Module 0: Introduction

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Econ 470 & HLTH 470

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Motivation

Does wealth improve health?

Health and Health Care

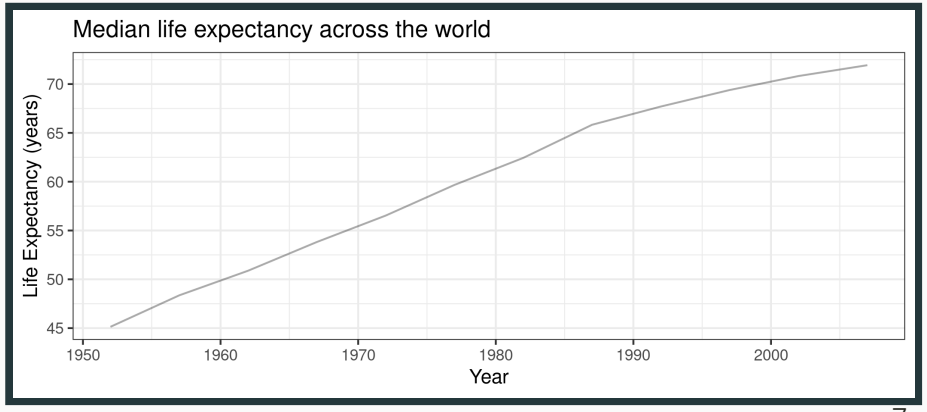
Health and Health Care

We've made *major* improvements in life expectancy (and many other measures of health) across the world

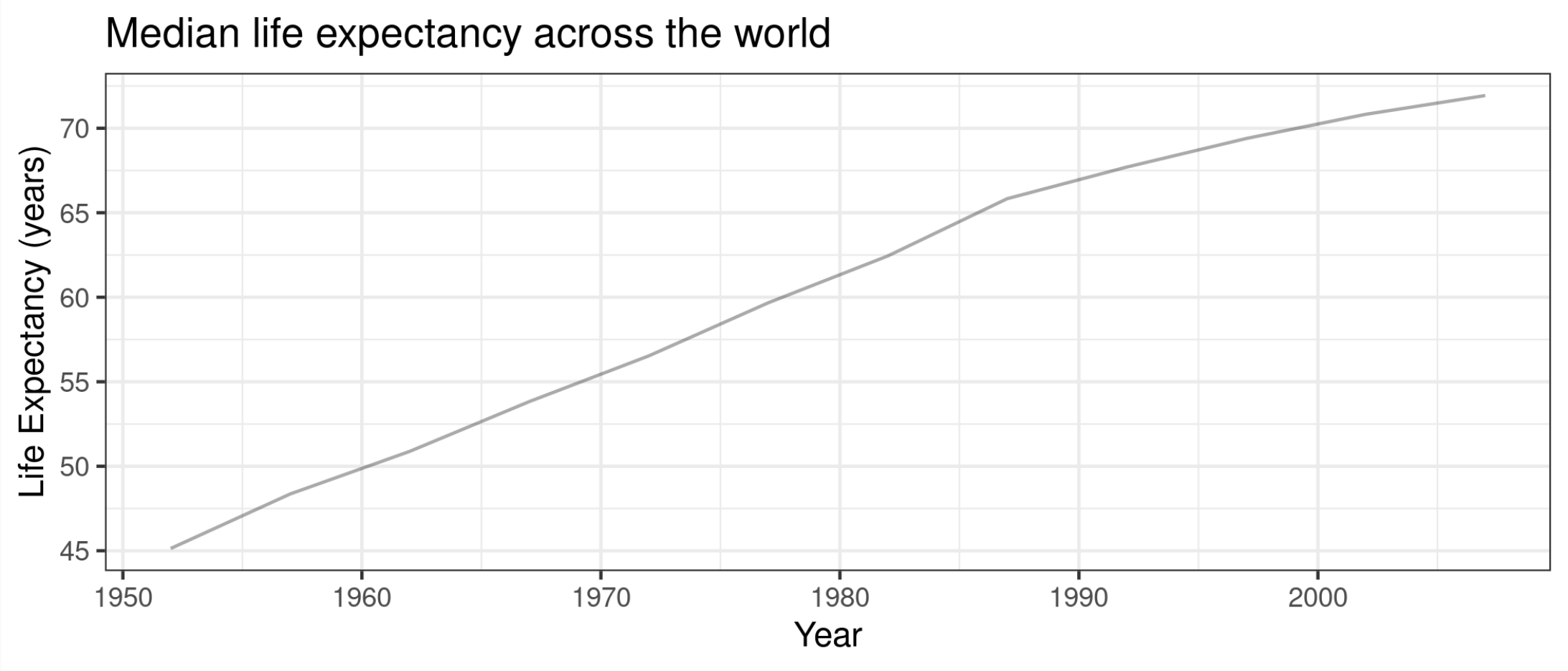
- Poverty reduction
- Technology development and innovation
- Technology diffusion and adoption
- Access to better services, including health care

Evidence of better health

```
library(gapminder)
gapminder %>%
  group_by(year) %>%
  summarize(lifeExp = median(lifeExp),
            gdpMed = median(gdpPercap)) %>%
  ggplot(aes(year,lifeExp)) + geom_line(alpha = 1/3) + theme_bw() +
    labs(x = "Year",
         y = "Life Expectancy (years)",
         title = "Median life expectancy across the world")
```

