# Week 3: Introduction to the Tidyverse

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### What is the Tidyverse?

- · The tidyverse is a set of packages developed to make the data science process easier
  - They share a common philosophy and grammar
  - Intended to make code more understandable
- · Several different people contributing code can lead to confusion and overlapping ideas
- · Tidyverse packages have different purposes, but read easily into each other

## **Download the Packages**

- · The tidyverse consists of several different packages
- · We can download them all at once, though
  - Just call install.packages("tidyverse")

### Data Manipulation with dplyr

- dplyr has several functions for manipulating data frames
  - Note that they output tibbles rather than data frames
- We can chain functions together using %>%
  - Also known as the pipe operator
  - Note that if for some reason we needed to refer back to the tibble we are manipulating, we would specify it with a .

```
# Format would look like this

tibble_x %>%
  function_x() %>%
  function_y() %>%
  function_z()
```

### Some Common dplyr Functions

- select(): pick certain variables
- filter(): filter your data on a given condition
- arrange(): order your data
- rename(): rename your columns
- mutate(): create a new column

#### select

- · With select, we just specify the variables we want to choose
  - We can instead specify what columns we don't want by putting a before the column

```
initial.starwars %>%

# Select height and birth year of all characters
select(height, birth_year) %>%
head()
```

```
## # A tibble: 6 x 2
    height birth_year
##
                <dbl>
     <int>
##
       172
                 19
## 1
## 2
       167
                112
## 3
       96
                 33
## 4
       202
             41.9
## 5
       150
                 19
## 6
       178
                 52
```

#### filter

- · With filter, we subset the tibble on some condition
  - == means equal to and != means not equal to

```
initial.starwars %>%

# Filter on only males

filter(gender == "male") %>%

head()
```

```
## # A tibble: 6 x 10
    name height mass hair_color skin_color eye_color birth_year gender
    <chr> <int> <dbl> <chr>
                                 <chr>>
                                            <chr>>
                                                          <dbl> <chr>
                   77 blond
                             fair
                                           blue
                                                                male
## 1 Luke~
             172
                                                           19
## 2 Dart~
                            white
                                           yellow
                                                           41.9 male
                   136 none
             202
## 3 Owen~
                   120 brown, gr~ light
                                           blue
                                                           52
                                                                male
             178
                   84 black
                                                           24 male
## 4 Bigg~
             183
                                 light
                                           brown
## 5 Obi-~
             182
                   77 auburn, w∼ fair
                                           blue-gray
                                                               male
                                                           57
                    84 blond
                                 fair
                                            blue
                                                           41.9 male
## 6 Anak~
             188
## # ... with 2 more variables: homeworld <chr>, species <chr>
```

#### arrange

- · With arrange, we order the tibble by some column
  - By default, order is ascending; to get descending order, we use desc()

```
initial.starwars %>%

# Arrange by descending height
arrange(desc(height)) %>%
head()
```

```
## # A tibble: 6 x 10
    name height mass hair_color skin_color eye_color birth_year gender
    <chr> <int> <dbl> <chr>
                                 <chr>>
                                            <chr>>
                                                          <dbl> <chr>
                                            yellow
## 1 Yara~
             264
                    NA none
                                 white
                                                             NA male
## 2 Tarf~
                   136 brown
                                            blue
                                                             NA male
             234
                                 brown
## 3 Lama~
                   88 none
                                            black
                                                         NA male
             229
                                 grey
                                 unknown
                                            blue
                                                            200 male
## 4 Chew~
             228
                   112 brown
## 5 Roos~
             224
                 82 none
                                                    NA male
                                 grey
                                            orange
## 6 Grie~
             216
                   159 none
                                 brown, wh~ green, y~
                                                             NA male
## # ... with 2 more variables: homeworld <chr>, species <chr>
```

#### rename

- rename allows us to change the name of a column
  - First specify the new name, than specify what column is being renamed

```
initial.starwars %>%

# Rename name to full_name
rename(full_name = name) %>%
head()
```

```
## # A tibble: 6 x 10
    full_name height mass hair_color skin_color eye_color birth_year gender
               <int> <dbl> <chr>
    <chr>>
                                    <chr>>
                                              <chr>>
                                                             <dbl> <chr>
                       77 blond
                                    fair
                                              blue
## 1 Luke Sky~
                172
                                                              19
                                                                  male
## 2 C-3P0
                167 75 <NA>
                                    gold
                                              yellow
                                                             112
                                                                  <NA>
## 3 R2-D2
           96
                     32 <NA>
                                    white, bl~ red
                                                                  <NA>
                                    white
                                              yellow
                                                             41.9 male
## 4 Darth Va~ 202
                      136 none
## 5 Leia Org~
                150
                     49 brown
                                    light
                                              brown
                                                                  female
                                                              19
                                              blue
## 6 Owen Lars
                178
                      120 brown, gr~ light
                                                              52
                                                                  male
## # ... with 2 more variables: homeworld <chr>, species <chr>
```

#### mutate

- mutate allows us to create new columns
  - Remember to save your output if you want to keep it!

```
new.starwars <- initial.starwars %>%

# Creating a new column that is the square root of height
mutate(sqrt_height = sqrt(height))

new.starwars %>%
select(height, sqrt_height) %>%
head(3)
```

#### summarize

· With summarize, we can perform certain summary statistics, like finding the average, or getting a count

```
initial.starwars %>%

# Here we are finding the average height of all characters
summarize(avg.height = mean(height, na.rm=T))
```

### group\_by

- group\_by allows us to perform different operations by groups
  - If we want to end the grouping, use ungroup()

```
initial.starwars %>%

# Grouping by gender
group_by(gender) %>%

# Finding average height
summarize(avg.height = mean(height, na.rm=T)) %>%

# Remember to ungroup if you want to perform non-grouped functions
# after grouping
ungroup()
```

```
## # A tibble: 5 x 2
    gender
                  avg.height
##
    <chr>
                       <dbl>
## 1 female
                        165.
## 2 hermaphrodite
                   175
                        179.
## 3 male
                        200
## 4 none
## 5 <NA>
                        120
```

## **Chaining Functions**

· With the %>% operator, we can easily chain several functions together

```
initial.starwars %>%

# Filter characters taller than 200 cm
filter(height > 200) %>%

# Select the mass column
select(mass) %>%

# Order mass in descending order
arrange(desc(mass))
```

```
## # A tibble: 10 x 1
## mass
## <dbl>
## 1 159
## 2 136
## 3 136
## 4 112
## 5 88
## 6 82
## 7 80
## 8 NA
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```

### **Joining Data**

- We can use several functions that work similarly to **SQL** joins
- inner\_join: return all rows from x where there are matching values in y
- · left\_join: return all rows from x
- right\_join: return all rows from y
- full\_join: return all rows from x and y

# **Reshaping Data**

- · Data generally comes in wide and long format
- · Wide format makes every variable a column
- · Long format puts multiple variables in one column (key) and their values in a second column
- · Consider the goal of your analysis when choosing which format

### Reshaping with tidyr

- tidyr allows us to easily go from long to wide or vice-versa
- The main functions we'll use are gather() and spread()
  - These change data sets into long and wide formats, respectively

### **Going from Wide to Long**

- · We'll use gather() to go from wide to long
- We need to first specify what columns we are gathering
  - The names of the columns will become our key, their values become our value
  - We need to also specify the new key and value column names

### **Going from Long to Wide**

- We'll use spread() to go from long to wide
- · We need to first specify where the data is coming from, followed by the key and value columns

```
spread(data = long.tb, key = Year, value = Cases)
```