

Introduction to R

Module 3: Conditionals, loops, functions, and joins

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Intro



Goals for Today

- ① Basics of programming in R—learn how to write and use:
 - Iterations (loops and map functions)
 - Conditional statements
 - Basic functions
- ② Learn how to perform different types of dataset joins.



Iteration



For loops

For tasks that you want to iterate over multiple data frames/variables/elements, you may want to think about creating a **loop**.

- A loop performs a function/functions multiple times, across either a list of objects or a set of index values.

Syntax:

```
for(indexname in range) {  
  do stuff  
}
```



For loop across numeric values

```
for (i in 1:4){  
  print(i^2)  
}
```

```
## [1] 1  
## [1] 4  
## [1] 9  
## [1] 16
```



For loop across named elements

You can also loop over elements instead of values.

- In the last module exercises, you had to convert many columns to numeric. Here's how you could do that with a loop:

```
wdi_data <- import("wdi_data.rds")

vars <- colnames(wdi_data)
indicators <- vars[-(1:4)]

for (i in indicators){
  wdi_data[,i] <- as.numeric(wdi_data[,i])
}
```



The map() function

For iterations over vectors and dataframes, the `map()` function is a great alternative to the for loop.

Map functions take a user-supplied function and iterate it over:

- Elements for a vector
- Objects of a list
- Columns of a data frame

Map functions are much simpler to write than loops and are also generally a good bit faster.

- **Sidenote:** Map is a part of the tidyverse collection of packages. In base R, the `apply()` family of functions does roughly the same thing, but `map()` simplifies and improves this task.



Using the map() function

Syntax:

```
map(data, fxn, option1, option2...)
```

Example:

```
wdi_data[,4:15] <- map(wdi_data[,4:15], as.numeric)
```

Applies the function as.numeric to the columns 4-15 of wdi_data.



Using class-specific map variants

There are multiple map variants that enforce a given data type on results. You should use these whenever you want output of a certain class.

- `map_lgl` for logical vector
- `map_dbl` for numeric vector
- `map_chr` for character vector
- `map_df` for a data frame

```
map_dbl(wdi_data, IQR, na.rm=TRUE)
```

##	GDP_pc	pop	pov
##	10548.89	40860583.00	65.40



Using `map()` with anonymous functions

`map()` works with not only predefined functions, but also “anonymous functions”—unnamed functions defined inside of `map()`.

- In the example below, I z-standardize the values of year, GDP, population, and poverty.

```
ztransform <- map_df(wdi_data, function(x)
  (x - mean(x, na.rm=TRUE)) / sd(x, na.rm=TRUE)
)
```



Did it work?

```
map_dbl(ztransform, function(x)
  round(mean(x, na.rm=TRUE), 10))
```

```
##      year GDP_pc      pop      pov
##           0       0       0       0
```

```
map_dbl(ztransform, function(x)
  round(sd(x, na.rm=TRUE), 10))
```

```
##      year GDP_pc      pop      pov
##           1       1       1       1
```



Conditional Statements



If statement

“If statements” are also a useful part of programming, either in conjunction with iteration or seperately.

- An if statement performs operations only if a specified condition is met.
 - An important thing to know, however, is that if statements evaluate conditions of length one (ie non-vector arguments).
 - We will cover a vector equivalent to the if statement shortly.

Syntax

```
if(condition){  
  do stuff  
}
```



Example of an if statement

- In the for loop example, the loop was indexed over only the columns of indicator codes.
- Equally, the loop could be done over all columns with an if-statement to change only the indicator codes.

```
for (j in colnames(wdi_data)){  
  
  if(j %in% indicators){  
    wdi_data[,j] <- as.numeric(wdi_data[,j])  
  }  
}
```



Multiple conditions

You can encompass several conditions using the **else if** and catch-all **else** control statements.

```
if (condition1) {  
  do stuff  
} else if (condition2) {  
  do other stuff  
} else {  
  do other other stuff  
}
```



Vectorized if statements

- As alluded to earlier, if statements can't test-and-do for vectors, but only single-valued objects.
- Most of the time, you probably want to use conditional statements on vectors. The vector equivalent to the if statement is `ifelse()`

Syntax:

```
ifelse(condition, true_statement, false_statement)
```

The statements returned can be simple values, but they can also be functions or even further conditions. You can easily nest multiple `ifelses` if desired.



An ifelse example

```
numbers <- sample(1:30, 7); numbers
```

```
## [1] 17 20 29 24 22 15 18
```

```
ifelse(numbers %% 2 == 0, "even", "odd")
```

```
## [1] "odd" "even" "odd" "even" "even" "odd" "even"
```

Note: What if we tried a normal if statement instead?

```
if(numbers %% 2 == 0){  
  print("even")} else{  
  print("odd")}
```

```
## [1] "odd"
```



Multiple vectorized if statements

A better alternative to multiple nested `ifelse` statements is the tidyverse `case_when` function.

