

Session 3: Data Wrangling

R for Stata Users

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The World Bank – DIME | [WB Github](#)

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Introduction

Introduction

Initial Setup

If You Created an RStudio Project in Session 2

If You Did Not Attend Session 2

1. Go to the `dime-r-training-mar2024` folder that you created yesterday, and open the `dime-r-training-mar2024` R project that you created there.

Introduction

Initial Setup

If You Created an RStudio Project in Session 2

If You Did Not Attend Session 2

1. Open RStudio.
2. Type in the following lines, replacing “YOURFILEPATHHERE” (use forward slashes only: "/") with a file path where the file path where you will place this R project.

```
install.packages("usethis")  
library(usethis)  
usethis::use_course(  
  "https://github.com/worldbank/dime-r-training/archive/main.zip",  
  destdir = "YOURFILEPATHHERE"  
)
```

3. In the console, type in the requisite number to delete the .zip file (we don't need it anymore)
4. A new RStudio environment will open. Use this for the session today.

Introduction

Goals of this session

- To organize data in a manner that makes it easier to analyze and communicate.

Things to keep in mind

- We'll take you through the same steps we've taken when we were preparing the datasets used in this course.
- In most cases, your datasets won't be **tidy**.

Tidy data: A dataset is said to be tidy if it satisfies the following conditions:

“**TIDY DATA** is a standard way of mapping the meaning of a dataset to its structure.”
—HADLEY WICKHAM

In tidy data:

- each **variable** forms a **column**
- each **observation** forms a **row**
- each **cell** is a **single measurement**

each column a variable

id	name	color
1	floof	gray
2	max	black
3	cat	orange
4	donut	gray
5	merlin	black
6	panda	calico

each row an observation

Therefore, messy data is any other arrangement of the data.

Introduction

In this session, you'll be introduced to some basic concepts of data cleaning in R. We will cover:

1. Exploring a dataset;
2. Creating new variables;
3. Filtering and subsetting datasets;
4. Merging datasets;
5. Dealing with factor variables;
6. Saving data.

There are many other tasks that we usually perform as part of data cleaning that are beyond the scope of this session.

Before we start, let's make sure we are ready:

1. Start a fresh RStudio session.
2. Open the RStudio project you created yesterday.
3. Create a new R Script called `exercises-session3.R`

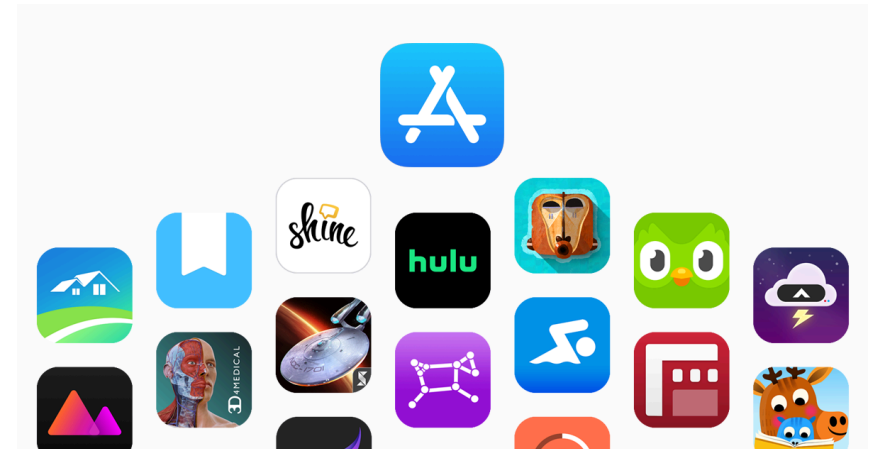
Introduction: Packages

Another important aspect to consider is R packages. Consider the following:

R is a new phone



R packages are apps on your phone



RECAP: Packages

To install a package you can run the following command:

```
# To install  
install.packages("dplyr")
```

- Unlike Stata, R packages need to be loaded in each R session that will use them.
- That means that, for example, a function that comes from the `dplyr` package cannot be used if the package has not been installed and loaded first.

To load a package you can run the following command:

```
# To load  
library(dplyr)
```

RECAP: loading packages

```
# If you haven't installed the packages uncomment the next line  
# install.packages("tidyverse")  
# install.packages("here")  
# install.packages("janitor")  
library(tidyverse) # To wrangle data  
library(here) # A package to work with relative file paths  
library(janitor) # Additional data cleaning tools
```

```
## Warning: package 'janitor' was built under R version 4.3.1
```

Notes: Remember you should always load your packages before your start coding.

RECAP: File paths

The `here` package allows you to interact with your working directory. It will look for the closest R Project and set its location as the working directory. That's why it is important to set your RStudio project correctly.

The goal of this package is to:

- Easily reference your files in project-oriented workflows.

Using `here`:

- Load the library.
- Use `here()` for relative file paths.

```
path <- here("data", "raw", "data-file.csv")
df <- read.csv(path)
```



RECAP: Loading a dataset in R

Before we start wrangling our data, let's read our dataset. In R, we can use the `read.csv` function from Base R, or `read_csv` from the `readr` package if we want to load a CSV file. For this exercise, we are going to use the World Happiness Report (2015-2018)

Exercise 1: Loading data using the `here` package:

Use either of the functions mentioned above and load the three WHR datasets from the `DataWork/DataSets/Raw/UnWHR` folder. Use the following notation for each dataset: `whrYY`, e.g. `WHR2015.csv` becomes the `whr15` dataset.

