

Part 2: Basics of Data Management in R

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### The tidyverse

- Suite of packages collectively known as the tidyverse
- Different from base R in many ways
- The tidyverse with pipes<sup>1</sup> is more intuitive to me

<sup>&</sup>lt;sup>1</sup> We'll talk about pipes very soon!

#### What is Tidy data?

#### Resources:

- Paper: Tidy Data (Hadley Wickham, 2014 JSS)
- Vignette: Tidy data (from the tidyr package)

#### Essentially:

- 1. Variables are columns
- 2. Observations are rows
- 3. Variables and observations make a table

#### Intro to Tidy data

Let's load the tidyverse package and check the output:

```
library(tidyverse)
```

Comes with lots of other packages like ggplot2, tibble, dplyr, etc.

#### Pipes: %>%

- The pipe operator is denoted %>% and is automatically loaded with the tidyverse.
- Pipes are awesome!

```
## These next two lines of code do exactly the same thing.
mpg %>% filter(manufacturer="audi") %>% group_by(model) %>% summarise(hwy_mean = mean(hwy summarise(group_by(filter(mpg, manufacturer="audi"), model), hwy_mean = mean(hwy))
```

The first line reads from left to right and from data to operation. The Base R version (line 2) works in the opposite order.

#### Pipes: %>%

## 2 a4 quattro

## 3 a6 quattro

Helps to break the pipes over several lines

25.8

24

#### The dplyr package:

- 1. filter(): Find or exclude certian rows
- 2. arrange(): Sort your observations
- 3. select(): Select specific variables
- 4. mutate(): Create new variables
- 5. summarise(): Collapse multiple rows into a single summary value

### 1) dplyr::filter()

Multiple filters separated by commas:

```
starwars %>%
  filter(
    species = "Human",
    height ≥ 190
    ) %>% head(5)
## # A tibble: 4 × 14
         height mass hair color skin color eye color birth year sex
                                                                 gender
##
    name
    <chr>
         <int> <dbl> <chr>
                                <chr> <chr>
                                                       <dbl> <chr> <chr>
###
## 1 Darth Va... 202 136 none white yellow 41.9 male mascu...
## 2 Qui-Gon ... 193 89 brown fair
                                          blue
                                                        92
                                                           male mascu…
         193 80 white
                             fair
  3 Dooku
                                          brown
                                                       102
                                                            male mascu…
## 4 Bail Pre... 191
                    NA black
                                tan
                                          brown
                                                   67
                                                            male
                                                                 mascu...
## # ... with 5 more variables: homeworld <chr>, species <chr>, films <list>,
    vehicles <list>, starships <list>
```

### 1) dplyr::filter()

Common to use filter() to check or remove missing values

```
starwars %>%
  filter(is.na(height)) %>% head(5)
## # A tibble: 5 × 14
             height mass hair color skin color eye color birth year sex
                                                                        gender
##
    name
          <int> <dbl> <chr>
                                              <chr>
    <chr>
                                    <chr>
                                                             <dbl> <chr> <chr>
##
## 1 Arvel C...
                                   fair
                                              brown
                 NΑ
                      NA brown
                                                               NA male mascul…
## 2 Finn
                NA NA black
                                    dark
                                              dark
                                                               NA male mascul...
## 3 Rey
                 NA NA brown
                               light
                                              hazel
                                                               NA fema... femini...
                                   light
                                                               NA male mascul...
## 4 Poe Dam...
              NA NA brown
                                              brown
                                              black
## 5 BB8
                 NΑ
                      NA none
                                                               NA none
                                                                        mascul...
                                    none
## # ... with 5 more variables: homeworld <chr>, species <chr>, films <list>,
     vehicles <list>, starships <list>
## #
```

## 1) dplyr::filter()

To remove missing observations, use filter(!is.na(height)). Also try the drop\_na() function from tidyr.

