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Data Frame Manipulation with dplyr

Overview

Teaching: 40 min Exercises: 15 min Questions

· How can I manipulate data frames without repeating myself?

Objectives

- . To be able to use the six main data frame manipulation 'verbs' with pipes in dplyr.
- To understand how group_by() and summarize() can be combined to summarize datasets.
- · Be able to analyze a subset of data using logical filtering.

Manipulation of data frames means many things to many researchers, we often select certain observations (rows) or variables (columns), we often group the data by a certain variable(s), or we even calculate summary statistics. We can do these operations using the normal base R operations:

R

mean(gapminder[gapminder\$continent == "Africa", "gdpPercap"])

Output

[1] 2193.755

R

mean(gapminder[gapminder\$continent == "Americas", "gdpPercap"])

Output

[1] 7136.11

R

mean(gapminder[gapminder\$continent == "Asia", "gdpPercap"])

Output

[1] 7902.15

But this isn't very nice because there is a fair bit of repetition. Repeating yourself will cost you time, both now and later, and potentially introduce some nasty bugs.

The dplyr package

Luckily, the dplyr (https://cran.r-project.org/package=dplyr) package provides a number of very useful functions for manipulating data frames in a way that will reduce the above repetition, reduce the probability of making errors, and probably even save you some typing. As an added bonus, you might even find the dplyr grammar easier to read.

🖈 Tip: Tidyverse

dplyr package belongs to a broader family of opinionated R packages designed for data science called the "Tidyverse". These packages are specifically designed to work harmoniously together. Some of these packages will be covered along this course, but you can find more complete information here: https://www.tidyverse.org/ (https://www.tidyverse.org/).

Here we're going to cover 5 of the most commonly used functions as well as using pipes (%>%) to combine them.

- 2. filter()
- 3. group_by()
- 4. summarize()
- 5. mutate()

If you have have not installed this package earlier, please do so:

```
R
install.packages('dplyr')
```

Now let's load the package:

R

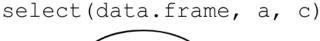
library("dplyr")

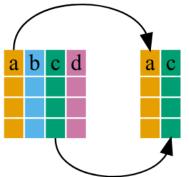
Using select()

If, for example, we wanted to move forward with only a few of the variables in our data frame we could use the select() function. This will keep only the variables you select.

R

year_country_gdp <- select(gapminder, year, country, gdpPercap)</pre>





If we want to remove one column only from the gapminder data, for example, removing the continent column.

```
R
smaller_gapminder_data <- select(gapminder, -continent)</pre>
```

If we open up <code>year_country_gdp</code> we'll see that it only contains the year, country and <code>gdpPercap</code>. Above we used 'normal' grammar, but the strengths of <code>dplyr</code> lie in combining several functions using pipes. Since the pipes grammar is unlike anything we've seen in R before, let's repeat what we've done above using pipes.

```
R

year_country_gdp <- gapminder %>% select(year, country, gdpPercap)
```

To help you understand why we wrote that in that way, let's walk through it step by step. First we summon the gapminder data frame and pass it on, using the pipe symbol %>%, to the next step, which is the select() function. In this case we don't specify which data object we use in the select() function since in gets that from the previous pipe. Fun Fact: There is a good chance you have encountered pipes before in the shell. In R, a pipe symbol is %>% while in the shell it is | but the concept is the same!

★ Tip: Renaming data frame columns in dplyr

In Chapter 4 we covered how you can rename columns with base R by assigning a value to the output of the names() function. Just like select, this is a bit cumbersome, but thankfully dplyr has a rename() function.

Within a pipeline, the syntax is rename (new_name = old_name) . For example, we may want to rename the gdpPercap column name from our select() statement above.

```
R
```

```
tidy_gdp <- year_country_gdp %>% rename(gdp_per_capita = gdpPercap)
```

head(tidy_gdp)

Output

```
year country gdp_per_capita
1 1952 Afghanistan 779.4453
2 1957 Afghanistan 820.8530
3 1962 Afghanistan 853.1007
4 1967 Afghanistan 836.1971
5 1972 Afghanistan 739.9811
6 1977 Afghanistan 786.1134
```

Using filter()

If we now want to move forward with the above, but only with European countries, we can combine select and filter

```
R
year_country_gdp_euro <- gapminder %>%
  filter(continent == "Europe") %>%
  select(year, country, gdpPercap)
```

If we now want to show life expectancy of European countries but only for a specific year (e.g., 2007), we can do as below.

```
R
europe_lifeExp_2007 <- gapminder %>%
filter(continent == "Europe", year == 2007) %>%
select(country, lifeExp)
```

Challenge 1

Write a single command (which can span multiple lines and includes pipes) that will produce a data frame that has the African values for lifeExp, country and year, but not for other Continents. How many rows does your data frame have and why?