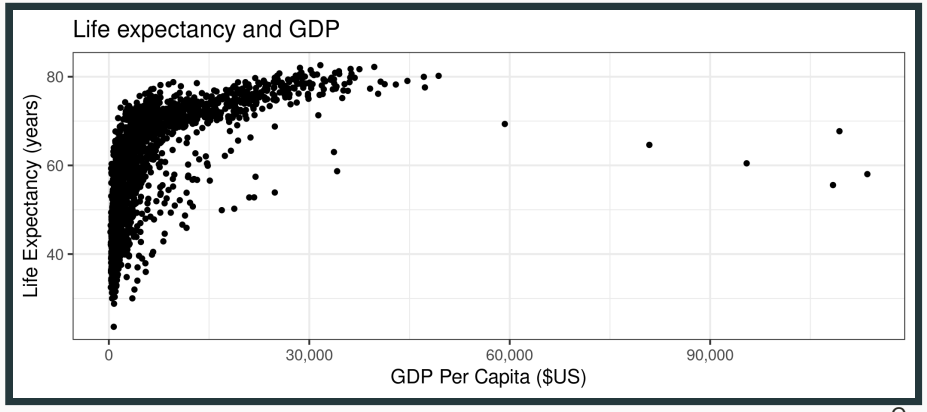


Evidence of better health

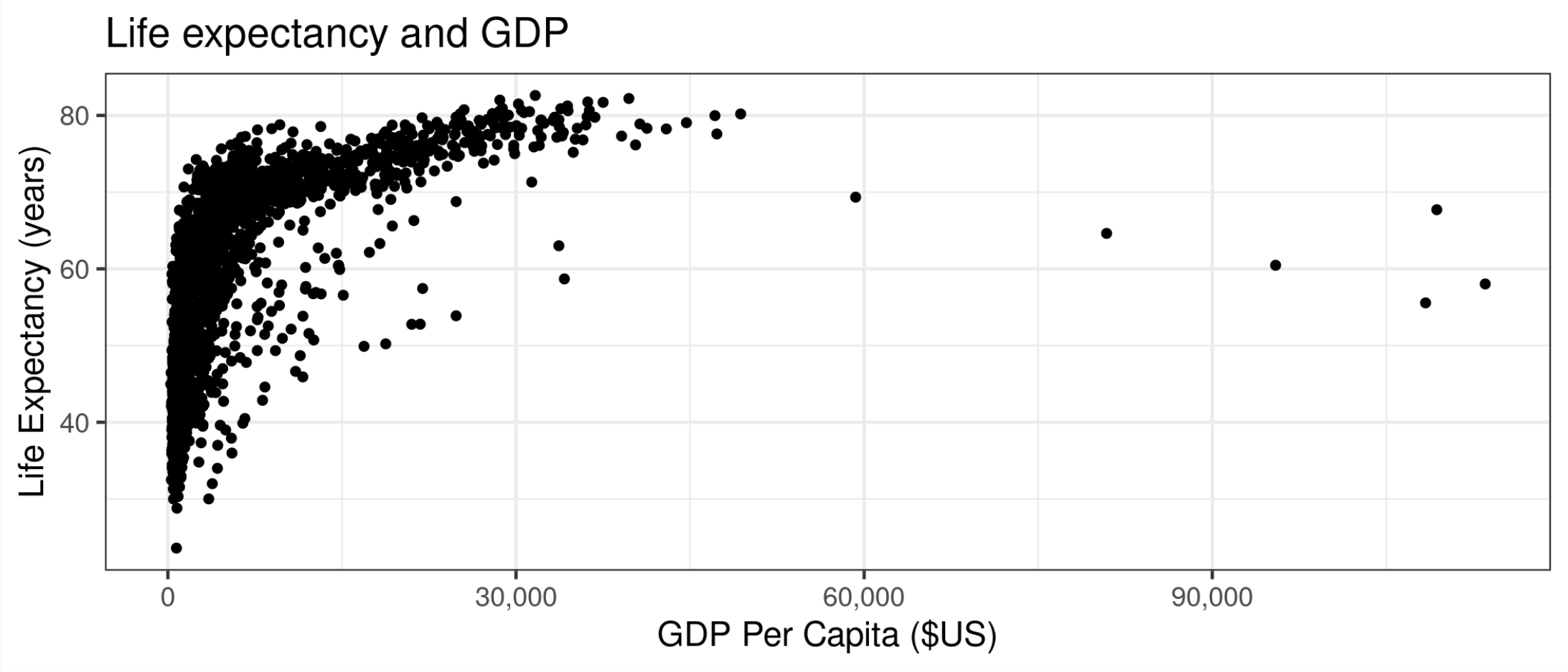


GDP and Health

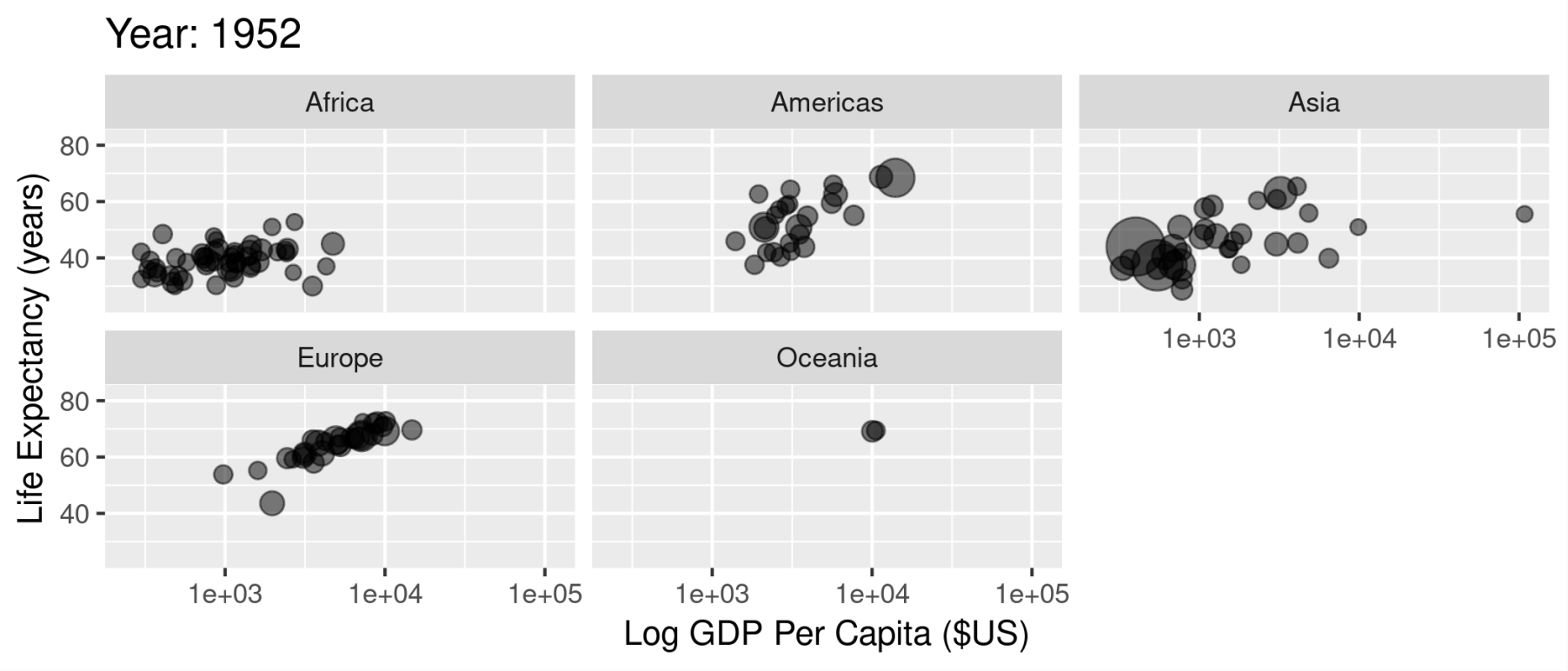
```
library(gapminder)
ggplot(data = gapminder, mapping = aes(x = gdpPercap, y = lifeExp)) +
  geom_point(size = 1) + theme_bw() + scale_x_continuous(label = comma) +
  labs(x = "GDP Per Capita ($US)",
       y = "Life Expectancy (years)",
       title = "Life expectancy and GDP")
```