## 2) dplyr::arrange()

Arrange in ascending order:

```
starwars %>%
  arrange(birth year) %>% head(5)
## # A tibble: 5 × 14
              height mass hair color skin color eye color birth year sex
                                                                          gender
##
    name
          <int> <dbl> <chr>
                                                <chr>
                                                               <dbl> <chr> <chr>
    <chr>
                                     <chr>
##
## 1 Wicket S...
                                                brown
               88
                        20 brown
                                     brown
                                                                  8 male
                                                                          mascu...
                                     metal
## 2 IG-88
          200 140 none
                                                red
                                                                 15 none
                                                                          mascu...
  3 Luke Sky... 172 77 blond
                                     fair
                                                blue
                                                                 19 male mascu...
## 4 Leia Org... 150 49 brown
                                     light
                                                brown
                                                                 19 fema... femin...
  5 Wedge An... 170
                       77 brown
                                     fair
                                                hazel
                                                                 21 male
                                                                          mascu...
## # ... with 5 more variables: homeworld <chr>, species <chr>, films <list>,
## #
     vehicles <list>, starships <list>
```

## 2) dplyr::arrange()

Arrange descending order using arrange(desc()):

```
starwars %>%
  arrange(desc(birth year)) %>% head(5)
## # A tibble: 5 × 14
            height mass hair color skin color eye color birth year sex
                                                                           gender
##
    name
          <int> <dbl> <chr>
                                                <chr>
    <chr>
                                    <chr>
                                                               <dbl> <chr>
                                                                           <chr>
##
                66
## 1 Yoda
                      17 white
                                                brown
                                                                 896 male
                                    green
                                                                           mascu...
                                                                 600 herma... mascu...
## 2 Jabba ... 175 1358 <NA>
                                    green-tan,... orange
                     112 brown
                                   unknown
                                                blue
  3 Chewba...
             228
                                                                 200 male
                                                                           mascu...
## 4 C-3PO
          167 75 <NA>
                                   gold
                                                vellow
                                                                112 none
                                                                           mascu...
## 5 Dooku
          193 80 white
                                    fair
                                                brown
                                                                 102 male
                                                                           mascu...
## # ... with 5 more variables: homeworld <chr>, species <chr>, films <list>,
## #
     vehicles <list>, starships <list>
```

- Use commas to select multiple columns
- Use "first:last" for consecutive columns
- Deselect a column with "-"

```
starwars %>%
  select(name:skin_color, species, -height) %>% head(5)
## # A tibble: 5 × 5
##
                     mass hair color skin color
                                                  species
     name
                    <dbl> <chr>
                                     <chr>
                                                 <chr>
     <chr>
###
## 1 Luke Skywalker
                       77 blond
                                     fair
                                                 Human
                                                 Droid
## 2 C-3P0
                       75 <NA>
                                     gold
  3 R2-D2
                       32 <NA>
                                     white, blue Droid
                                     white
## 4 Darth Vader
                      136 none
                                                 Human
                                     light
                                                 Human
## 5 Leia Organa
                       49 brown
```

#### Rename within select():

```
starwars %>%
  select(alias=name, crib=homeworld, sex=gender) %>% head(5)
## # A tibble: 5 × 3
    alias
          crib
###
                          sex
   <chr>
          <chr> <chr>
###
## 1 Luke Skywalker Tatooine masculine
## 2 C-3P0
                 Tatooine masculine
                 Naboo masculine
## 3 R2-D2
## 4 Darth Vader Tatooine masculine
## 5 Leia Organa Alderaan feminine
```

Use select(contains(PATTERN)) to find rows that contain some strings of interest

```
starwars %>%
  select(name, contains("color")) %>% head(5)
## # A tibble: 5 × 4
                  hair color skin color eye color
##
    name
                  <chr>
                            <chr>
                                       <chr>
###
    <chr>
## 1 Luke Skywalker blond fair blue
## 2 C-3P0
                  <NA>
                            gold yellow
                            white, blue red
## 3 R2-D2
                  <NA>
## 4 Darth Vader
                            white
                                  yellow
                  none
## 5 Leia Organa
                  brown
                            light
                                       brown
```

Also look into the stringr package.