Solution to Challenge 1

```
R
```

As with last time, first we pass the gapminder data frame to the filter() function, then we pass the filtered version of the gapminder data frame to the select() function. **Note:** The order of operations is very important in this case. If we used 'select' first, filter would not be able to find the variable continent since we would have removed it in the previous step.

Using group_by()

Now, we were supposed to be reducing the error prone repetitiveness of what can be done with base R, but up to now we haven't done that since we would have to repeat the above for each continent. Instead of filter(), which will only pass observations that meet your criteria (in the above: continent=="Europe"), we can use group_by(), which will essentially use every unique criteria that you could have used in filter.

```
R
```

str(gapminder)

```
Output

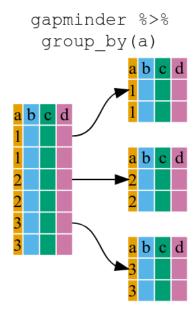
'data.frame': 1704 obs. of 6 variables:
$ country : chr "Afghanistan" "Afghanistan" "Afghanistan" ...
$ year : int 1952 1957 1962 1967 1972 1977 1982 1987 1992 1997 ...
$ pop : num 8425333 9240934 10267083 11537966 13079460 ...
$ continent: chr "Asia" "Asia" "Asia" "Asia" "Asia" ...
$ lifeExp : num 28.8 30.3 32 34 36.1 ...
$ gdpPercap: num 779 821 853 836 740 ...
```

R

str(gapminder %>% group_by(continent))

```
Output
\label{eq:grouped_df/tbl_df/tbl/data.frame)} groupd\_df \ [\textbf{1,704} \times \textbf{6}] \ (S3: grouped\_df/tbl\_df/tbl/data.frame)
 $ country : chr [1:1704] "Afghanistan" "Afghanistan" "Afghanistan" ...
           : int [1:1704] 1952 1957 1962 1967 1972 1977 1982 1987 1992 1997 ...
           : num [1:1704] 8425333 9240934 10267083 11537966 13079460 ...
$ pop
 $ continent: chr [1:1704] "Asia" "Asia" "Asia" "Asia" ...
 $ lifeExp : num [1:1704] 28.8 30.3 32 34 36.1 ...
 $ gdpPercap: num [1:1704] 779 821 853 836 740 ...
 - attr(*, "groups")= tibble [5 x 2] (S3: tbl_df/tbl/data.frame)
  ..$ continent: chr [1:5] "Africa" "Americas" "Asia" "Europe" ...
  ..$ .rows : list<int> [1:5]
  ....$ : int [1:624] 25 26 27 28 29 30 31 32 33 34 ...
  ....$ : int [1:300] 49 50 51 52 53 54 55 56 57 58 ...
  ....$: int [1:396] 1 2 3 4 5 6 7 8 9 10 ...
  ....$: int [1:360] 13 14 15 16 17 18 19 20 21 22 ...
  .. ..$ : int [1:24] 61 62 63 64 65 66 67 68 69 70 ...
  .. ..@ ptype: int(0)
  ..- attr(*, ".drop")= logi TRUE
```

You will notice that the structure of the data frame where we used <code>group_by()</code> (<code>grouped_df</code>) is not the same as the original <code>gapminder</code> (<code>data.frame</code>). A <code>grouped_df</code> can be thought of as a <code>list</code> where each item in the <code>list</code> is a <code>data.frame</code> which contains only the rows that correspond to the a particular value <code>continent</code> (at least in the example above).



