

Class 5 Data Wrangling with R (Part II)

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October 18, 2023

Section 1

Data Wrangling

Recap: filter(), arrange(), and mutate()

- dataset %>%`filter(criteria): pick observations by their values

Variable 1	Variable 2	Variable 3		Variable 1	Variable 2	Variable 3
A				A		
B				C		
C						

- `dataset %>%`arrange(variable):` reorder the rows
- `dataset %>%`mutate(newvariable = ...):` create new variables with functions of existing variables

Make New Variables



Pipe Operator

- The pipe has a huge advantage over any other method of processing data in R or Python: It makes data wrangling processes easy to read. If we read `%>%` as “then”, the code will be very easy to interpret as a set of instructions in plain English:

```

1 mtcars %>%
2   filter(cyl>=5) %>%
3   mutate(sqrt_wt=sqrt(wt)) %>%
4   filter(sqrt_wt>1.5) %>%
5   arrange(hp)
6 ## can go on and on

```

- 1 take the mtcars data, THEN
- 2 find cars cyl ≥ 5, THEN
- 3 mutate a new variable sqrt_wt, THEN
- 4 find cars sqrt_wt > 1.5 THEN
- 5 reorder all cars based on hp
- 6 chain more operations ...

Without Pipe Operator

- As a comparison, without using pipe operators, the previous data cleaning steps need to be done as follows. Overwriting our output dataframe `new_data` in every line is problematic.
 - First, doing this for a procedure with lots of steps isn't efficient and creates unnecessary repetition in the code.
 - Second, this repetition also makes it harder to identify exactly what is changing on each line in some cases.

```
1 new_data <- filter(mtcars, cyl >=5)
2 new_data <- mutate(new_data, sqrt_wt=sqrt(wt))
3 new_data <- filter(new_data, sqrt_wt>1.5)
4 new_data <- arrange(new_data, hp)
```

Select Variables: select

- `select()` can select variables into a smaller dataset.

```
1 # Select two columns: hp and cyl
2 mtcars%>%
3   select(hp, cyl) %>%
4   head()
```

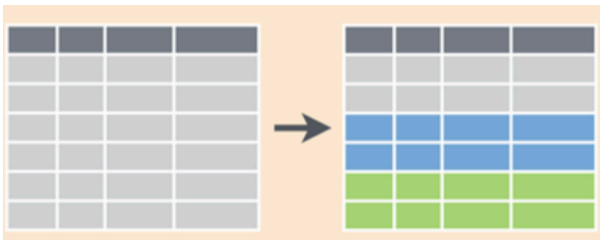
Error in mtcars %>% select(hp, cyl) %>% head(): could not find function

Aggregation by Groups: `group_by`

- `group_by()` allows us to aggregate data by group and compute statistics for each group

```
1 # group by cyl
2 mtcars %>%
3   group_by(cyl)
```

- Although nothing seemingly happens to the dataset, internally, the dataset is already grouped based on the specified variable(s).



Aggregation by Groups: `group_by()` + `summarise()`

- `summarise()` creates a new data frame after aggregating data. The final dataset
 - has one row for each pair of grouping variables (for each `cyl` value)
 - contains one column for each grouping variable (`cyl`)
 - contains one column for each new summarised variable (`avg_mpg`)

```
1 # compute the average mpg for each cyl group
2 mtcars %>%
3   group_by(cyl) %>% # group by cyl
4   summarise(avg_mpg = mean(mpg)) %>% # compute the average mpg
5   ungroup()
```

```
Error in mtcars %>% group_by(cyl) %>% summarise(avg_mpg = mean(mpg)) %>%
```


Aggregation by Groups: `group_by()` + `summarise()`

- We can have multiple group variables for `group_by`

```
1 # compute the average mpg for each cyl,vs group
2 mtcars %>%
3   group_by(cyl,vs) %>% # group by cyl
4   summarise(avg_mpg = mean(mpg)) %>% # compute the average mpg
5   ungroup()
```

Aggregation by Groups: `group_by()` + `mutate()`

- Try the following code by replacing `summarise()` with `mutate()`, what do you get now?

```
1 # compute the average mpg for each cyl,vs group
2 mtcars %>%
3   group_by(cyl,vs) %>% # group by cyl
4   mutate(avg_mpg = mean(mpg)) %>% # compute the average mpg
5   ungroup()
```

- A new column is added to the original dataset, the value of which is from the group-by aggregation.

