

Data Wrangling

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STA 209, 2/6/23

Outline

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- Efficiently summarize data with the `summarize` function
- Discuss data wrangling and survey the `dplyr` verbs
- Practice decomposing data using the “grammar of wrangling”

Section 1

Summarizing with dplyr

The dplyr package



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mean(biketown$Distance_Miles)
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- But it would be nice to have an easy way to store multiple summary statistics in a data frame

The `summarize` function

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The summarize function

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```
library(dplyr)
summarize(
  biketown,
  Mean_Distance = mean(Distance_Miles),
  SD_Distance = sd(Distance_Miles),
  Median_StartHour = median(StartHour),
  IQR_StartHour = IQR(StartHour)
)
```

```
## # A tibble: 1 x 4
##   Mean_Distance SD_Distance Median_StartHour IQR_StartHour
##           <dbl>         <dbl>           <int>         <dbl>
## 1           2.04           1.95             15             7
```

- Note that code is separated by line breaks for improved readability
- New column names can be arbitrary (but it's nice if they are informative)

The summarize function

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```
library(dplyr)
summarize(
  biketown,
  These = mean(Distance_Miles),
  Can = sd(Distance_Miles),
  Be = median(StartHour),
  Whatever = IQR(StartHour)
)
```

```
## # A tibble: 1 x 4
##   These    Can    Be Whatever
##   <dbl> <dbl> <int>    <dbl>
## 1  2.04  1.95    15      7
```

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distance_summary <- summarise(biketown,  
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- It's helpful to save the summarize dataframe for later access:

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distance_summary <- summarise(biketown,  
                               mean_dist = mean(Distance_Miles),  
                               sd_dist = sd(Distance_Miles))
```

```
distance_summary$mean_dist
```

```
## [1] 2.044768
```

```
distance_summary$sd_dist
```

```
## [1] 1.950804
```

Section 2

Data Wrangling

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- In addition to tidying a data set, data wrangling also allows us to explore components of the data.
- Data analysts and survey statisticians spend about 50 – 80% of work-time on data wrangling.
- As such, it is important to have *consistent* and *efficient* tools for the job.

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 - ④ mutate
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 - ⑥ select
- Each verb takes a data frame and returns a data frame
- Verbs can be chained together using a special operator `%>%` to perform complicated manipulations.
- These verbs form a “grammar” of Data Manipulation.
 - So even if you aren't using R, they represent the basic components you would think about when manipulating data.

A long time ago, in a galaxy far, far away...

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Star Wars: The Rise of Skywalker

We'll investigate the starwars data set from the dplyr package

```
head(starwars)
```

```
## # A tibble: 6 x 14
##   name          height  mass hair_~1 skin_~2 eye_c~3 birth~4 sex  gender homew~5
##   <chr>          <int> <dbl> <chr>   <chr>   <chr>   <dbl> <chr> <chr>  <chr>
## 1 Luke Skywal~    172    77 blond   fair    blue     19  male  mascu~ Tatooi~
## 2 C-3P0          167    75 <NA>    gold    yellow   112  none  mascu~ Tatooi~
## 3 R2-D2           96    32 <NA>    white,~ red      33  none  mascu~ Naboo
## 4 Darth Vader    202   136 none    white    yellow   41.9 male  mascu~ Tatooi~
## 5 Leia Organa    150    49 brown   light    brown     19  fema~ femin~ Aldera~
## 6 Owen Lars      178   120 brown,~ light    blue     52  male  mascu~ Tatooi~
## # ... with 4 more variables: species <chr>, films <list>, vehicles <list>,
## #   starships <list>, and abbreviated variable names 1: hair_color,
## #   2: skin_color, 3: eye_color, 4: birth_year, 5: homeworld
## # i Use `colnames()` to see all variable names
```


`filter()`

Subset Observations (Rows)



`filter()`

Subset Observations (Rows)



```
filter(starwars, height < 100)
```

```
## # A tibble: 7 x 14
##   name      height  mass hair_~1 skin_~2 eye_c~3 birth~4 sex  gender homew~5
##   <chr>      <int> <dbl> <chr>   <chr>   <chr>   <dbl> <chr> <chr> <chr>
## 1 R2-D2         96    32 <NA>   white,~ red      33 none mascul~ Naboo
## 2 R5-D4         97    32 <NA>   white,~ red      NA none mascul~ Tatooi-
## 3 Yoda          66    17 white  green   brown    896 male mascul~ <NA>
## 4 Wicket Syst~  88    20 brown  brown   brown      8 male mascul~ Endor
## 5 Dud Bolt      94    45 none   blue, ~ yellow  NA male mascul~ Vulpter
## 6 Ratts Tyere~  79    15 none   grey, ~ unknown NA male mascul~ Aleen ~
## 7 R4-P17        96    NA none   silver~ red, b~  NA none femin~ <NA>
## # ... with 4 more variables: species <chr>, films <list>, vehicles <list>,
## #   starships <list>, and abbreviated variable names 1: hair_color,
## #   2: skin_color, 3: eye_color, 4: birth_year, 5: homeworld
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`select()`