Create new variables with mutate()

```
starwars %>%
  select(name, birth year) %>%
  mutate(dog years = birth year * 7) %>%
  mutate(comment = paste0(name, " is ", dog years, " in dog years.")) %>% head(5)
## # A tibble: 5 × 4
                  birth year dog years comment
###
    name
  <chr>
                       <dbl> <dbl> <chr>
###
                    19
## 1 Luke Skywalker
                                 133 Luke Skywalker is 133 in dog years.
## 2 C-3P0
                       112
                                 784 C-3PO is 784 in dog years.
                        33
                                 231 R2-D2 is 231 in dog years.
## 3 R2-D2
                   41.9
## 4 Darth Vader
                                 293. Darth Vader is 293.3 in dog years.
## 5 Leia Organa
                        19
                                     Leia Organa is 133 in dog years.
```

mutate() processes in order, so you can put dependent mutates one after another:

```
## # A tibble: 5 × 4
                birth year dog years comment
###
    name
  <chr>
                      <dbl> <dbl> <chr>
###
                   19
## 1 Luke Skywalker
                                133 Luke Skywalker is 133 in dog years.
## 2 C-3P0
                      112
                                784 C-3PO is 784 in dog years.
                       33
## 3 R2-D2
                                231 R2-D2 is 231 in dog years.
                  41.9
## 4 Darth Vader
                                293. Darth Vader is 293.3 in dog years.
## 5 Leia Organa
                       19
                                   Leia Organa is 133 in dog years.
```

Other handy ways to use mutate():

```
starwars %>%
  select(name, height) %>%
  filter(name %in% c("Luke Skywalker", "Anakin Skywalker")) %>%
  mutate(tall1 = height > 180) %>%
  mutate(tall2 = ifelse(height > 180, "Tall", "Short")) ## Same effect, but can choose lab
## # A tibble: 2 × 4
                    height tall1 tall2
###
    name
    <chr>
          <int> <lgl> <chr>
###
## 1 Luke Skywalker 172 FALSE Short
## 2 Anakin Skywalker 188 TRUE Tall
```

Note the "scoped" variants of mutate() that work on a subset of variables:

- mutate\_all() affects every variable
- mutate\_at() affects named or selected variables
- mutate\_if() affects variables that meet some criteria (e.g. are numeric)

These have since been replaced with <code>across()</code>. So you would use <code>mutate(across())</code> instead.

## 5) dplyr::summarise() with group\_by()

```
starwars %>%
  group by(species, gender) %>%
  summarise(mean_height = mean(height, na.rm = T)) %>% head(5)
## # A tibble: 5 × 3
## # Groups: species [5]
   species gender mean height
##
   <chr> <chr> <dbl>
###
## 1 Aleena masculine
                           79
## 2 Besalisk masculine
                             198
## 3 Cerean
            masculine
                             198
## 4 Chagrian masculine
                       196
## 5 Clawdite feminine
                             168
```

Note: na.rm = T is usually a good idea, otherwise your summary will be NA too.

## 5) dplyr::summarise()

```
starwars %>% group_by(species) %>% summarise(across(where(is.numeric), mean, na.rm=TRUE))
## # A tibble: 5 × 4
###
    species height mass birth year
    <chr> <dbl> <dbl>
                               <dbl>
##
## 1 Aleena
                 79
                       15
                                 NaN
## 2 Besalisk
                198
                      102
                                 NaN
                                  92
## 3 Cerean
                198
                     82
  4 Chagrian
                196
                      NaN
                                 NaN
## 5 Clawdite
                168
                       55
                                 NaN
```

## Joining operations

Central feature of the dplyr package invovles merging data from multiple tables with join operations.