Solution:

```
whr15 <- read_csv(here("DataWork", "DataSets", "Raw", "Un WHR", "WHR2015.csv"))
whr16 <- read_csv(here("DataWork", "DataSets", "Raw", "Un WHR", "WHR2016.csv"))
whr17 <- read_csv(here("DataWork", "DataSets", "Raw", "Un WHR", "WHR2017.csv"))
```

00:45

The pipe %>% (or |>) operator

- "Piping" in R can be seen as "**chaining**." This means that we are invoking multiple method calls.
- Every time you have invoked a method (a function) this return an object that then is going to be used in the next pipe.

```
rony %>%  
  wake_up(time = "5:30") %>%  
  get_out_of_bed() %>%  
  do_exercise() %>%  
  shower() %>%  
  get_dressed() %>%  
  eat(meal = "breakfast", coffee = TRUE) %>%  
  brush_teeth() %>%  
  work(effort = "minimum")
```

```
work(  
  brush_teeth(  
    eat(  
      get_dressed(  
        shower(  
          do_exercise(  
            get_out_of_bed(  
              wake_up(rony, time = "5:30")  
            ),  
          )  
        )  
      ), meal = "breakfast", coffee = TRUE  
    )  
  ), effort = "minimum"  
)
```

The pipe %>% operator

From R for Data Science by Wickham & Grolemund:

Pipes are a powerful tool for clearly expressing a sequence of multiple operations. The point of the pipe is to help you write code in a way that is easier to read and understand. [...] It focusses on verbs, not nouns. You can read this series of function compositions like it's a set of imperative actions.

(only for 🧐 nerds:)

- The `%>%` pipe is part of the `magrittr` package. `R v4.1.0` adds a *native pipe* via `|>`. you could use it like

```
whr15|> mean(variable, na.rm = T)
```

Janitor package: The `clean_names()` function

The `clean_names()` function helps us when our variables names are pretty bad. For example, if we have a variable that is called `GDP_per_Capita_2015`, the `clean_names()` function will help us fix those messy names.

Tip: Pipe the `clean_names()` function after you load a dataset.

```
whr15 <- whr15 %>%  
  clean_names()  
whr16 <- whr16 %>%  
  clean_names()  
whr17 <- whr17 %>%  
  clean_names()
```

However, if we want to rename our variable manually, we could use:

```
whr15 <- whr15 %>%  
  rename(  
    var_newname = var_oldname  
  )
```

Exploring your data

Exploring a data set

These are some useful functions from base R:

- `View()`: open the data set.
- `class()`: reports object type of type of data stored.
- `dim()`: reports the size of each one of an object's dimension.
- `names()`: returns the variable names of a dataset.
- `str()`: general information on an R object.
- `summary()`: summary information about the variables in a data frame.
- `head()`: shows the first few observations in the dataset.
- `tail()`: shows the last few observations in the dataset.

Some other useful functions from the tidyverse:

- `glimpse()`: get a glimpse of your data.

Load and show a dataset

We can just show our dataset using the name of the object; in this case, `whr15`.

```
whr15
```

```
## # A tibble: 158 × 12
##   country      region      happiness_rank happiness_score standard_error
##   <chr>        <chr>          <dbl>          <dbl>          <dbl>
## 1 Switzerland Western Europe      1            7.59            0.0341
## 2 Iceland      Western Europe      2            7.56            0.0488
## 3 Denmark      Western Europe      3            7.53            0.0333
## 4 Norway       Western Europe      4            7.52            0.0388
## 5 Canada       North America      5            7.43            0.0355
## 6 Finland      Western Europe      6            7.41            0.0314
## 7 Netherlands Western Europe      7            7.38            0.0280
## 8 Sweden       Western Europe      8            7.36            0.0316
## 9 New Zealand  Australia and New ... 9            7.29            0.0337
## 10 Australia   Australia and New ... 10           7.28            0.0408
## # i 148 more rows
## # i 7 more variables: economy_gdp_per_capita <dbl>, family <dbl>,
## #   health_life_expectancy <dbl>, freedom <dbl>,
## #   trust_government_corruption <dbl>, generosity <dbl>,
```

Glimpse your data

Use `glimpse()` to get information about your variables (e.g., type, row, columns,)

```
whr15 %>%
```

```
glimpse()
```

```
## Rows: 158
## Columns: 12
## $ country      <chr> "Switzerland", "Iceland", "Denmark", "Norw...
## $ region       <chr> "Western Europe", "Western Europe", "Weste...
## $ happiness_rank <dbl> 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13,...
## $ happiness_score <dbl> 7.587, 7.561, 7.527, 7.522, 7.427, 7.406, ...
## $ standard_error <dbl> 0.03411, 0.04884, 0.03328, 0.03880, 0.0355...
## $ economy_gdp_per_capita <dbl> 1.39651, 1.30232, 1.32548, 1.45900, 1.3262...
## $ family       <dbl> 1.34951, 1.40223, 1.36058, 1.33095, 1.3226...
## $ health_life_expectancy <dbl> 0.94143, 0.94784, 0.87464, 0.88521, 0.9056...
## $ freedom      <dbl> 0.66557, 0.62877, 0.64938, 0.66973, 0.6329...
## $ trust_government_corruption <dbl> 0.41978, 0.14145, 0.48357, 0.36503, 0.3295...
## $ generosity  <dbl> 0.29678, 0.43630, 0.34139, 0.34699, 0.4581...
## $ dystopia_residual <dbl> 2.51738, 2.70201, 2.49204, 2.46531, 2.4517...
```