GDP and Health



GDP and Health over Time



What does this mean?

- Pretty clear relationship between wealth and health (at least by some measures)
- How does this relationship emerge? What are the "mechanisms"?

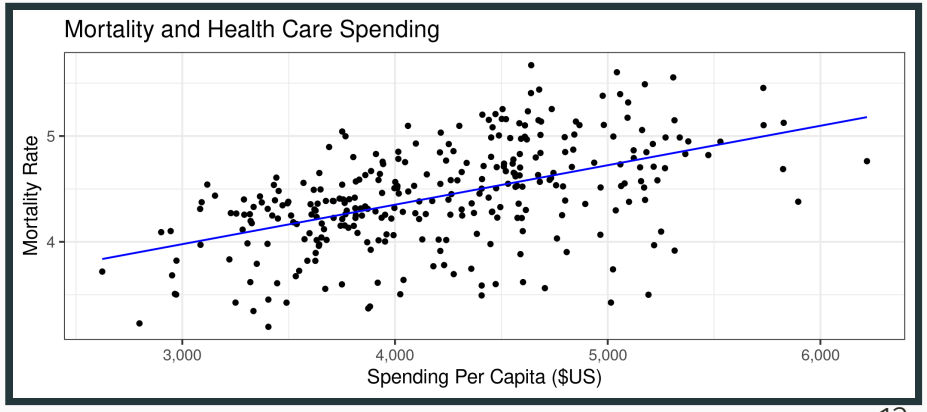
- Classify these into two areas:

  - Health behaviors and related things like education
  - Health care consumption

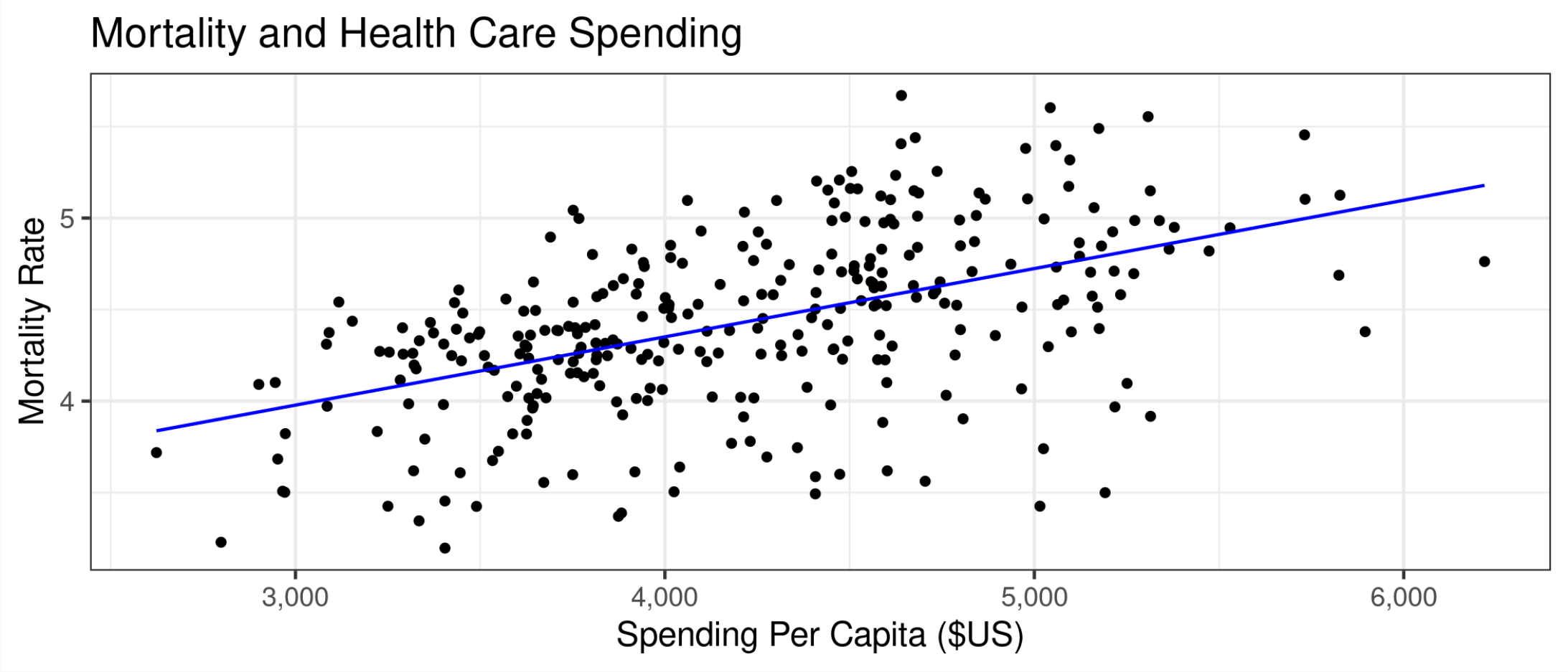
- Does health care spending improve health?