- inner\_join(df1, df2)
- left\_join(df1, df2)
- right\_join(df1, df2)
- full\_join(df1, df2)
- semi\_join(df1, df2)
- anti\_join(df1, df2)

## Joining operations

- For some simple examples, we'll need some data sets that come bundled with the nycflights13 package.
- Load it now and then inspect these data frames in your own console.

```
library(nycflights13)
flights
planes
```

Let's perform a left join on the flights and planes datasets.

```
left join(flights, planes) %>%
  select(year, month, day, dep time, arr time, carrier, flight, tailnum, type, model) %>%
## # A tibble: 5 × 10
                  day dep time arr time carrier flight tailnum type model
###
     vear month
     <int> <int> <int>
                         <int>
                                  <int> <chr>
                                              <int> <chr>
                                                              <chr> <chr>
##
## 1
     2013
                           517
                                    830 UA
                                                  1545 N14228 <NA>
                                                                    <NA>
## 2
     2013
                           533
                                    850 UA
                                                  1714 N24211 <NA>
                                                                    <NA>
                           542
                                923 AA
                                                  1141 N619AA <NA>
## 3
     2013
                                                                    <NA>
                           544
                                   1004 B6
                                                   725 N804JB
## 4
     2013
                                                              <NA>
                                                                    <NA>
## 5
     2013
                           554
                                    812 DL
                                                   461 N668DN
                                                              <NA>
                                                                    <NA>
```

dplyr guessed about which columns to join on (i.e. columns that share the same name). It also told us its choices:

```
## Joining, by = c("year", "tailnum")
```

Problem: the variable "year" does not have a consistent meaning across our joining datasets!

year of flight versus year of construction

Luckily, there's an easy way to avoid this problem.

- See if you can figure it out before turning to the next slide.
- Try ?dplyr::join.

Let's be more explicit with the by = argument:

```
left_join(
  flights,
  planes %>% rename(year_built = year), ## Not necessary w/ below line, but helpful
  by = "tailnum" ## Be specific about the joining column
  ) %>%
  select(year, month, day, dep_time, arr_time, carrier, flight, tailnum, year_built, type,
  head(5) ## Just to save vertical space on the slide
```

```
## # A tibble: 5 × 11
     year month day dep time arr time carrier flight tailnum year built type
##
    <int> <int> <int> <int> <int> <chr>
                                                           <int> <chr>
###
                                           1545 N14228
                                                            1999 Fixed w...
     2013
                        517
                               830 UA
## 1
                                                            1998 Fixed w...
## 2
    2013
                       533
                           850 UA
                                           1714 N24211
                        542
                                                            1990 Fixed w...
## 3
    2013
                           923 AA
                                           1141 N619AA
                        544
                              1004 B6
                                            725 N804JB
                                                            2012 Fixed w...
     2013
## 4
```

#### Be specific

What happens if we again specify the join column but don't rename the ambiguous "year"?

```
left_join(flights, planes, ## Not renaming "year" to "year_built" this time
by = "tailnum") %>%
select(contains("year"), month, day, dep_time, arr_time, carrier, flight, tailnum, type,
head(5)
```

```
## # A tibble: 5 × 11
                         day dep time arr time carrier flight tailnum type model
##
     vear.x year.y month
      <int> <int> <int> <int>
                               <int>
                                           <int> <chr>
                                                        <int> <chr> <chr> <chr>
##
                                             830 UA
## 1
      2013
             1999
                                    517
                                                           1545 N14228 Fixe... 737-...
                                                           1714 N24211 Fixe... 737-...
## 2
      2013
              1998
                                    533
                                             850 UA
                                    542
                                        923 AA
                                                           1141 N619AA Fixe... 757-...
      2013
              1990
## 3
      2013
              2012
                                    544
                                            1004 B6
                                                            725 N804JB Fixe... A320...
## 4
## 5
      2013
              1991
                                    554
                                             812 DL
                                                            461 N668DN
                                                                        Fixe... 757-...
```

#### Other dplyr functions

pull(): Extract a column from a data frame as a vector or scalar.

• e.g. starwars %>% filter(gender="female") %>% pull(height)

count() and distinct(): Number and isolate unique observations.

- e.g. starwars %>% count(species), Of starwars %>% distinct(species)
- You could also use a combination of mutate(), group\_by(), and n(), e.g. starwars
   %>% group\_by(species) %>% mutate(num = n())
- Built-in combination using add\_count().

#### Other dplyr functions

There is also a whole class of window functions for getting leads and lags, ranking, creating cumulative aggregates, etc.

See vignette("window-functions") for more.