Dimensions of your data

Dimensions of your data

Let's see first how many columns and observations the dataset has:

- **Dimensions of your data** (Rows and Columns):

```
dim(whr15)
```

```
## [1] 158 12
```

- **The number of distinct values of a particular variable:**

```
n_distinct(DATASET$variable, na.rm = TRUE)
```

The `$` sign is a subsetting operator. In R, we have three subsetting operators (`[[`, `[`, and `$`). It is often used to access variables in a dataframe.

The `n_distinct` function allows us to count the number of unique values of a variable length of a vector. We included `na.rm = TRUE`, so we don't count missing values.

Dimensions of your data

Exercise 2: Identify distinct values of a variable in a dataset. Using the `n_distinct` function, can you tell how many unique values these variables in the `whr15` dataset have?

1. Country
2. Region

Solution:

```
n_distinct(whr15$country, na.rm = TRUE)
```

```
## [1] 158
```

```
n_distinct(whr15$region, na.rm = TRUE)
```

```
## [1] 10
```

01:00

Dimensions of your data

We can also test whether the number of rows is equal to the number of distinct values in a specific variable as follows:

```
nrow(whr15)
```

```
## [1] 158
```

We can use the two functions (`nrow` and `n_distinct`) together to test if their result is the same.

```
n_distinct(whr15$country, na.rm = TRUE) == nrow(whr15)
```

```
## [1] TRUE
```

```
n_distinct(whr16$country, na.rm = TRUE) == nrow(whr16)
```

```
## [1] TRUE
```

```
n_distinct(whr17$country, na.rm = TRUE) == nrow(whr17)
```

```
## [1] TRUE
```

Wrangling your data

Wrangling vs Cleaning

Cleaning:

- Detecting and addressing inconsistencies in a dataset. Removing erroneous data from your data.

Wrangling:

- Translating raw data into a more useful form. Unifying messy and complex data.

Wrangling your data

For data wrangling you will frequently use one tidyverse package called dplyr.

- `dplyr` is part of the `tidyverse` package family.
- You are **highly encouraged** to read through [Hadley Wickham's chapter](#). It's clear and concise.
- Also check out this great "cheatsheet" [here](#).
- The package is organized around a set of **verbs**, i.e. *actions* to be taken.
- We operate on `data.frames` or `tibbles` (*nicer looking data.frames.*)
- All *verbs* work as follows:

$$\text{verb}(\underbrace{\text{data.frame}}_{\text{1st argument}}, \underbrace{\text{what to do}}_{\text{2nd argument}})$$

- Alternatively you can (should) use the `pipe` operator `%>%`:

$$\underbrace{\text{data.frame}}_{\text{1st argument}} \underbrace{\%>\%}_{\text{"pipe" operator}} \underbrace{\text{verb}(\text{what to do})}_{\text{2nd argument}}$$

dplyr::filter

- The `filter` function is used to subset rows in a dataset.

```
whr15 %>% filter(region == "Western Europe")
```

```
## # A tibble: 21 × 12
##   country      region      happiness_rank happiness_score standard_error
##   <chr>        <chr>          <dbl>          <dbl>          <dbl>
## 1 Switzerland Western Europe      1            7.59           0.0341
## 2 Iceland      Western Europe      2            7.56           0.0488
## 3 Denmark      Western Europe      3            7.53           0.0333
## 4 Norway       Western Europe      4            7.52           0.0388
## 5 Finland      Western Europe      6            7.41           0.0314
## 6 Netherlands Western Europe      7            7.38           0.0280
## 7 Sweden       Western Europe      8            7.36           0.0316
## 8 Austria      Western Europe     13            7.2            0.0375
## 9 Luxembourg   Western Europe     17            6.95           0.0350
## 10 Ireland     Western Europe     18            6.94           0.0368
## # i 11 more rows
## # i 7 more variables: economy_gdp_per_capita <dbl>, family <dbl>,
## #   health_life_expectancy <dbl>, freedom <dbl>,
## #   trust_government_corruption <dbl>, generosity <dbl>,
```

dplyr::filter

Exercise 3: Use `filter()` to extract rows in these regions: (1) Eastern Asia and (2) North America. Hint: use the **or** operator (`|`):

Solution:

```
whr15 %>%  
  filter(region == "Eastern Asia" | region == "North America")
```

A more elegant approach would be to use the `%in%` operator (equivalent to `inlist()` in Stata):

```
whr15 %>%  
  filter(region %in% c("Eastern Asia", "North America"))
```

01:00

dplyr::filter

Code ↻ Start Over ▶ Run Code

```
1  
2  
3
```

dplyr::filter missing cases

If you want to remove (or identify) missing cases for a specific variable, you can use `is.na()`.