Spending and Health

```
ggplot(data = (dartmouth.data %>% filter(Year==2015)),
       mapping = aes(x = Expenditures, y = Total_Mortality)) +
  geom_point(size = 1) + theme_bw() + scale_x_continuous(label = comma) +
  geom_smooth(method="lm", se=FALSE, color="blue", size=1/2) +
  labs(x = "Spending Per Capita ($US)",
       y = "Mortality Rate",
       title = "Mortality and Health Care Spending")
```

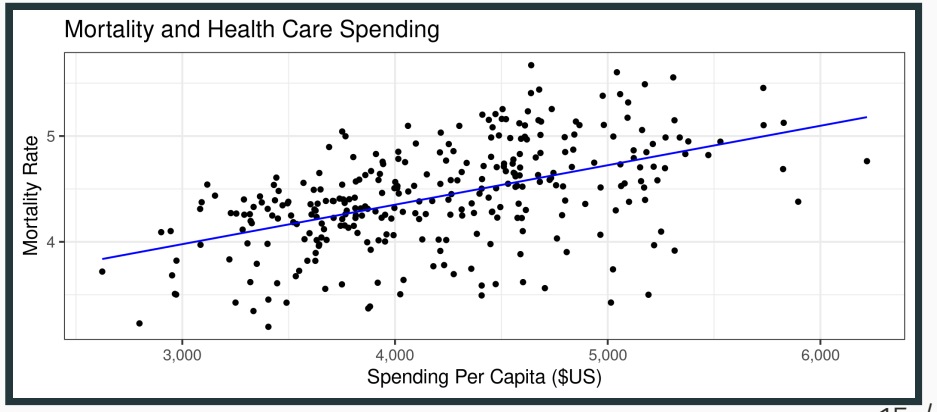


Spending and Health



Spending and Health

- Does medical spending make us sicker?
- What else might explain this relationship?



Goals of this course.

1. Understand and implement selected methods for causal inference
2. Along the way...data management and version control with real data
3. Summarize, visualize, and explain research results

Syllabus highlights

(Read the full document here.)

Why this course?

1. Working with data is hard

2. Health and health care are constantly changing

3. $3.5 trillion, 17.9% of GDP, $10,739 per person

> *"Nobody knew health care could be so complicated"*
>
> --- Donald Trump

Structure

- Very applied in nature
- *Methods* for causal inference
  - Selection on observables (matching, propensity scores)
  - Instrumental variables
  - Regression discontinuity
  - Difference-in-differences

Structure

- *Substantive* areas
  - Hospital pricing, policy, and competition
  - Cigarette taxes and demand
  - Medicare Advantage and quality disclosure
  - Medicaid expansion and health insurance

Structure

- *Datasets* from the real world
  - Hospital Cost Report Information System (HCRIS)
  - Centers for Disease Control (CDC)
  - Medicare Advantage data
  - Behavioral Risk Factor Surveillance System (BRFSS), Medicaid, Health Insurance Exchanges

Assignments

- Homework (x5)
- Research project
- Participation

Grading

Component	Weight
5 × homework assignments (11% each)	55%
Research project	40%
Participation	5%

Software Installation

Software Installation

1. Download R
2. Download RStudio
3. Download Git
4. Create an account on GitHub

For help and troubleshooting with Git and GitHub, take a look at Jenny Bryan's http://happygitwithr.com.

Checklist

☑ Do you have the most recent version of R?

```
version$version.string
```

```
## [1] "R version 4.1.2 (2021-11-01)"
```

☑ Do you have the most recent version of RStudio? (The preview version is fine.)

```
RStudio.Version()$version
## Requires an interactive session but should return something like "[1] ‘1.4.1717’"
```

☑ Have you updated all of your R packages?

```
update.packages(ask = FALSE, checkBuilt = TRUE)
```

Checklist

- Open up the shell
- Windows users, make sure that you installed a Bash-compatible version of the shell. If you installed Git for Windows, then you should be good to go.

Checklist

☑ Which version of Git have you installed?

```
git --version
```

☑ Did you introduce yourself to Git? (Substitute in your details.)

```
git config --global user.name 'Ian McCarthy'
git config --global user.email 'ian.mccarthy@emory.edu'
git config --global --list
```

☑ Did you register an account in GitHub?

Altnernative setup...

Just use the cloud!