Consolidate Multiple Data Frames

- When consolidating multiple data frames, we usually have 4 types of joining methods

Combine Data Sets

a		b	
x1	x2	x1	x3
A	1	A	T
B	2	B	F
C	3	D	T

+

=

Mutating Joins

x1	x2	x3
A	1	T
B	2	F
C	3	NA

dplyr::left_join(a, b, by = "x1")

Join matching rows from b to a.

x1	x3	x2
A	T	1
B	F	2
D	T	NA

dplyr::right_join(a, b, by = "x1")

Join matching rows from a to b.

x1	x2	x3
A	1	T
B	2	F

dplyr::inner_join(a, b, by = "x1")

Join data. Retain only rows in both sets.

x1	x2	x3
A	1	T
B	2	F
C	3	NA
D	NA	T

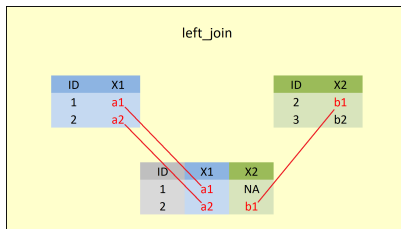
dplyr::full_join(a, b, by = "x1")

Join data. Retain all values, all rows.

left_join

- `left_join` keeps everything from the left data frame and matches as much as it can from the right data frame.
 - All IDs **in the left data frame** will be retained
 - If a match can be found, value from the right data frame will be filled in
 - If a match cannot be found, a missing value will be filled in

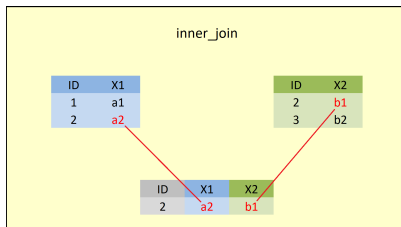
```
1 # Method 1 without pipe operator
2 left_join(df_left, df_right, by = 'ID')
3 # Method 2 with pipe operator
4 df_left %>%
5   left_join(df_right, by = 'ID')
```



inner_join

- `inner_join` only keeps the observations that appear in both data frames
 - Only common IDs in **both data frames** will be retained
 - If a match can be found, values will be filled in from both data frames

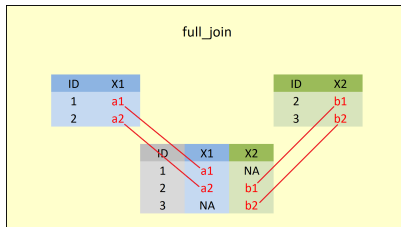
```
1 # Method 1 without pipe operator
2 inner_join(df_left, df_right, by = 'ID')
3 # Method 2 with pipe operator
4 df_left %>%
5   inner_join(df_right, by = 'ID')
6 # Method 3: order of data frames should not matter. Why?
7 df_right %>%
8   inner_join(df_left, by = 'ID')
```



full_join

- `full_join` keeps all observations from both data frames
 - All IDs in **either data frames** will be retained
 - If a match can be found, values will be filled in from both data frames

```
1 # Method 1 without pipe operator
2 full_join(df_left, df_right, by = 'ID')
3 # Method 2 with pipe operator
4 df_left %>%
5   full_join(df_right, by = 'ID')
6 # Method 3: order of data frames should not matter. Why?
7 df_right %>%
8   full_join(df_left, by = 'ID')
```