- This function returns a value of true and false for each value in a data set.
- If the value is `NA` the `is.na()` function returns `TRUE`, otherwise, it returns `FALSE`.
- In this way, we can check `NA` values that can be used for other functions.
- We can also negate the function using `!is.na()` which indicates that we want to return those observations with no missing values in a variable.

The function syntax in a pipeline is as follows:

```
DATA %>%  
  filter(  
    is.na(VAR)  
  )
```

What are we returning here?

The observations with missing values for the variable `VAR`.

dplyr::filter missing cases

Let's try filtering the `whr15` data. Let's keep those observations that have information per region, i.e., no missing values.

```
whr15 %>%  
  filter(!is.na(region)) %>%  
  head(5)  
  
## # A tibble: 5 × 12  
##   country      region      happiness_rank happiness_score standard_error  
##   <chr>      <chr>          <dbl>          <dbl>          <dbl>  
## 1 Switzerland Western Europe      1            7.59            0.0341  
## 2 Iceland     Western Europe      2            7.56            0.0488  
## 3 Denmark     Western Europe      3            7.53            0.0333  
## 4 Norway      Western Europe      4            7.52            0.0388  
## 5 Canada      North America       5            7.43            0.0355  
## # i 7 more variables: economy_gdp_per_capita <dbl>, family <dbl>,  
## #   health_life_expectancy <dbl>, freedom <dbl>,  
## #   trust_government_corruption <dbl>, generosity <dbl>,  
## #   dystopia_residual <dbl>
```

Notice that we are negating the function, i.e., `!is.na()`

In case we want to keep the observations that contains missing information we will only use `is.na()`.

dplyr::filter missing cases

Code Start Over Run Code

```
1  
2  
3
```


Creating new variables

Creating new variables

We use `dplyr::mutate()` to create new variables. For example:

```
whr15 %>%  
  mutate(  
    hap_hle = happiness_score * health_life_expectancy  
  )
```

This will add a new variable called `hap_hle` which is the interaction of happiness score and health life expectancy.

Creating new variables: Dummy variables

```
whr15 %>%  
  mutate(happiness_score_6 = (happiness_score > 6))
```

What do you think is happening to this variable?

This new variable contains either `TRUE` or `FALSE`. To have it as a numeric variable (1 or 0, respectively), we include the `as.numeric()` function.

```
whr15 %>%  
  mutate(happiness_score_6 = as.numeric((happiness_score > 6)))
```

Finally, instead of using a random number, such as 6, we can do the following:

```
whr15 %>%  
  mutate(  
    happiness_high_mean = as.numeric((happiness_score > mean(happiness_score, na.rm = TRUE)))  
  )
```

Creating variables by groups

In R, we can use `dplyr::group_by()` before we mutate to group an estimation. For example, we are going to pipe the following functions:

1. Group our data by the `region` variable.
2. Create a variable that would be the mean of `happiness_score` by each region.
3. Select the variables `country, region, happiness_score, mean_hap`.

Example	With variables	Output
---------	----------------	--------

```
DATASET %>%  
  group_by(GROUPING VARIABLE) %>%  
  mutate(  
    NAME OF NEW VAR = mean(VARIABLE, na.rm = TRUE)  
  ) %>%  
  select(VAR1, VAR2, VAR3, VAR4)
```

Creating variables by groups

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1. Group our data by the `region` variable.
2. Create a variable that would be the mean of `happiness_score` by each region.
3. Select the variables `country`, `region`, `happiness_score`, `mean_hap`.

Example

With variables

Output

```
whr15 %>%  
  group_by(region) %>%  
  mutate(  
    mean_hap = mean(happiness_score, na.rm = TRUE)  
  ) %>%  
  select(country, region, happiness_score, mean_hap)
```

Creating variables by groups

In R, we can use `dplyr::group_by()` before we mutate to group an estimation. For example, we are going to pipe the following functions:

1. Group our data by the `region` variable.
2. Create a variable that would be the mean of `happiness_score` by each region.
3. Select the variables `country`, `region`, `happiness_score`, `mean_hap`.

Example	With variables	Output
---------	----------------	--------

```
## # A tibble: 7 × 4
## # Groups:   region [2]
##   country      region      happiness_score mean_hap
##   <chr>        <chr>          <dbl>     <dbl>
## 1 Switzerland Western Europe      7.59      6.69
## 2 Iceland      Western Europe      7.56      6.69
## 3 Denmark      Western Europe      7.53      6.69
## 4 Norway       Western Europe      7.52      6.69
## 5 Canada       North America      7.43      7.27
## 6 Finland      Western Europe      7.41      6.69
## 7 Netherlands Western Europe      7.38      6.69
```

Creating multiple variables at the same time

We can create multiple variables in an easy way. Let's imagine that we want to estimate the mean value for the variables:

`happiness_score`, `health_life_expectancy`, and `trust_government_corruption`.

How we can do it?