- We have our own virtual computer via AWS
- This computer has all the space you need and the data are already available
- Downside: the computer will be "on" during designated times only
- Details of this will be on Canvas shortly

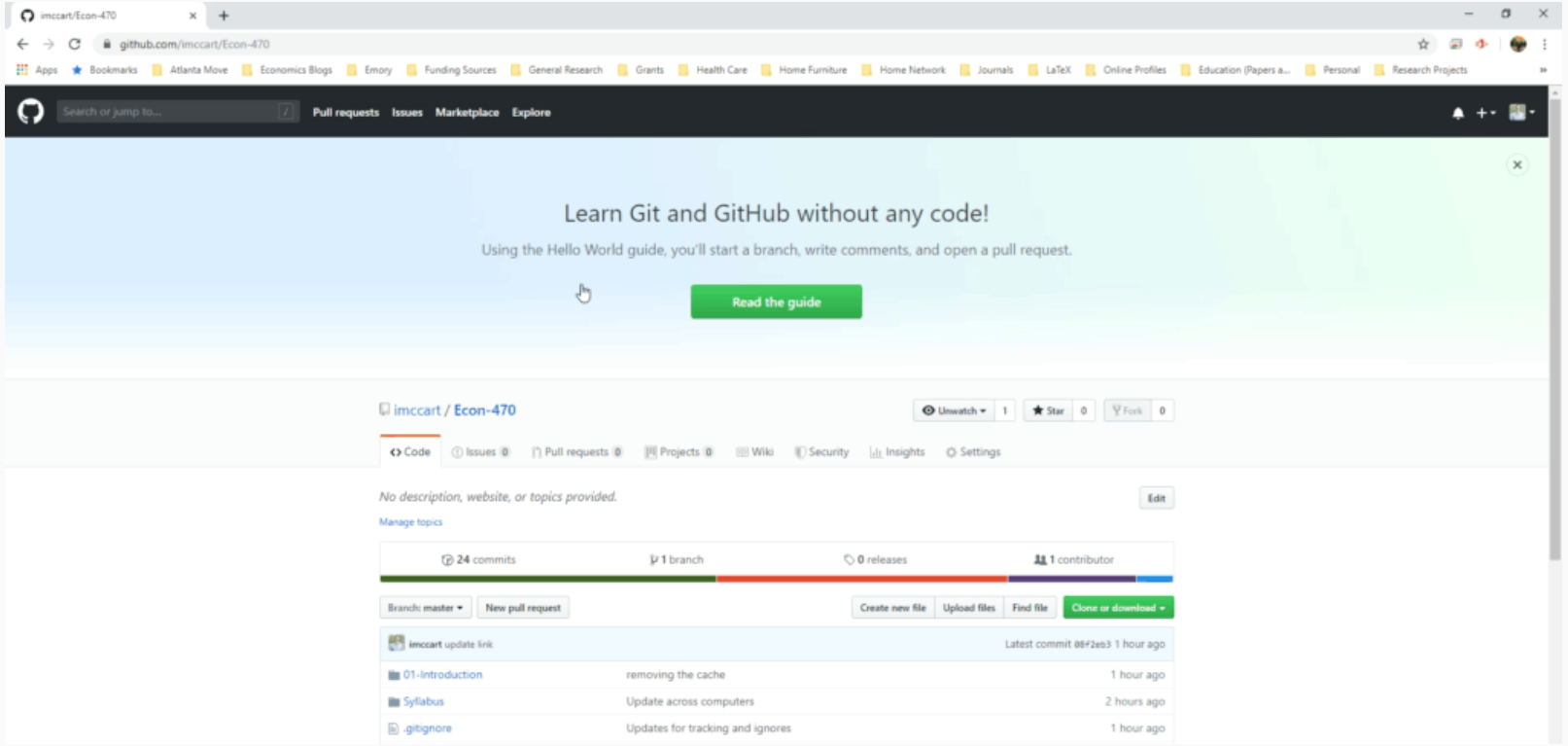
Practice with Git and RStudio

Before next class (see http://happygitwithr.com)

1. Download R
2. Download RStudio
3. Download Git
4. Create an account on GitHub
5. Connect RStudio to Git and GitHub
6. Start/clone/fork a repository for this class

Setting things up

Now we're going to clone a GitHub repository (repo) using RStudio.



Some common mistakes for windows users

- Windows folders are *not* files...there is no content without a file. You can't commit or push changes without content.
- Let RStudio/GitHub create the directory (main folder) for you.
- If you're working across devices on your own repo, be sure to pull before starting and push afterward.
- Avoid spaces in file names. Avoid them at all costs. *DO NOT PUT SPACES IN YOUR FILE NAMES.*

> *"A space in a file name is a space in your soul."*

Ideal workflow

Until you are a Git(Hub) expert...

1. Start project on GitHub (fork from another repo if needed)

2. Clone to desktop with RStudio

3. See http://happygitwithr.com for instructions on linking your local repo with a new upstream remote

Tidy Data

The tidyverse

- Suite of packages collectively known as the tidyverse
- Different from `base` R in many ways
- The tidyverse with pipes[1] is more intuitive to me

[1] 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: %>%

Helps to break the pipes over several lines

```
mpg %>%
  filter(manufacturer=="audi") %>%
  group_by(model) %>%
  summarise(hwy_mean = mean(hwy))
```

```
## # A tibble: 3 × 2
##   model      hwy_mean
##   <chr>         <dbl>
## 1 a4             28.3
## 2 a4 quattro     25.8
## 3 a6 quattro     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
##   name      height  mass hair_color skin_color eye_color birth_year sex   gender
##   <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…
## 3 Dooku        193    80 white      fair       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 vlaues

```
starwars %>%
  filter(is.na(height)) %>% head(5)
```

```
## # A tibble: 5 × 14
##   name     height  mass hair_color skin_color eye_color birth_year sex   gender
##   <chr>     <int> <dbl> <chr>      <chr>      <chr>           <dbl> <chr> <chr>
## 1 Arvel C…     NA    NA brown      fair       brown              NA male  mascul…
## 2 Finn         NA    NA black      dark       dark               NA male  mascul…
## 3 Rey          NA    NA brown      light      hazel              NA fema… femini…
## 4 Poe Dam…     NA    NA brown      light      brown              NA male  mascul…
## 5 BB8          NA    NA none       none       black              NA none  mascul…
## # … with 5 more variables: homeworld <chr>, species <chr>, films <list>,
## #   vehicles <list>, starships <list>
```

2) dplyr::arrange()

Arrange in ascending order:

