# GSP Workshop: Intermediate R: Statistics for Graduate Students

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#### Part I

- Conditionals and Control Work flows
  - ► Equality (or not)
  - & and
  - ► The if statement

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- Loops
  - Write a while loop
  - Write a for loop
  - Mix up loops with control flow

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#### Part II

► Functions and Arguments: Review

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#### Part II

- Functions and Arguments: Review
- Subset functions in R
  - Demonstration with data set
  - Resources for developing packages for tidying data

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  - & and
  - ▶ The if statement
- Loops
  - ► Write a while loop
  - Write a for loop
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Link to slides: Intermediate-R

#### Part II

- Functions and Arguments: Review
- Subset functions in R
  - Demonstration with data set
  - Resources for developing packages for tidying data
- Advanced: Tidyverse

## Prelude: R Markdown

R Markdown provides an patform for data science. In an R Markdown file you can both

- Both save and execute code
- create reports and/or articles that can be shared to various audiences.

#### First:

```
'''{r}
install.packages("rmarkdown")
'''
```

Here is a link to the code used: R Code
Here is a link to an R Markdown cheet sheet: R Markdown

A **logical statement** is a declarative sentence which conveys information about the truth of a statement.



Figure: Figure credit: Boolean

A **logical statement** is a declarative sentence which conveys information about the truth of a statement.

Examples 1 and 2: Equality (or not): If x = 3, y = 4 then we can assess the truth about the following statements:

```
> #Example 1: Equality
>
> x=3
> y=4
> x==y
[1] FALSE
```

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```
> #Example 1: Equality
>
> x=3
> y=4
> x==y
[1] FALSE
```

```
> #Example 2: Not equality
>
> x!=y
[1] TRUE
> |
```

- Example 3: Equality (or not): Comparison of strings.
  - Set the two vectors below equal to each other
  - ► This creates a logical vector (with TRUE/FALSE)

```
user1<-c("userR", "user", "UserR")
user2<-c("userR", "useR", "UserR")
```

user1==user2 #this is a logical vector
[1] TRUE FALSE TRUE

- Example 3: Equality (or not): Comparison of strings.
  - Set the two vectors below equal to each other
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user1<-c("userR", "user", "UserR")
user2<-c("userR", "useR", "UserR")
```

user1==user2 #this is a logical vector [1] TRUE FALSE TRUE

**Demonstration:** Let's use logical operators to subset our data.

► Example 5: The greater than (>), and less than (<) operators. Dealing with unusual points:

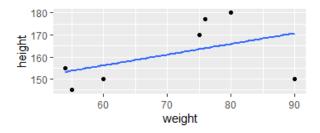


Figure: There appears to be an unusual point at (90, 150)

► Example 5: The greater than (>), and less than (<) operators. Dealing with unusual points:

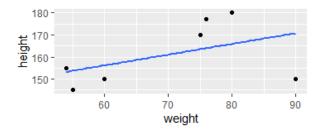


Figure: There appears to be an unusual point at (90, 150)

. Demonstration: Let's use logical operators to subset our data.

► Example 6: The & and | operators. Dealing with unusual points:

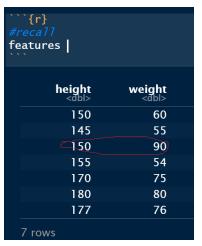


Figure: There appears to be an unusual point at (90, 150)

Example 6: The & and | operators: outlier <-features[ which(weight==90 & height==150), ] outlier

```
features3<-features[ which(weight==90 | height==150), ]
features3</pre>
```

Example 6: The & and | operators: outlier <-features[ which(weight==90 & height==150), ] outlier

features3<-features[ which(weight==90 | height==150), ]
features3</pre>

(a) The subset with & operator

(b) The subset with | operator.

Figure: Data frames created from the AND and OR logical operators

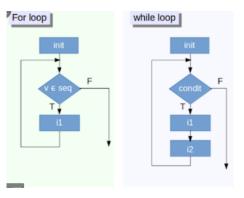


Figure: Credit: A figure of a for and while loop loops

A **loop** is a replication of instructions. It's a mini-step process that organizes a sequence of actions into parts that need to be repeated. Diamonds, on the other hand, are called **decision symbols**, and translate into questions which only have two possible logical answers; TRUE or FALSE.