#### Other summaries

- Summary statistics can get cumbersome with summarize()
- Quick summary of selected variables: sumtable() from the vtable package

#### Some dplyr tips

- Any group\_by() statement stays until ungroup()
- Look out for plyr package. Do not use plyr and dplyr together. Just don't do it.



# tidyr

### Key tidyr verbs

- 1. pivot\_wider() and pivot\_longer() to reshape data between wide and long format
- 2. separate(): Split one column into multiple columns
- 3. unite(): Combine multiple columns into one

## 1) tidyr::pivot\_longer()

```
stocks 		 tibble(
  time = as.Date('2009-01-01') + 0:1,
  X = rnorm(2, 0, 1),
  Y = rnorm(2, 0, 2),
  Z = rnorm(2, 0, 4)
  )
stocks
```

## 1) tidyr::pivot\_longer()

```
## # A tibble: 6 × 3
    time stock price
###
   <date> <chr> <dbl>
##
## 1 2009-01-01 X
                   -0.456
## 2 2009-01-01 Y -2.89
## 3 2009-01-01 Z
                   -0.221
## 4 2009-01-02 X
                 1.36
  5 2009-01-02 Y
                 0.482
## 6 2009-01-02 Z
                   -6.97
```

## 1) tidyr::pivot\_longer()

Aside: Remembering the syntax

There's a long-running joke about no-one being able to remember Stata's "reshape" command. (Exhibit A.)

It's easy to see this happening with pivot\_wider() and pivot\_longer() too.

## 1) tidyr::pivot\_wider()

## 2 2009-01-02 1.36 0.482 -6.97

# 2) tidyr::separate()

<chr>

## 1 Abhijit Banerjee

## 2 Esther Duflo

## 3 Michael Kremer

##

```
economists ← tibble(
  name = c("Abhijit Banerjee", "Esther Duflo", "Michael Kremer")
)
economists

## # A tibble: 3 × 1
## name
```

## 2) tidyr::separate()

Should also specify the separation character with separate( ..., sep=" ").

## 3) tidyr::separate\_rows()

A related function is separate\_rows() for splitting into new rows

```
jobs ← tibble(
  name = c("Jack", "Jill"),
  occupation = c("Homemaker", "Philosopher, Philanthropist, Troublemaker")
  )
jobs
```

```
## # A tibble: 2 × 2
## name occupation
## <chr> <chr>
## 1 Jack Homemaker
## 2 Jill Philosopher, Philanthropist, Troublemaker
```

## 3) tidyr::separate\_rows()

```
## Now split out Jill's various occupations into different rows
jobs %>% separate_rows(occupation)
```

```
## # A tibble: 4 × 2
## name occupation
## <chr> <chr> <hr> ## 1 Jack Homemaker
## 2 Jill Philosopher
## 3 Jill Philanthropist
## 4 Jill Troublemaker
```

```
gdp 		 data.frame(
    yr = rep(2016, times = 4),
    mnth = rep(1, times = 4),
    dy = 1:4,
    gdp = rnorm(4, mean = 100, sd = 2)
    )
gdp
```

```
## Combine "yr", "mnth", and "dy" into one "date" column
gdp %>% unite(date, c("yr", "mnth", "dy"), sep = "-")
```

```
## date gdp
## 1 2016-1-1 99.04880
## 2 2016-1-2 103.95492
## 3 2016-1-3 99.72489
## 4 2016-1-4 100.43369
```

unite() automatically creates character variable:

Use mutate() with lubridate date functions to change the variable type.

```
library(lubridate)
gdp_u %>% mutate(date = ymd(date))

## # A tibble: 4 × 2

## date gdp

## <date> <dbl>
## 1 2016-01-01 99.0

## 2 2016-01-02 104.

## 3 2016-01-03 99.7

## 4 2016-01-04 100.
```

## Other tidyr functions

- drop\_na() to drop missing values among specified columns
- fill() to impute missing values from past/future values
- replace\_na() to replace missing values with known value

### Summary

#### dplyr

- 1. filter()
- 2. arrange()
- 3. select()
- 4. mutate()
- 5. summarise()

#### tidyr

- 1. pivot\_longer()
- 2. pivot\_wider()
- 3. separate()
- 4. unite()

Other useful items include: pipes (%>%), grouping (group\_by()), joining functions (left\_join(), inner\_join, etc.).