```
starwars %>%
  arrange(birth_year) %>% head(5)
```

```
## # A tibble: 5 × 14
##   name      height  mass hair_color skin_color eye_color birth_year sex   gender
##   <chr>      <int> <dbl> <chr>      <chr>      <chr>           <dbl> <chr> <chr>
## 1 Wicket S…     88    20 brown      brown      brown               8 male  mascu…
## 2 IG-88        200   140 none       metal      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
##   name    height  mass hair_color skin_color  eye_color birth_year sex    gender
##   <chr>    <int> <dbl> <chr>      <chr>       <chr>           <dbl> <chr>  <chr>
## 1 Yoda        66    17 white      green       brown             896 male   mascu…
## 2 Jabba …    175  1358 <NA>       green-tan,… orange            600 herma… mascu…
## 3 Chewba…    228   112 brown      unknown     blue              200 male   mascu…
## 4 C-3PO      167    75 <NA>       gold        yellow            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>
```

3) dplyr::select()

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

3) dplyr::select()

```
starwars %>%
  select(name:skin_color, species, -height) %>% head(5)
```

```
## # A tibble: 5 × 5
##   name             mass hair_color skin_color  species
##   <chr>           <dbl> <chr>      <chr>       <chr>
## 1 Luke Skywalker    77 blond      fair        Human
## 2 C-3PO             75 <NA>       gold        Droid
## 3 R2-D2             32 <NA>       white, blue Droid
## 4 Darth Vader      136 none       white       Human
## 5 Leia Organa       49 brown      light       Human
```

3) dplyr::select()

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-3PO          Tatooine masculine
## 3 R2-D2          Naboo    masculine
## 4 Darth Vader    Tatooine masculine
## 5 Leia Organa    Alderaan feminine
```

3) dplyr::select()

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

```
starwars %>%
  select(name, contains("color")) %>% head(5)
```

```
## # A tibble: 5 × 4
##   name           hair_color skin_color  eye_color
##   <chr>          <chr>      <chr>       <chr>
## 1 Luke Skywalker blond      fair        blue
## 2 C-3PO          <NA>       gold        yellow
## 3 R2-D2          <NA>       white, blue red
## 4 Darth Vader    none       white       yellow
## 5 Leia Organa    brown      light       brown
```

Also look into the `stringr` package.

4) dplyr::mutate()

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

4) dplyr::mutate()

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

```
starwars %>%
  select(name, birth_year) %>%
  mutate(dog_years = birth_year * 7, ## Separate with a comma
         comment = paste0(name, " is ", dog_years, " in dog years.")) %>% head(5)
```

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

4) dplyr::mutate()

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
##   name             height tall1 tall2
##   <chr>             <int> <lgl> <chr>
## 1 Luke Skywalker      172 FALSE Short
## 2 Anakin Skywalker    188 TRUE  Tall
```

4) dplyr::mutate()

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)

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()

"scoped" variants also work with `summarise()`

- `summarise_all()` affects every variable
- `summarise_at()` affects named or selected variables
- `summarise_if()` affects variables that meet some criteria (e.g. are numeric)

```
starwars %>% group_by(species, gender) %>% summarise_if(is.numeric, list(avg=mean), na.rm=
```

```
## # A tibble: 5 × 5
## # Groups:   species [5]
##   species  gender    height_avg mass_avg birth_year_avg
##   <chr>    <chr>          <dbl>    <dbl>          <dbl>
## 1 Aleena   masculine         79       15            NaN
## 2 Besalisk masculine        198      102            NaN
## 3 Cerean   masculine        198       82             92
## 4 Chagrian masculine        196      NaN            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
```

Left join

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
##    year month   day dep_time arr_time carrier flight tailnum type  model
##   <int> <int> <int>    <int>    <int> <chr>    <int> <chr>   <chr> <chr>
## 1  2013     1     1      517      830 UA        1545 N14228  <NA>  <NA>
## 2  2013     1     1      533      850 UA        1714 N24211  <NA>  <NA>
## 3  2013     1     1      542      923 AA        1141 N619AA  <NA>  <NA>
## 4  2013     1     1      544     1004 B6         725 N804JB  <NA>  <NA>
## 5  2013     1     1      554      812 DL         461 N668DN  <NA>  <NA>
```

Left join

`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*

Left join

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`.

Left 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>        <int> <chr>
## 1  2013     1     1      517      830 UA        1545 N14228        1999 Fixed w…
## 2  2013     1     1      533      850 UA        1714 N24211        1998 Fixed w…
## 3  2013     1     1      542      923 AA        1141 N619AA        1990 Fixed w…
## 4  2013     1     1      544     1004 B6         725 N804JB        2012 Fixed w…
```

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

Other dplyr goodies

`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)`, or `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 goodies

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.

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
```

```
## # A tibble: 2 × 4
##   time             X      Y      Z
##   <date>       <dbl>  <dbl>  <dbl>
## 1 2009-01-01  0.192 -1.62  -0.780
## 2 2009-01-02 -0.907 -0.332  6.00
```

1) tidyr::pivot_longer()

```
tidy_stocks ← stocks %>%
  pivot_longer(cols=c("X","Y","Z"),
               names_to="stock", values_to="price")
tidy_stocks
```

```
## # A tibble: 6 × 3
##   time       stock  price
##   <date>     <chr>  <dbl>
## 1 2009-01-01 X      0.192
## 2 2009-01-01 Y     -1.62
## 3 2009-01-01 Z     -0.780
## 4 2009-01-02 X     -0.907
## 5 2009-01-02 Y     -0.332
## 6 2009-01-02 Z      6.00
```

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()

```
tidy_stocks %>%
  pivot_wider(values_from="price",names_from="stock")
```

```
## # A tibble: 2 × 4
##   time             X      Y      Z
##   <date>       <dbl>  <dbl>  <dbl>
## 1 2009-01-01   0.192 -1.62  -0.780
## 2 2009-01-02  -0.907 -0.332  6.00
```

2) tidyr::separate()

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

```
## # A tibble: 3 × 1
##   name
##   <chr>
## 1 Abhijit Banerjee
## 2 Esther Duflo
## 3 Michael Kremer
```