- We can use the function `across()`.
Syntax: `across(VARS that you want to transform, FUNCTION to execute)`.
- `across()` should be always use inside `summarise()` or `mutate()`.

Across	Output
--------	--------

```
vars <- c("happiness_score", "health_life_expectancy", "trust_government_corruption")
```

```
whr15 %>%  
  group_by(region) %>%  
  summarize(  
    across(  
      all_of(vars), mean  
    )  
  ) %>%  
  head(3)
```

Creating multiple variables at the same time

We can create multiple variables in an easy way. Let's imagine that we want to estimate the mean value for the variables:

`happiness_score`, `health_life_expectancy`, and `trust_government_corruption`.

How we can do it?

- We can use the function `across()`.
Syntax: `across(VARS that you want to transform, FUNCTION to execute)`.
- `across()` should be always use inside `summarise()` or `mutate()`.

Across	Output
--------	--------

```
## # A tibble: 3 × 4
##   region      happiness_score health_life_expectancy trust_government_cor...1
##   <chr>          <dbl>                <dbl>                <dbl>
## 1 Australia and N...      7.28                0.920                0.393
## 2 Central and Eas...      5.33                0.719                0.0867
## 3 Eastern Asia           5.63                0.877                0.128
## # i abbreviated name: 1trust_government_corruption
```


Exercise 4: Create a variable called `year` that equals to the year of each dataframe using the `mutate()`. Remember to assign it to the same dataframe.

Solution:

```
whr15 <- whr15 %>%  
  mutate(year = 2015)  
  
whr16 <- whr16 %>%  
  mutate(year = 2016)  
  
whr17 <- whr17 %>%  
  mutate(year = 2017)
```

Creating variables

Code ↻ Start Over ▶ Run Code

```
1  
2  
3
```

Appending and merging data sets

Appending dataframes

Now that we can identify the observations, we can combine the data set. Here are two functions to append objects by row

```
rbind(df1, df2, df3) # The base R function
```

```
bind_rows(df1, df2, df3) # The dplyr function, making some improvements to base R
```

Exercise 5: Append data sets. Use the function `bind_rows` to append the three WHR datasets:

Solution:

```
bind_rows(whr15, whr16, whr17)
```

Notes

- One of the problems with binding rows like this is that, sometimes, some columns are not *fully* compatible

00:45

Appending dataframes

Code ↻ Start Over ▶ Run Code

```
1  
2  
3
```

Appending and merging data sets

Exercise 6: Fixing our variables and appending the data frames correctly.

Exercise 6a

- Load the R data set `regions.RDS` from `DataWork/DataSets/Raw/Un WHR` using the `read_rds` function.

Solution:

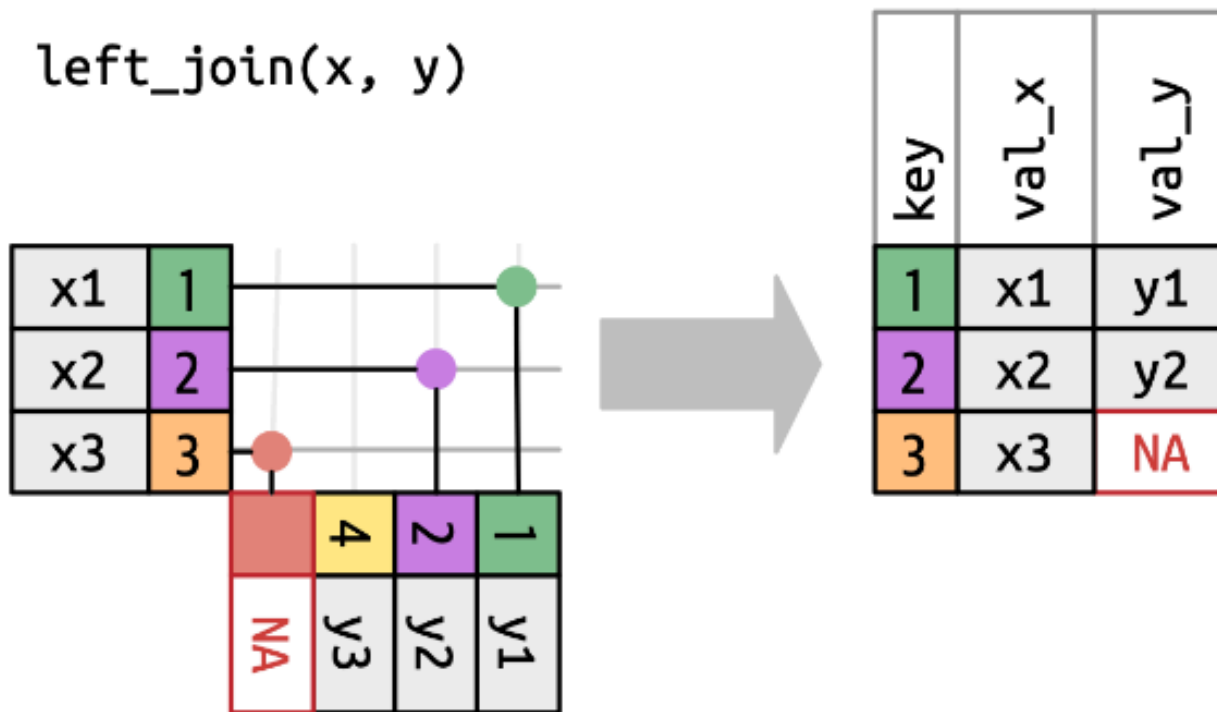
```
regions <- read_rds(here("DataWork", "DataSets", "Raw", "Un WHR", "regions.RDS"))
```

00:45

Appending and merging data sets

We can use the `dplyr::left_join()` function to merge two dataframes. The function syntax is: `left_join(a_df, another_df, by = c("id_coll"))`.