## Real World

## Practice data versus the real world



## Advice 1: Be patient and careful in your coding



## Advice 2: Comment, comment, comment

You don't want to end up like this guy...



# Medicare Advantage

## Medicare Advantage

Let's work with the Medicare Advantage GitHub repository

#### Access the data

First step is to download the raw data that we'll be using, or work in AWS:

- Monthly Enrollment
- Plan Characteristics
- Service Areas

Lots more out there, but this is enough for now.

```
for (y in 2006:2015) {
  monthlist=get(paste0("monthlist ",y))
  for (m in monthlist) {
    ## Basic contract/plan information
    ma.path=paste0("...CPSC_Contract_Info_",y,"_",m,".csv")
    contract.info=read csv(ma.path,
                            skip=1,
                            col names = c("contractid", "planid", "org type", "plan type",
                                          "partd", "snp", "eghp", "org_name", "org_marketing_na
                                          "plan name", "parent org", "contract date"),
                            col types = cols(
                              contractid = col character(),
                              planid = col_double(),
```

```
## Clean the contract level data
contract.info = contract.info %>%
    group_by(contractid, planid) %>%
    mutate(id_count=row_number())

contract.info = contract.info %>%
    filter(id_count=1) %>%
    select(-id_count)
```

```
## Enrollments per plan
ma.path=paste0("...CPSC_Enrollment_Info_",y,"_",m,".csv")
enroll.info=read csv(ma.path,
                     skip=1,
                     col_names = c("contractid", "planid", "ssa", "fips", "state", "county"
                     col types = cols(
                       contractid = col character(),
                       planid = col double(),
                       ssa = col double(),
                       fips = col double(),
                       state = col character(),
                       county = col character(),
                       enrollment = col double()
                     ),na="*")
```

```
## Merge contract info with enrollment info
plan.data = contract.info %>%
   left_join(enroll.info, by=c("contractid", "planid")) %>%
   mutate(month=as.numeric(m),year=y)
assign(paste0("plan.data.",m),plan.data)
}
```

```
## Fill in missing fips codes (by state and county)
plan.month = plan.month %>%
  group by(state, county) %>%
  fill(fips)
## Fill in missing plan characteristics by contract and plan id
plan.month = plan.month %>%
  group by(contractid, planid) %>%
  fill(plan type, partd, snp, eghp, plan name)
## Fill in missing contract characteristics by contractid
plan.month = plan.month %>%
  group by(contractid) %>%
 fill(org_type,org_name,org_marketing_name,parent_org)
```

```
## Collapse from monthly data to yearly
plan.year = plan.month %>%
  group by(contractid, planid, fips) %>%
  arrange(contractid,planid,fips,month) %>%
  summarize(avg enrollment=mean(enrollment),sd enrollment=sd(enrollment),
            min enrollment=min(enrollment), max enrollment=max(enrollment),
            first enrollment=first(enrollment), last enrollment=last(enrollment),
            state=last(state), county=last(county), org type=last(org type),
            plan type=last(plan type),partd=last(partd),snp=last(snp),
            eghp=last(eghp),org name=last(org name),org marketing name=last(org marketin
            plan name=last(plan name), parent org=last(parent org), contract date=last(con
            vear=last(vear))
write_rds(plan.year,paste0(path.data.final,"/ma_data_",y,".rds"))
```

```
full.ma.data 
    readRDS(paste0(path.data.final,"/ma_data_2006.rds"))

for (y in 2007:2015) {
    full.ma.data 
    rbind(full.ma.data, paste0(path.data.final,"/ma_data_",y,".rds"))
}
```

### All together now

Now let's move to AWS and try some of this together.