#### Example 7: For loop

- Start with an intialization (e.g. vector or matrix) of empties.
- ► Tests to whether a current value is within a specified defined range (i.e. within 1:100).
- ▶ If the condition is not met and the resulting outcome is False, the loop is never executed.

#### Example 7: For loop

- Start with an intialization (e.g. vector or matrix) of empties.
- ► Tests to whether a current value is within a specified defined range (i.e. within 1:100).
- ▶ If the condition is not met and the resulting outcome is False, the loop is never executed.

```
set.seed(1)
rv <- rnorm(1000, 0, 1)
usq<-matrix("NA", 100, 1)

for(i in 1:100) {
  usq[i] <- rv[i]*rv[i]
  print(usq[i])
}
  usq<-data.frame(usq=as.numeric(unlist(usq)))
attach(usq)</pre>
```

#### **Example 7: For loop**

Once the condition is past 100, the evaluation is False, and the loop ends:

```
> dim(usq)
[1] 100 1
>
```

▶ The data frame is entries for the \_\_\_\_\_ distribution.

#### Example 7: For loop

Once the condition is past 100, the evaluation is False, and the loop ends:

```
> dim(usq)
[1] 100 1
>
```

- •
- ▶ The data frame is entries for the \_\_\_\_\_ distribution.
- ▶ The data frame is entries for the <u>chi</u> distribution (denoted  $\chi_1^2$ ).

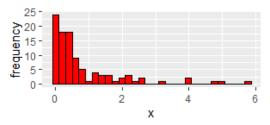


Figure: The square of a standard normal is chi

#### **Example 8: While loop**

- ▶ Start with an intialization (e.g. vector or matrix) of empties.
- Now followed by a logical comparison between a control variable and a value (using earlier defined conditionals).
- ▶ If the condition is not met and the resulting outcome is False, the loop is never executed.

#### **Example 8: While loop**

- ▶ Start with an intialization (e.g. vector or matrix) of empties.
- Now followed by a logical comparison between a control variable and a value (using earlier defined conditionals).
- ▶ If the condition is not met and the resulting outcome is False, the loop is never executed.

```
set.seed(1)
usq<-matrix("NA", 100, 1)
n = length(usq)
x=0
while (x \le n) {
x < -sum(x, 1)
i<-sum(i, 1)
usq[i] <- rv[i]*rv[i]
print(usq[i])
usq<-data.frame(usq=as.numeric(unlist(usq)))</pre>
                                   attach(usq)
```

#### Example 8: While loop

Once the condition is past 100, the evaluation is False, and the loop ends:

```
> dim(usq)
[1] 100 :
```

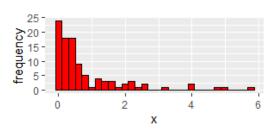


Figure: The square of a standard normal is chi

Note: The while loop was constructed to yield the same output as the for loop. This was less efficient.

#### Example 8: Mix up loops with control flow

Example 4 (revisted): Outlier point in weight and height.

```
weight
height
# Code the for loop with conditionals
for (i in weight) {
if (i > 80) {
print("You're an outlier!")
else {
print("Nothing to see here!")
print(i)
```

#### Example 8: Mix up loops with control flow

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weight
height
# Code the for loop with conditionals
for (i in weight) {
if (i > 80) {
print("You're an outlier!")
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```

**Demonstration:** This is a bit silly but let's see what we get.

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Question: What have we examined and why?

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- 2. Loops and the role of logical statements/condititons
- 3. The role of if statements in executing logicals

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Answer: All of these components are important features of tidying data in R.

#### Functions: Review

An **R function** is created by using the keyword \*function\*. The basic syntax of an R function definition is as follows:

```
function_name <- function(arg_1,..) {
Function_body
}</pre>
```

The different parts of a function are:

- ► Function Name = This is the actual name of the function. It is stored in the R environment as its name.
- Arguments This is an placeholder to pass an input value into your function (i.e. f(x)).
- ► Function Body The function contains the statements that determines what the function does.