A left join takes all the values from the first table, and looks for matches in the second table. If it finds a match, it adds the data from the second table; if not, it adds missing values. It is the equivalent of `merge, keep(master matched)` in Stata.



Appending and merging data sets

Exercise 6b: Join the dataframes: `regions` and `whr17`.

Solution:

```
whr17 <- whr17 %>%  
  left_join(regions, by = "country") %>%  
  select(country, region, everything())
```

Notes:

Look at the `everything()` function. It takes all the variables from the dataframe and put them after country and region. In this way, select can be use to **order** columns!

01:00

Exercise 6c: Check if there is any other countries in `whr17` without region info:

- Only use pipes `%>%`
- And `filter()`
- Do not assign it to an object.

Solution:

```
whr17 %>%  
  filter(is.na(region))
```

```
## # A tibble: 2 × 14  
##   country          region happiness_rank happiness_score whisker_high whisker_low  
##   <chr>            <chr>          <dbl>          <dbl>          <dbl>        <dbl>  
## 1 Taiwan Provinc... <NA>             33             6.42           6.49         6.35  
## 2 Hong Kong S.A.... <NA>             71             5.47           5.55         5.39  
## # i 8 more variables: economy_gdp_per_capita <dbl>, family <dbl>,  
## #   health_life_expectancy <dbl>, freedom <dbl>, generosity <dbl>,  
## #   trust_government_corruption <dbl>, dystopia_residual <dbl>, year <dbl>
```

So we ended up with two countries with NAs

This is due to the name of the countries. The regions dataset doesn't have "Taiwan Province of China" nor "Hong Kong S.A.R., China" but "Taiwan" and "Hong Kong."

How do you think we should solve this?

- My approach would be to:
 1. fix the names of these countries in the `whr17` dataset (a data cleaning task) and;
 2. merge (`left_join`) it with the regions dataset.

Appendix: [case_when](#) and [mutate](#) for more information.

Appending and merging data sets

01:00

Finally, let's keep those relevant variables first and bind those rows.

Exercise 7: Bind all rows and create a panel called: `whr_panel`.

- Select the variables: `country`, `region`, `year`, `happiness_rank`, `happiness_score`, `economy_gdp_per_capita`, `health_life_expectancy`, `freedom` for each df, i.e., `whr15`, `whr16`, `whr17`.
- Use `rbind()`

Solution:

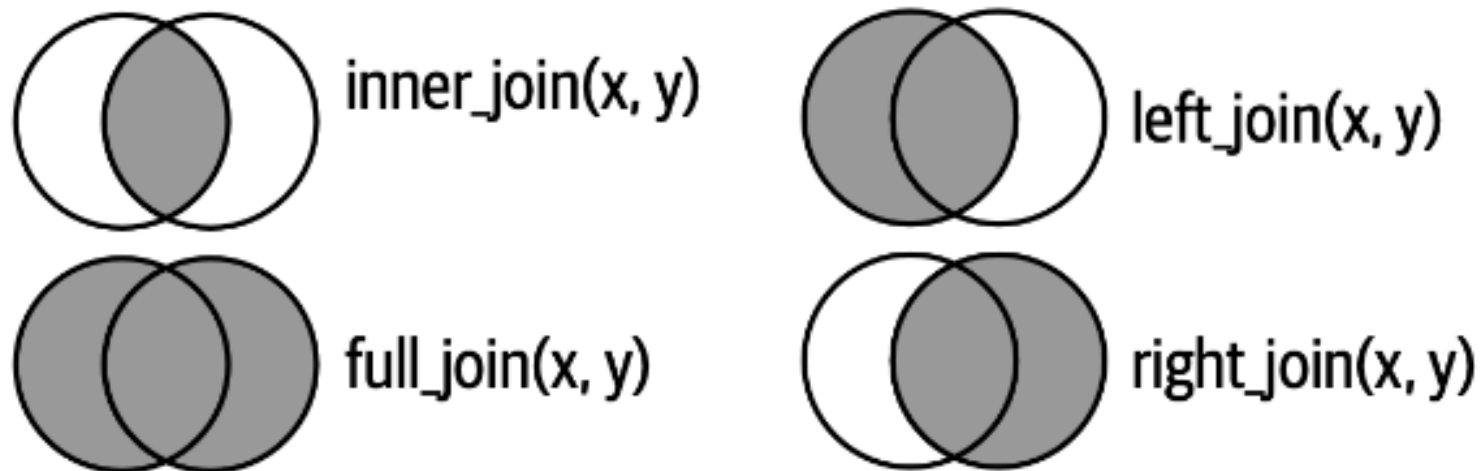
```
vars_to_keep <- c("country", "region", "year", "happiness_rank",  
                 "happiness_score", "economy_gdp_per_capita",  
                 "health_life_expectancy", "freedom")
```

```
whr15 <- select(whr15, all_of(vars_to_keep))  
whr16 <- select(whr16, all_of(vars_to_keep))  
whr17 <- select(whr17, all_of(vars_to_keep))
```

```
whr_panel <- rbind(whr15, whr16, whr17)   # or bind_rows
```

Appending and merging data sets

There are other types of joins in the `dplyr` package. We won't get into detail, but here are some examples.