Using summarize()

The above was a bit on the uneventful side but $group_by()$ is much more exciting in conjunction with summarize(). This will allow us to create new variable(s) by using functions that repeat for each of the continent-specific data frames. That is to say, using the $group_by()$ function, we split our original data frame into multiple pieces, then we can run functions (e.g. mean() or sd()) within summarize().

```
R
gdp_bycontinents <- gapminder %>%
    group_by(continent) %>%
    summarize(mean_gdpPercap = mean(gdpPercap))
```

```
gapminder %>%
group_by(a) %>%
summarize(mean_b=mean(b))

a b c d

a b c d

a b c d

a b c d

a b c d

a b c d

a b c d

a b c d

a b c d

a b c d

a b c d
```

That allowed us to calculate the mean gdpPercap for each continent, but it gets even better.

🖍 Challenge 2

Calculate the average life expectancy per country. Which has the longest average life expectancy and which has the shortest average life expectancy?

Solution to Challenge 2

```
IfeExp_bycountry <- gapminder %>%
    group_by(country) %>%
    summarize(mean_lifeExp = mean(lifeExp))
lifeExp_bycountry %>%
    filter(mean_lifeExp == min(mean_lifeExp) | mean_lifeExp == max(mean_lifeExp))
```

Output

Another way to do this is to use the dplyr function arrange(), which arranges the rows in a data frame according to the order of one or more variables from the data frame. It has similar syntax to other functions from the dplyr package. You can use desc() inside arrange() to sort in descending order.

```
R
lifeExp_bycountry %>%
   arrange(mean_lifeExp) %>%
   head(1)
```

Output

R

```
lifeExp_bycountry %>%
  arrange(desc(mean_lifeExp)) %>%
  head(1)
```

Output

Alphabetical order works too

R

```
lifeExp_bycountry %>%
  arrange(desc(country)) %>%
  head(1)
```

Output

The function $group_by()$ allows us to group by multiple variables. Let's group by year and continent .

```
gdp_bycontinents_byyear <- gapminder %>%
group_by(continent, year) %>%
summarize(mean_gdpPercap = mean(gdpPercap))
```

Output

```
`summarise()` has grouped output by 'continent'. You can override using the `.groups` argument.
```

That is already quite powerful, but it gets even better! You're not limited to defining 1 new variable in summarize().

```
gdp_pop_bycontinents_byyear <- gapminder %>%
    group_by(continent, year) %>%
    summarize(mean_gdpPercap = mean(gdpPercap),
        sd_gdpPercap = sd(gdpPercap),
        mean_pop = mean(pop),
        sd_pop = sd(pop))
```

Output

```
`summarise()` has grouped output by 'continent'. You can override using the `.groups` argument.
```

count() and n()

A very common operation is to count the number of observations for each group. The dplyr package comes with two related functions that help with this.

For instance, if we wanted to check the number of countries included in the dataset for the year 2002, we can use the <code>count()</code> function. It takes the name of one or more columns that contain the groups we are interested in, and we can optionally sort the results in descending order by adding <code>sort=TRUE</code>:

```
R
gapminder %>%
  filter(year == 2002) %>%
  count(continent, sort = TRUE)
```

```
Output

continent n
1 Africa 52
2 Asia 33
3 Europe 30
4 Americas 25
5 Oceania 2
```

If we need to use the number of observations in calculations, the n() function is useful. It will return the total number of observations in the current group rather than counting the number of observations in each group within a specific column. For instance, if we wanted to get the standard error of the life expectency per continent:

```
R
gapminder %>%
    group_by(continent) %>%
    summarize(se_le = sd(lifeExp)/sqrt(n()))
```

```
# A tibble: 5 × 2
continent se_le
<chr> <dbl>
1 Africa 0.366
2 Americas 0.540
3 Asia 0.596
4 Europe 0.286
5 Oceania 0.775
```

You can also chain together several summary operations; in this case calculating the minimum, maximum, mean and se of each continent's per-country life-expectancy:

```
gapminder %>%
   group_by(continent) %>%
   summarize(
   mean_le = mean(lifeExp),
   min_le = min(lifeExp),
   max_le = max(lifeExp),
   se_le = sd(lifeExp)/sqrt(n()))
```

```
Output
# A tibble: 5 \times 5
  continent mean_le min_le max_le se_le
  <chr>
             <dbl> <dbl> <dbl> <dbl>
1 Africa
              48.9
                    23.6
                           76.4 0.366
              64.7 37.6
                           80.7 0.540
2 Americas
              60.1 28.8 82.6 0.596
3 Asia
4 Europe
              71.9 43.6 81.8 0.286
5 Oceania
              74.3 69.1 81.2 0.775
```