2) tidyr::separate()

```
economists %>% separate(name, c("first_name", "last_name"))
```

```
## # A tibble: 3 × 2
##   first_name last_name
##   <chr>      <chr>
## 1 Abhijit    Banerjee
## 2 Esther     Duflo
## 3 Michael    Kremer
```

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>
## 1 Jack  Homemaker
## 2 Jill  Philosopher
## 3 Jill  Philanthropist
## 4 Jill  Troublemaker
```

4) tidyr::unite()

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

```
##     yr mnth dy       gdp
## 1 2016    1  1  98.72684
## 2 2016    1  2  98.91063
## 3 2016    1  3  99.23657
## 4 2016    1  4 105.45198
```

4) tidyr::unite()

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

```
##       date       gdp
## 1 2016-1-1  98.72684
## 2 2016-1-2  98.91063
## 3 2016-1-3  99.23657
## 4 2016-1-4 105.45198
```

4) tidyr::unite()

`unite()` automatically creates character variable:

```
gdp_u ← gdp %>% unite(date, c("yr", "mnth", "dy"), sep = "-") %>% as_tibble()
gdp_u
```

```
## # A tibble: 4 × 2
##   date       gdp
##   <chr>    <dbl>
## 1 2016-1-1  98.7
## 2 2016-1-2  98.9
## 3 2016-1-3  99.2
## 4 2016-1-4 105.
```

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

4) tidyr::unite()

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

```
## # A tibble: 4 × 2
##   date         gdp
##   <date>     <dbl>
## 1 2016-01-01  98.7
## 2 2016-01-02  98.9
## 3 2016-01-03  99.2
## 4 2016-01-04 105.
```

Other tidyr goodies

- `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.).

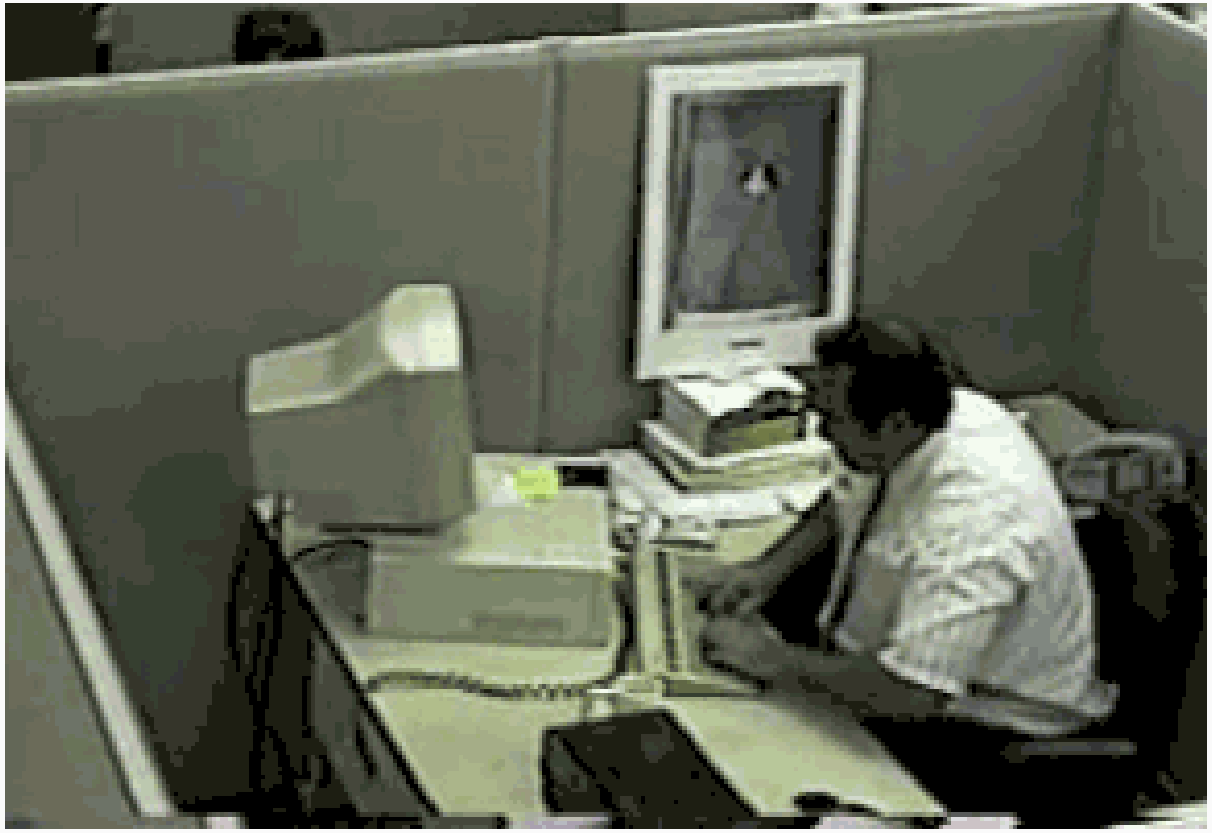
Real World

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

Download the data

First step is to download the raw data that we'll be using.

- Monthly Enrollment
- Plan Characteristics
- Service Areas

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

1) Contract/enrollment info

```
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(),
                             org_type = col_character(),
                             plan_type = col_character(),
                             partd = col_character(),
                             snp = col_character(),
```

1) Contract/enrollment info

```
## 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)
```

1) Contract/enrollment info

```
## 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="*")
```

1) Contract/enrollment info

```
  ## 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.",y,".",m),plan.data)
}
```

1) Contract/enrollment info