- 1. Clone repository
- 2. Create your own repository
- 3. Remove remote, git remote rm origin
- 4. Set new remote git remote add origin NEW\_URL (where NEW\_URL is your own repository)
- 5. git push origin HEAD:master
- 6. git push --set-upstream origin master

## All together now

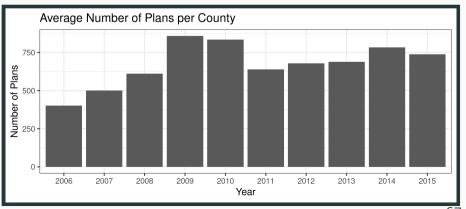
Next, you'll need to make some changes to get things to work:

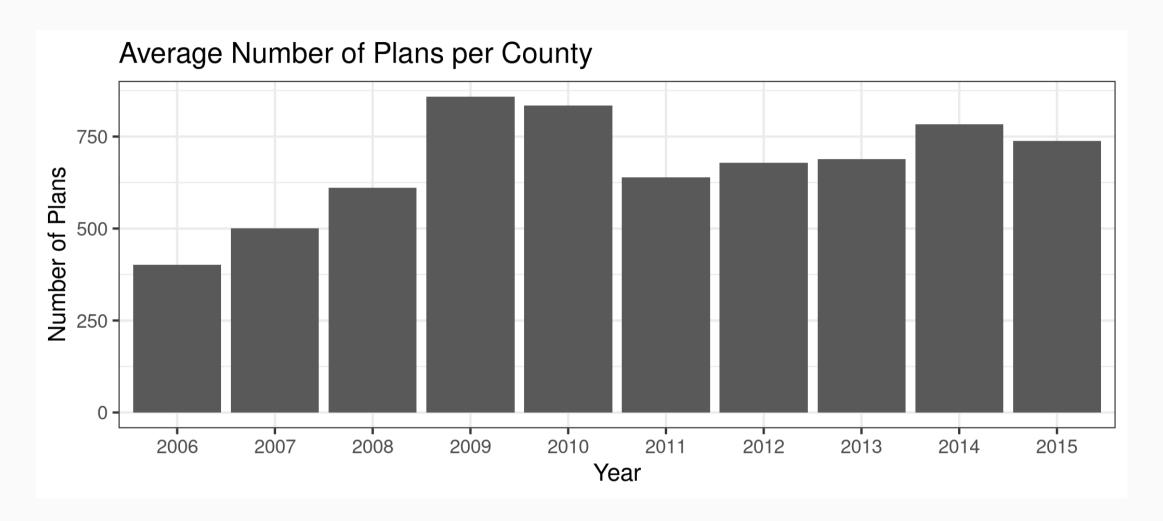
1. Create new paths.R script with:

```
path.code="R_code"
path.data.ma="/data/medicare-advantage"
path.data.ffs="/data/medicare-ffs"
```

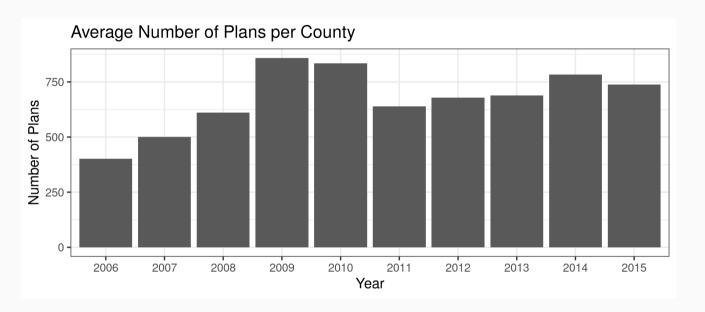
- 1. Update file paths
- 2. Remove "path.data.final" and rely on here() package

```
full.ma.data %>% group_by(fips, year) %>% select(fips, year) %>% summarize(plan_count=n())
    ggplot(aes(x=as.factor(year),y=plan_count)) +
    stat_summary(fun.y="mean", geom="bar") +
    labs(
         x="Year",
         y="Number of Plans",
         title="Average Number of Plans per County"
    ) + scale_y_continuous(labels=comma) +
    theme_bw()
```