Subset Variables (Columns)



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```
## # A tibble: 87 x 4
##   name          height mass homeworld
##   <chr>         <int> <dbl> <chr>
## 1 Luke Skywalker    172    77 Tatooine
## 2 C-3P0             167    75 Tatooine
## 3 R2-D2              96    32 Naboo
## 4 Darth Vader       202   136 Tatooine
## 5 Leia Organa       150    49 Alderaan
## 6 Owen Lars         178   120 Tatooine
## 7 Beru Whitesun lars 165    75 Tatooine
## 8 R5-D4              97    32 Tatooine
## 9 Biggs Darklighter 183    84 Tatooine
## 10 Obi-Wan Kenobi    182    77 Stewjon
## # ... with 77 more rows
```

`summarize()`

Summarise Data



`summarize()`

Summarise Data



```
summarize(starwars,  
  Avg_Height = mean(height, na.rm = T),  
  Median_Height = median(height, na.rm = T))
```

```
## # A tibble: 1 x 2  
##   Avg_Height Median_Height  
##   <dbl>         <int>  
## 1      174.           180
```

group_by()

Link data according to levels of a variable. Usually followed by `summarize()`



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```
grouped_sw <- group_by(starwars, gender)
summarize(grouped_sw, Avg_Height = mean(height, na.rm = T))
```

```
## # A tibble: 3 x 2
##   gender    Avg_Height
##   <chr>      <dbl>
## 1 feminine    165.
## 2 masculine    177.
## 3 <NA>        181.
```


`mutate()`

Make New Variables



mutate()

Make New Variables



```
mutated_sw <- mutate(starwars, height_ft = height/30.48)
select(mutated_sw, name, height_ft, everything())
```

```
## # A tibble: 87 x 15
##   name      heigh~1 height  mass hair_~2 skin_~3 eye_c~4 birth~5 sex  gender
##   <chr>      <dbl>  <int> <dbl> <chr>  <chr>  <chr>  <dbl> <chr> <chr>
## 1 Luke Skywa~  5.64    172    77 blond  fair   blue    19  male  mascu~
## 2 C-3P0      5.48    167    75 <NA>  gold  yellow  112 none  mascu~
## 3 R2-D2      3.15     96    32 <NA>  white,~ red    33  none  mascu~
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## 5 Leia Organa 4.92    150    49 brown  light  brown   19  fema~ femin~
## 6 Owen Lars   5.84    178   120 brown,~ light  blue    52  male  mascu~
## 7 Beru White~ 5.41    165    75 brown  light  blue    47  fema~ femin~
## 8 R5-D4       3.18     97    32 <NA>  white,~ red    NA  none  mascu~
## 9 Biggs Dark~ 6.00    183    84 black  light  brown   24  male  mascu~
## 10 Obi-Wan Ke~ 5.97    182    77 auburn~ fair   blue-g~ 57  male  mascu~
## # ... with 77 more rows, 5 more variables: homeworld <chr>, species <chr>,
```

arrange()

Sort the rows



arrange()

Sort the rows



```
arrange(starwars, mass)
```

```
## # A tibble: 87 x 14
##   name      height  mass hair_~1 skin_~2 eye_c~3 birth~4 sex  gender homew~5
##   <chr>      <int> <dbl> <chr>   <chr>   <chr>   <dbl> <chr> <chr>  <chr>
## 1 Ratts Tyer~    79    15 none   grey, ~ unknown NA male mascul~ Aleen ~
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```
arrange(filter(select(my_data, var_1) %in% range))
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- This method has two primary problems:

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```

- This method has two primary problems:
 - ① Code quickly become overwhelming to read and review (especially as number of functions and arguments increases)
 - ② The operations (as read from left to right) appear in the *opposite* order to how they are performed

Pipe Composition

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```

- Reading %>% as “then”, this sequence translates to

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- ② Use this output as input of `select()` *then*
- ③ Use this output as input of `filter()` *then*
- ④ Use this output as input of `arrange()`

- Advantages:

- The pipe sequence is much more readable.
- Much easier to add more functions to the mix at a later time (since they can be tacked on at the end of the sequence)