```
## Append monthly enrollment info for each year
if (y==2006) {
  plan.month=rbind(plan.data.07, plan.data.08, plan.data.09, plan.data.10,
                   plan.data.11, plan.data.12)
} else {
  plan.month=rbind(plan.data.01, plan.data.02, plan.data.03, plan.data.04,
                   plan.data.05, plan.data.06, plan.data.07, plan.data.08,
                   plan.data.09, plan.data.10, plan.data.11, plan.data.12)
}
```

1) Contract/enrollment info

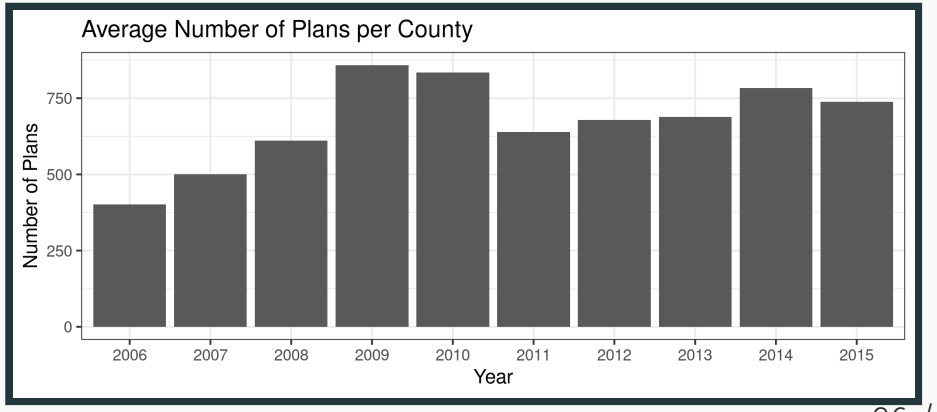
```
## Fill in missing fips codes (by state and county)
plan.month = plan.month %>%
  group_by(state, county) %>%
  fill(fips)

## 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)
```

1) Contract/enrollment info

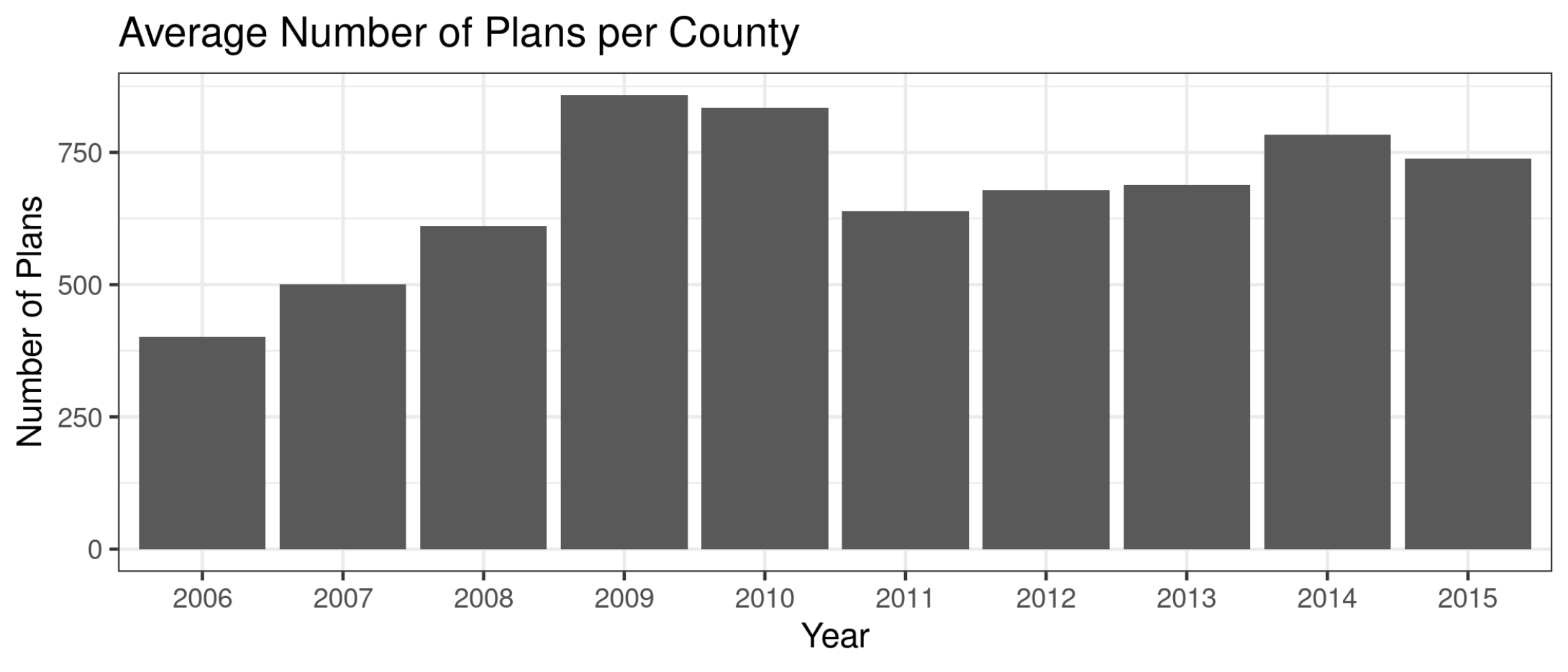
```
  ## Collapse from monthly data to yearly
  plan.year = plan.month %>%
    group_by(contractid, planid, fips) %>%
    arrange(contractid,planid,fips,month) %>%
    summarize(avg_enrollment=mean(enrollment),sd_enrollment=sd(enrollment),
              min_enrollment=min(enrollment),max_enrollment=max(enrollment),
              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
              year=last(year))

  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"))
```

Number of plans available

```
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()
```



Number of plans available



Number of plans available

```
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()
```

Number of plans available

```
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=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()
```

Number of plans available

```
full.ma.data %>%
  filter(snp=="No" & eghp=="No") %>%
  filter(planid < 800 | planid ≥ 900) %>%
  filter(!is.na(planid)) %>%
  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()
```

Number of plans available

```
full.ma.data %>%
  filter(snp=="No" & eghp=="No") %>%
  filter(planid < 800 | planid ≥ 900) %>%
  filter(!is.na(planid)) %>%
  inner_join(service.area %>%
               select(contractid, fips, year),
             by=c("contractid", "fips", "year")) %>%
  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()
```

Interactive plot