Section 2

Data Cleaning

Variable Types

- Non-metric
 - Categorical (gender, region, brand, religion)
 - Ordinal (Business Week rankings, NCAA rankings)
- Metric
 - Continuous (age, height, sales, rainfall)
- Different types of variables are handled in different ways in statistics
 - Can talk about an average age, but not an average color
 - Some statistical techniques only work with one type of variable
- We need to make sure the variables are of the correct data types. Or we may need to convert them to the correct types.
 - e.g., from character to date time using `lubridate` package

Missing Values

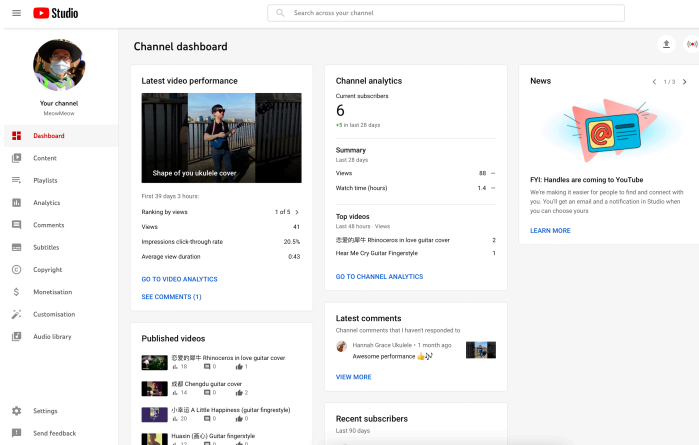
- In R, missing values are represented by the symbol NA (i.e., *not available*).
- Most statistical models cannot handle missing values, so we need to deal with them in R.
 - Few missing values: remove them from analysis.
 - Many missing values: need to replace them with appropriate values: mean/median/**imputation**

Section 3

Descriptive Analytics

Descriptive Analytics

- You can think of descriptive analytics as **creating a dashboard** to display the key information you would like to know for your business.



Summary Statistics

- In descriptive analytics, **summary statistics** are used to summarize a set of observations, in order to communicate the largest amount of information as simply as possible.
- There are two main types of summary statistics used in evaluation: **measures of central tendency** and **measures of dispersion**.
 - Measures of central tendency provide different versions of the average, including the mean, the median, 25 percentile, 75 percentile, the mode, etc.
 - Measures of dispersion provide information about how much variation there is in the data, including the range and the standard deviation.
- It's good to include summary statistics table in your dissertation before any statistical analysis!
 - Commonly reported summary statistics include *mean*, *standard deviation*, *number of observations*, *min*, *25 percentile*, *median*, *75 percentile*, and *max*.
 - Then describe the distribution of the variable, dispersion of the variable, etc.

Summary Statistics with R

- In R, a nice package to report summary statistics is `modelsummary`.
`datasummary_skim()` is a shortcut to conduct basic summary statistics
- For more features, refer to the package tutorial [here](#), especially `datasummary()` function.
 - `datasummary_skim()` is a special case of more general `datasummary()`, which outputs a pre-determined set of summary statistics

```
1  pacman::p_load(modelsummary)
2  mtcars %>%
3    datasummary_skim()
```

Correlation Matrix

- Correlation matrix helps us understand the co-movement of any two variables in the data
- `datasummary_correlation()` reports the pairwise correlation coefficient
- In general, in a statistical model, variables of high correlation should not be included together, which leads to instability

```
1 mtcars %>%  
2   datasummary_correlation()
```

Section 4

Preliminary Customer Analysis

Preliminary Customer Analysis

- Spend 30 min-ish to work on the case study in a group.
- There are 7 questions in total. At the end of the discussion, each group selects a group leader to answer one question.
- To show your codes, join the Zoom link under “Module Overview” on Moodle.
- The group that did not answer correctly needs to do a performance next week!

After-Class Exercise

- What percent of customers are single? Try alternative ways to do the calculation.
- Is the average total spending by responders and non-responders statistically different? Answer this question using a t-test.
- Is income and total spending correlated?
- Are PhDs more likely to respond to marketing offers than Graduation? Use a statistical test to answer the question. Is the result what you expected?
- What would be the other useful descriptive analytics you would like to know for Tesco?