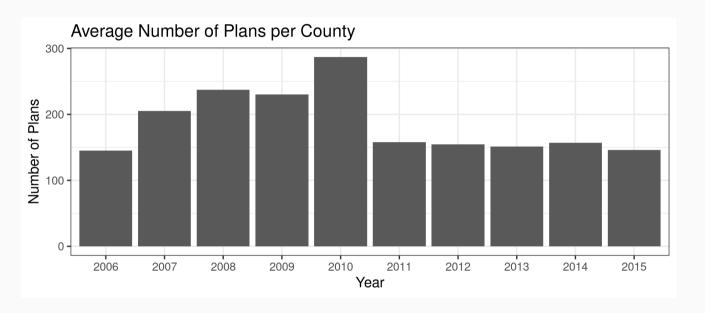




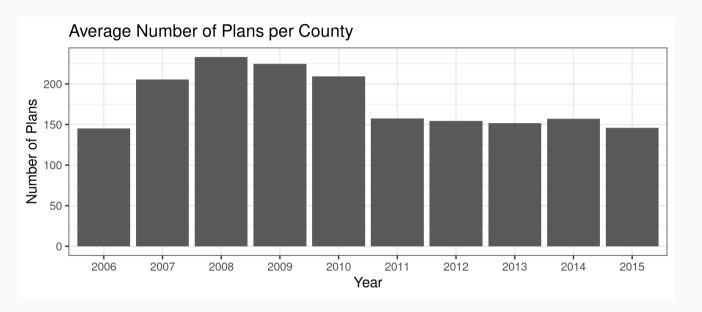
```
full.ma.data %>%
  group_by(fips, year) %>%
  select(fips, year) %>%
  summarize(plan_count=n()) %>%
  ggplot(aes(x=as.factor(year),y=p
  stat_summary(fun.y="mean", geom=
  labs(x="Year", y="Number of Plar
  scale_y_continuous(labels=comma)
  theme_bw()
```



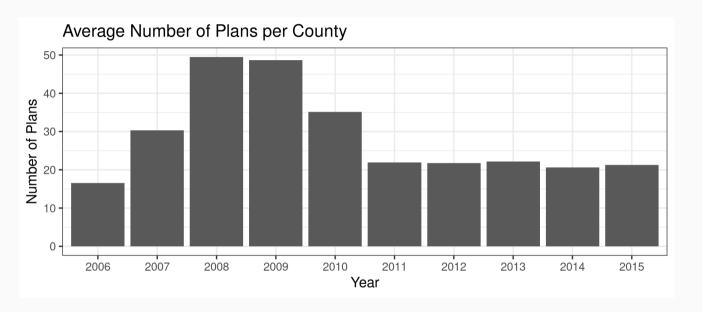
```
full.ma.data %>%
  filter(snp="No" & eghp="No") %
  group_by(fips, year) %>%
  select(fips, year) %>%
  summarize(plan_count=n()) %>%
  ggplot(aes(x=as.factor(year),y=p)
  stat_summary(fun.y="mean", geom=labs(x="Year", y="Number of Plar)
  scale_y_continuous(labels=comma)
  theme_bw()
```



```
full.ma.data %>%
  filter(snp="No" & eghp="No") %
  filter(planid < 800 | planid ≥
  filter(!is.na(planid)) %>%
  group by(fips, year) %>%
  select(fips, year) %>%
  summarize(plan count=n()) %>%
  ggplot(aes(x=as.factor(year),y=r
  stat summary(fun.y="mean", geom=
  labs(x="Year", y="Number of Plar
  scale y continuous(labels=comma)
  theme bw()
```



```
full.ma.data %>%
  filter(snp="No" & eghp="No") %
  filter(planid < 800 | planid ≥
  filter(!is.na(planid)) %>%
  inner join(service.area %>%
               select(contractid,
             by=c("contractid", "f
  group by(fips, year) %>%
  select(fips, year) %>%
  summarize(plan count=n()) %>%
  ggplot(aes(x=as.factor(year),y=r
  stat summary(fun.y="mean", geom=
  labs(x="Year", y="Number of Plar
  scale y continuous(labels=comma)
  theme bw()
```



## Interactive plot