# Example 9: A perfect square in R

```
> perfect.squares<-function(x){for (x in 1:10){
+         if(x%x == 0)
+         b<-x^2
+         print (b)
+ }}
> perfect.squares(10)
```

Figure: A function in R to compute perfect squares of all the numbers in the range of 1 to 10

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+         if(x%x == 0)
+         b<-x^2
+         print (b)
+ }}
> perfect.squares(10)
```

Figure: A function in R to compute perfect squares of all the numbers in the range of 1 to 10

**Demonstration**: Let's see what we have printed.

# Example 9: A perfect square in R

Here is how the function is stored as an object in the R environment:

Figure: A function in R stored in the global environment.

# Example 10: Processing Grades in R

Here is a function that evaluates at two inputs:

```
translate<-function(x,y){
if (x=="NA"){x=0}
else if (x!="NA"){x=x}
print(x)
if (y== -1){y="NA"}
else if (y!=-1){y=y}
print(y)
}</pre>
```

- ▶ If x is "NA" then it will assign a value of 0.
- else if y is equal to -1 then it will assign a value of "NA"
- else let y=y.

# Example 10: Processing Grades in R

Here is the output at different pairs of (x, y).

```
#Testing the function with input values
 x<-"NA"
x2 < -1
v2<-0
translate(x, y)
11 "NA"
translate(x2, y2)
```

Figure: A function in R stored in the global environment.

**Question:** Does anyone see a problem with the way in which the function is evaluating the output?

# Example 10: Processing Grades in R

Here is the output at different pairs of (x, y).

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```

Figure: A function in R stored in the global environment.

**Question:** Does anyone see a problem with the way in which the function is evaluating the output?

Ans: It does so term by term, which is inefficient if we have a vector of entries that we want to process (let's see).

# Example 11: if else function in R

In order to understand the brilliance of this function let's go back to Example 10.

```
> x3<-c(0, "NA", 1)
> y3<-c(-1, "NA", 80)
> translate(x3,y3)
[1] "0" "NA" "1"
[1] "NA"
warning messages:
1: In if (x == "NA") { :
    the condition has length > 1 and only the first e
lement will be used
2: In if (x != "NA") { :
    the condition has length > 1 and only the first e
lement will be used
3: In if (y == -1) { :
    the condition has length > 1 and only the first e
lement will be used
3: In if (y == -1) { :
    the condition has length > 1 and only the first e
lement will be used
> |
```

Figure: The translate function in R with vector inputs.

# Example 11: if else function in R

The **'ifelse'** is a built in base R function that returns a value with the same length as the test, rather than the evaluation at the first element.

```
x3<-c(0, "NA", 1)
> z<-x3
> ifelse(z==0, "NA", z)
[1] "NA" "NA" "1"
>
```

#### Aside:

- ▶ In statistics education it is common to use placeholders of "-1" when a student has a valid documentation for absence.
- ▶ If no valid documentation is received, the student receives a value of 0.
- ► The ifelse function evaluates these condition term by term on the entire vector.

#### Subset function in R

library(myPackage)
attach(Grades)
view(Grades)

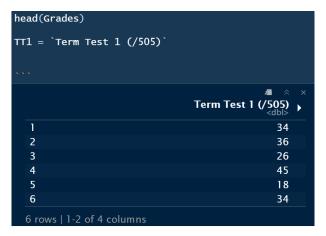


Figure: Processed marks for an undergraduate statistics course.

#### Subset function in R

The **subset** function is the easiest way to select variables and observations (at least within base R).

**Example 10-11 revisted: Grade Processing**: Suppose that I want to select only those students who have no noted absences in Term Test  $1 \, (TT1)$  and Term Test  $2 \, (TT2)$ 

```
dim(Grades)
```

```
newdata <-newdata <- subset(Grades, TT1 !=-1 | TT2!= -1,)
dim(newdata)</pre>
```

```
newgrades <- subset(Grades, TT1 !=-1 & TT2!= -1,)
dim(newgrades)</pre>
```

#### Subset function in R

The **subset** function is the easiest way to select variables and observations (at least within base R).

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dim(newdata)</pre>
```

```
newgrades <- subset(Grades, TT1 !=-1 & TT2!= -1,)
dim(newgrades)</pre>
```

**Demonstration:** What is the difference?

# new.function() function in myPackage

The data set *newgrades* is closer to what would use to process grades for undergraduate students.

This is still not the ideal realization of my grades vectors (I need to account for 0s).

Figure: The new grades function translates "-1" to "NA" and "NA" to "0".

**Demonstration:** Revisit the TT1 and TT2 grades in the Statistics data set.

# Tidyverse

Next information session: The **tidyverse** package is a collection of R packages designed for data science.

- Efficiency of data clean up
- ► Tidy pipeline structures
- Sophisticated plots (in fact I used ggplot here)

Resource: Tidyverse