You can also check [this chapter](#), which is very clear.

Saving a dataset

Saving a dataset

- The dataset you have is the same data set we've been using for earlier sessions, so we can save it now.
- To save a dataset we can use the `write_csv` function from the `readr` package, or `write.csv` from base R.

The function takes the following syntax:

```
write_csv(x, file, append = FALSE, row.names = FALSE, na = ""):
```

- `x`: the object (usually a data frame) you want to export to CSV
- `file`: the file path to where you want to save it, including the file name and the format (".csv")

Exercise 8: Save the dataset as csv format in the "Final" folder with the name `whr_panel_**YOUR INITIALS**.csv`

- Use `write_csv()`
- Use `here()`

Solution:

```
write_csv(
  whr_panel, here("DataWork", "DataSets", "Final", "whr_panel_MA.csv")
)
```

- The problem with CSVs is that they cannot differentiate between `strings` and `factors`
- They also don't save factor orders
- Data attributes (which are beyond the scope of this training, but also useful to document data sets) are also lost.

Saving a dataset

The R equivalent of a `.dta` file is a `.rds` file. It can be saved and loaded using the following commands:

- `write_rds(object, file = "")`: Writes a single R object to a file.
- `read_rds(file)`: Load a single R object from a file.

```
# Save the data set  
  
write_rds(  
  whr_panel,  
  here("DataWork", "DataSets", "Final", "whr_panel_MA.Rds")  
)
```


And that's it for this session. Join us tomorrow!!

Appendix

Missing values in R

Quick Note:

- Missings in R are treated differently than in Stata. They are represented by the NA symbol.
- Impossible values are represented by the symbol NaN which means 'not a number'
- R uses the same symbol for character and numeric data.
- NA is not a string or a numeric value, but an indicator of missingness.
- NAs are contagious. This means that if you compare a number with NAs you will get NAs.
- Therefore, always remember the `na.rm = TRUE` argument if needed.

Other relevant functions: slice, subset, select

Arrange

Slice

Select

Combining functions

Arrange: allows you to order by a specific column.

```
whr15 %>%
```

```
  arrange(region, country) %>%
```

```
  head(5)
```

```
## # A tibble: 5 × 8
```

```
##   country      region  year happiness_rank happiness_score economy_gdp_per_capita
```

```
##   <chr>        <chr> <dbl>          <dbl>          <dbl>          <dbl>
```

```
## 1 Australia  Austr... 2015           10            7.28            1.33
```

```
## 2 New Zealand Austr... 2015            9            7.29            1.25
```

```
## 3 Albania    Centr... 2015           95            4.96            0.879
```

```
## 4 Armenia    Centr... 2015          127            4.35            0.768
```

```
## 5 Azerbaijan Centr... 2015            80            5.21            1.02
```

```
## # i 2 more variables: health_life_expectancy <dbl>, freedom <dbl>
```

Other relevant functions: slice, subset, select

Arrange

Slice

Select

Combining functions

slice: allows you to select, remove, and duplicate rows.

```
whr15 %>%  
  slice(1:5) # to select the first 5 rows  
  
## # A tibble: 5 × 8  
##   country    region  year happiness_rank happiness_score economy_gdp_per_capita  
##   <chr>      <chr> <dbl>         <dbl>          <dbl>          <dbl>  
## 1 Switzerland Weste... 2015             1             7.59             1.40  
## 2 Iceland    Weste... 2015             2             7.56             1.30  
## 3 Denmark    Weste... 2015             3             7.53             1.33  
## 4 Norway     Weste... 2015             4             7.52             1.46  
## 5 Canada     North... 2015             5             7.43             1.33  
## # i 2 more variables: health_life_expectancy <dbl>, freedom <dbl>
```

You can also use **slice_head** and **slice_tail** to select the first or last rows respectively. Or **slice_sample** to randomly draw n rows.

