] 10
[1] 2
[1] 6
```

4 Data Manipulation via tidyverse

Pipes %>% operator:

- $x \gg f \gg g$ is equivalent to g(f(x)).
- $x \ \text{\sc{k}>\sc{k}} f(y, .)$ is equivalent to f(y, x).

dplyr functions: Some of the most important dplyr verbs (functions):

- filter(): subset rows based on a condition.
- group_by(): define groups of rows according to a column or specific condition.
- summarize(): apply computations across groups of rows.
- arrange(): order rows by value of a column.
- select(): pick out given columns.
- mutate(): create new columns.
- mutate_at(): apply a function to given columns.

tidyr functions: Two of the most important tidyr verbs (functions):

- pivot_longer(): make "wide" data longer.
- pivot_wider(): make "long" data wider.

5 Graphics in R

Data type and plot correspondence:

- One variable: factor: barplot, dot chart, line plot, etc. numeric: rug plot, density plot, histogram, boxplot, etc.
- Two variables:

Two factors: side-by-side bar plot, superposed line plots, mosaic plot, side-by-side dot chart, etc. Two numerics: scatter plot, line plot if one variable is time numeric + factor: super-posed density curves, side-by-side boxplots or violin plots, line plots, etc.

• More details can be found in the ggplot2 cheet sheet and the reference book.

Plotting principles:

- Make the data standout: symbols, over plotting, interference, jittering, scale, etc.
- Facilitate comparison: emphasize the important difference; don't jiggle the baseline; juxtapose, superpose and choice of scale; banking to 45 degrees; perception color, shape.
- Information rich: color; reference lines and markers; legends and labels; captions.

ggplot essentials:

- ggplot() initiates a plot. We can provide a data frame and a mapping between the variables in the data frame and plot aesthetics.
- Layers we add geoms/glyphs to the plot with geom_XXX() functions. Examples include

geom_point() for points in a scatter plot,

geom_line() for connecting dots to make line plots,

geom_smooth() for averaging the y-values with similar x-values to create a smooth curve as a function of x,

geom_bar() for making bars in a bar plot,

geom_boxplot() for making boxes in a box plot,

geom_density() for computing and drawing kernel density estimate, which is a smoothed version of the histogram,

geom_histogram() for making bars in a histogram.

In the layer, we can also provide the data frame and the mapping of the variables to aesthetics.

- Scales these are associated with the aesthetics. Each aesthetic has a scale. We use the scale_XX_xxxx()
 functions to specify details. The first argument in each scale function is name, which provides the axis
 or legend name. A short cut function called labs() allows use to specify the name to all of our aesthetics without having to call the scale functions, e.g., labs(x = "my x", y = "my y", color = "color
 legend title").
- Theme themes are used for general details, such as font type and size, background color, etc.

6 Writing Functions in R

Function signatures:

- Required parameters have no default value provided, usually come first in the signature.
- Optional parameters have default values (which may be computed from other parameters).
- Function calls R sets up the call frame with default values for parameters, then values are assigned for the arguments mentioned by name in the call. R will first look for the value of a variable by name in the function environment. If it can't be found, then it will resort to its parent environment, which is the global environment in this case, etc.
- <<- is used to assign values to variables in the parent environment.
- Lazy evaluation If an expression is provided for a parameter value, then this expression is not evaluated until the variable is referenced in the computations for the function call.
- Debugging tools and testing traceback(), browser(), debug(), etc.

Practice Problem: Write a function to compute the sum of the absolute deviations from the median for an input vector. For example, for a vector $\mathbf{x} = 1:3$, \mathbf{x} has a median of 2, and the absolute deviations from the median are 1, 0, and 1 so the sum of the absolute deviations from the median is 2.

Call the function \mathtt{sadm} (). This function has one parameters: x, which is required and holds the numeric vector that will be operated on.

```
sadm = function(x){
  res = sum(abs(x - median(x)))
  return(res)
}
```

Practice Problem: Update the function sadm() to handle NA values. Add a parameter: na.rm, which determines whether NAs are to be removed from the computation. The na.rm parameter has a default value of FALSE.

```
sadm = function(x, na.rm = FALSE){
    if (na.rm) {
        x = x[!is.na(x)]
    }
    res = sum(abs(x - median(x)))
    return(res)
}
```

7 Simulation Basics

- Simple random selection with sample()
- Replicating the experiment with replicate().

Practice Problem: Someone wants to study the distribution of the sum of three rolls of a die. To do this, he/she designs a simulation study. In the first step, he/she writes a function called sum3() to generate the sum of three random tosses of a fair die. In the second step, he/she uses this function to generate 1,000 of these sums.

1. Write the function for the first step.

```
sum3 = function(){
  res = sum(sample(1:6, 3, replace = TRUE))
  return(res)
}
```

2. Write one line of code that uses the function from the first step to generate the 1,000 random sums.

```
replicate(1000, sum3())
```

References

Y. Zhang. STAT 302 Lecture Slides, 2023. URL https://zhangyk8.github.io/teaching/stat302_uw.