Syntax:

```
case_when(  
  condition1 ~ statement1,  
  condition2 ~ statement2,  
  condition3 ~ statement3,  
)
```



A case_when example

```
nums_df <- numbers %>% as.tibble() %>%  
  mutate(interval = case_when(  
    (numbers > 0 & numbers <= 10) ~ "1-10",  
    (numbers > 10 & numbers <= 20) ~ "10-20",  
    (numbers > 20 & numbers <= 30) ~ "20-30"))  
nums_df[1:4,]
```

```
## # A tibble: 4 x 2  
##   value interval  
##   <int> <chr>  
## 1     17 10-20  
## 2     20 10-20  
## 3     29 20-30  
## 4     24 20-30
```



Functions



When you should write a function

If you find yourself performing the same specific steps more than a couple of times (perhaps with slight variations), then you should consider writing a function.

A function can serve essentially as a wrapper for a series of steps, where you define generalized inputs/arguments.



Writing a function

Ingredients:

- Function name
- Arguments
- Function body

Syntax:

```
function_name <- function(arg1, arg2, ...){  
  do stuff  
}
```



Function example

Let's turn the calculation of even or odd that was completed earlier into a function:

```
# Make odd function
odd <- function(obj){
  ifelse(obj %% 2 == 0, "even", "odd")
}
```

Notice that *obj* here is a descriptive placeholder name for the data object to be supplied as an argument for the function.

```
odd(numbers)
```

```
## [1] "odd" "even" "odd" "even" "even" "odd" "even"
```



RStudio's "Extract Function"

A useful way of writing simple functions when you've already written the code for a specific instance is to use RStudio's *Extract Function* option, which is available from the code menu.

- *Extract function* will take the code chunk and treat any data objects referenced but not created within the chunk as function arguments.



Joins



Merging data

Shifting gears from programming. . .

Another staple task in applied work is combining data from multiple data sets. The tidyverse set of packages includes several useful types of merges (or “joins”):

- **left_join()** Appends columns from dataset B to dataset A, keeping all observations in dataset A.
- **inner_join()** Appends columns together, keeping only observations that appear in both dataset A and B.
- **semi_join()** Keeps only columns of dataset A for observations that appear in both dataset A and B.
- **anti_join()** Keeps only columns of dataset A for observations that *do not* appear in both dataset A and B.



Joining using keys

The starting point for any merge is to enumerate the column or columns that uniquely identify observations in the dataset.

- For cross-sectional data, this might be a personal identifier or (for aggregate data) something like municipality, state, country, etc.
- For panel data, this will typically be both the personal/group identifier and a timing variable, for example Sweden in 2015 in a cross-country analysis.



Mismatched key names across datasets

Sometimes the names of the key variables are different across datasets.

- You could of course rename the key variables to be consistent.
- But mismatched key names are easily handled by the tidyverse join functions.

Syntax:

```
join_function(x, y, by = c("x_name" = "y_name"))
```



left_join

The **left_join()** is the most frequent type of join, corresponding to a standard **merge** in Stata.

- left_join simply appends additional variables from a second dataset to a main dataset, keeping all the observations (rows) of the first dataset.

Syntax:

```
left_join(x, y, by = "key")
```

If the key is multiple columns, use **c()** to list them.



left_join example

```
# Look at the datasets  
earnings
```

```
##    person_id wage  
## 1         001  150  
## 2         002   90  
## 3         003  270
```

```
educ
```

```
##    person_id schooling  
## 1         001        12  
## 2         003         8  
## 3         004        16
```



left_join example ctd

```
combined_data <- left_join(earnings, educ,  
                           by="person_id")  
  
combined_data
```

##	person_id	wage	schooling
## 1	001	150	12
## 2	002	90	NA
## 3	003	270	8

Notice that schooling is equal to NA for person '002' because that person does not appear in the *educ* dataset.



inner_join

If you want to combine the variables of two data sets, but only keep the observations present in both datasets, use the `inner_join()` function.

```
combined_data <- inner_join(earnings, educ,  
                             by="person_id")  
combined_data
```

```
##   person_id wage schooling  
## 1         001   150        12  
## 2         003   270         8
```



semi_join

To keep using only the variables in the first dataset, but where observations in the first dataset are matched in the second dataset, use `semi_join()`.

- `semi_join` is an example of a *filtering join*. Filtering joins don't add new columns, but instead just filter observations for matches in a second dataset.
- `left_join` and `inner_join` are instead known as *mutating joins*, because new variables are added to the dataset.

```
filtered_data <- semi_join(earnings, educ, by="person_id")  
filtered_data
```

```
##   person_id wage  
## 1         001  150  
## 2         003  270
```



anti_join

Another *filtering join* is `anti_join()`, which filters for observations that are *not matched* in a second dataset.

```
filtered_data <- anti_join(earnings, educ,  
                           by="person_id")  
filtered_data
```

```
##   person_id wage  
## 1         002   90
```

There are still other join types, which you can read about [here](#).