Other relevant functions: slice, subset, select

Arrange

Slice

Select

Combining functions

`Select`: allows you to select specific columns.

```
whr15 %>%  
  select(region, country, happiness_rank)
```

```
## # A tibble: 158 × 3
```

```
##   region          country happiness_rank  
##   <chr>          <chr>          <dbl>  
## 1 Western Europe Switzerland      1  
## 2 Western Europe Iceland          2  
## 3 Western Europe Denmark          3  
## 4 Western Europe Norway           4  
## 5 North America Canada            5  
## 6 Western Europe Finland           6  
## 7 Western Europe Netherlands       7  
## 8 Western Europe Sweden            8  
## 9 Australia and New Zealand New Zealand  9  
## 10 Australia and New Zealand Australia    10
```

Other relevant functions: slice, subset, select

Arrange

Slice

Select

Combining functions

Select: allows you to specific columns.

```
whr15 %>%  
  arrange(region, country) %>%           # Sort by region and country  
  filter(!is.na(region)) %>%           # Filter those non-missing obs for region if any  
  select(country, region, starts_with("happin")) %>% # Select country, year, and vars that stars with happin  
  slice_head()                          # Get the first row
```

```
## # A tibble: 1 × 4  
##   country      region      happiness_rank happiness_score  
##   <chr>      <chr>          <dbl>          <dbl>  
## 1 Australia Australia and New Zealand      10            7.28
```

Using ifelse when creating a variable

We can also create a dummy variable with the `ifelse()` function. The way we use this function is as: `ifelse(test, yes, no)`. We can also use another function called `case_when()`.

```
whr15 %>%
  mutate(
    latin_america_car = ifelse(region == "Latin America and Caribbean", 1, 0)
  ) %>%
  arrange(-latin_america_car) %>%
  head(5)

## # A tibble: 5 × 9
##   country    region  year happiness_rank happiness_score economy_gdp_per_capita
##   <chr>      <chr>  <dbl>         <dbl>          <dbl>          <dbl>
## 1 Costa Rica Latin ... 2015             12             7.23            0.956
## 2 Mexico     Latin ... 2015             14             7.19            1.02
## 3 Brazil     Latin ... 2015             16             6.98            0.981
## 4 Venezuela Latin ... 2015             23             6.81            1.04
## 5 Panama     Latin ... 2015             25             6.79            1.06
## # i 3 more variables: health_life_expectancy <dbl>, freedom <dbl>,
## #   latin_america_car <dbl>
```


Using case_when() to update a variable

Recall the problem we have with regions in the `whr17` data. We can fix it as follows:

```
whr17 <- whr17 %>%  
  mutate(  
    country = case_when(  
      country == "Hong Kong S.A.R., China" ~ "Hong Kong",  
      country == "Taiwan Province of China" ~ "Taiwan",  
      TRUE ~ country  
    )  
  )  
  
whr17 %>%  
  left_join(regions, by = "country") %>%  
  rename(region = region.y) %>%  
  select(-region.x) %>%  
  select(country, region, everything()) %>%  
  filter(is.na(region))
```

Factor variables

Factor variables

- When we imported this data set, we told R explicitly to not read strings as factor.
- We did that because we knew that we'd have to fix the country names.
- The region variable, however, should be a factor.

```
str(whr_panel$region)
```

```
## chr [1:470] "Western Europe" "Western Europe" "Western Europe" ...
```

Factor variables

To create a factor variable, we use the `factor()` function (or `as_factor()` from the `forcats` package).

- `factor(x, levels, labels)` : turns numeric or string vector `x` into a factor vector.
- `levels` : a vector containing the possible values of `x`.
- `labels` : a vector of strings containing the labels you want to apply to your factor variable
- `ordered` : logical flag to determine if the levels should be regarded as ordered (in the order given).

If your categorical variable does not need to be ordered, and your string variable already has the label you want, making the conversion is quite easy.

Factor variables

Extra exercise: Turn a string variable into a factor.

- Use the mutate function to create a variable called region_cat containing a categorical version of the region variable.
- TIP: to do this, you only need the first argument of the factor function.

Solution:

```
whr_panel <- mutate(whr_panel, region_cat = factor(region))
```

And now we can check the class of our variable.

```
class(whr_panel$region_cat)
```

```
## [1] "factor"
```

Reshaping a dataset

Reshaping a dataset

Finally, let's try to reshape our dataset using the tidyverse functions. No more `reshape` from Stata. We can use `pivot_wider` or `pivot_longer`. Let's assign our wide format panel to an object called `whr_panel_wide`.

Long to Wide

Wide to Long

```
whr_panel %>%
  select(country, region, year, happiness_score) %>%
  pivot_wider(
    names_from = year,
    values_from = happiness_score
  ) %>%
  head(3)
```

```
## # A tibble: 3 × 5
##   country      region    `2015` `2016` `2017`
##   <chr>        <chr>    <dbl>  <dbl>  <dbl>
## 1 Switzerland Western Europe  7.59   7.51   7.49
## 2 Iceland      Western Europe  7.56   7.50   7.50
## 3 Denmark      Western Europe  7.53   7.53   7.52
```

Reshaping a dataset

Finally, let's try to reshape our dataset using the tidyverse functions. No more `reshape` from Stata. We can use `pivot_wider` or `pivot_longer`. Let's assign our wide format panel to an object called `whr_panel_wide`.

Long to Wide

Wide to Long

```
whr_panel_wide %>%  
  pivot_longer(  
    cols = `2015`:`2017`,  
    names_to = "year",  
    values_to = "happiness_score"  
  ) %>%  
  head(3)
```

```
## # A tibble: 3 × 4  
##   country      region      year happiness_score  
##   <chr>        <chr>    <chr>          <dbl>  
## 1 Switzerland Western Europe 2015           7.59  
## 2 Switzerland Western Europe 2016           7.51  
## 3 Switzerland Western Europe 2017           7.49
```


Thank you!!