Using mutate()

We can also create new variables prior to (or even after) summarizing information using mutate().

```
gdp_pop_bycontinents_byyear <- gapminder %>%
    mutate(gdp_billion = gdpPercap*pop/10^9) %>%
    group_by(continent,year) %>%
    summarize(mean_gdpPercap = mean(gdpPercap),
        sd_gdpPercap = sd(gdpPercap),
        mean_pop = mean(pop),
        sd_pop = sd(pop),
        mean_gdp_billion = mean(gdp_billion),
        sd_gdp_billion = sd(gdp_billion))
```

```
Output

`summarise()` has grouped output by 'continent'. You can override using the
`.groups` argument.
```

Connect mutate with logical filtering: ifelse

When creating new variables, we can hook this with a logical condition. A simple combination of mutate() and ifelse() facilitates filtering right where it is needed: in the moment of creating something new. This easy-to-read statement is a fast and powerful way of discarding certain data (even though the overall dimension of the data frame will not change) or for updating values depending on this given condition.

```
## keeping all data but "filtering" after a certain condition
# calculate GDP only for people with a life expectation above 25
gdp_pop_bycontinents_byyear_above25 <- gapminder %>%
    mutate(gdp_billion = ifelse(lifeExp > 25, gdpPercap * pop / 10^9, NA)) %>%
    group_by(continent, year) %>%
    summarize(mean_gdpPercap = mean(gdpPercap),
        sd_gdpPercap = sd(gdpPercap),
        mean_pop = mean(pop),
        sd_pop = sd(pop),
        mean_gdp_billion = mean(gdp_billion),
        sd_gdp_billion = sd(gdp_billion))
```

```
Output
```

```
`summarise()` has grouped output by 'continent'. You can override using the `.groups` argument.
```

```
## updating only if certain condition is fullfilled
# for life expectations above 40 years, the gpd to be expected in the future is scaled
gdp_future_bycontinents_byyear_high_lifeExp <- gapminder %>%
    mutate(gdp_futureExpectation = ifelse(lifeExp > 40, gdpPercap * 1.5, gdpPercap)) %>%
    group_by(continent, year) %>%
    summarize(mean_gdpPercap = mean(gdpPercap),
        mean_gdpPercap_expected = mean(gdp_futureExpectation))
```

Output

`summarise()` has grouped output by 'continent'. You can override using the `.groups` argument.

Combining dplyr and ggplot2

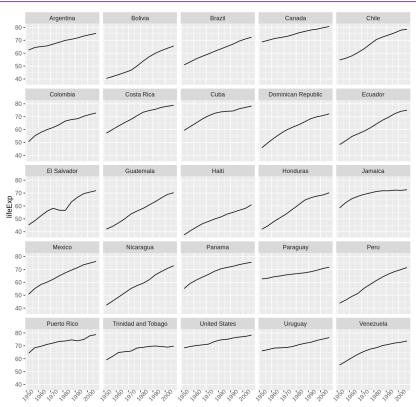
First install and load ggplot2:

```
R
install.packages('ggplot2')
```

R
library("ggplot2")

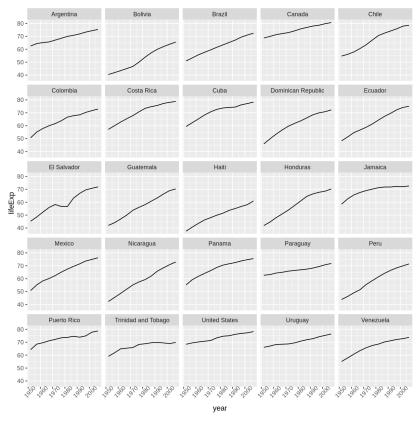
In the plotting lesson we looked at how to make a multi-panel figure by adding a layer of facet panels using ggplot2. Here is the code we used (with some extra comments):

```
# Filter countries located in the Americas
americas <- gapminder[gapminder$continent == "Americas", ]
# Make the plot
ggplot(data = americas, mapping = aes(x = year, y = lifeExp)) +
geom_line() +
facet_wrap( ~ country) +
theme(axis.text.x = element_text(angle = 45))
```



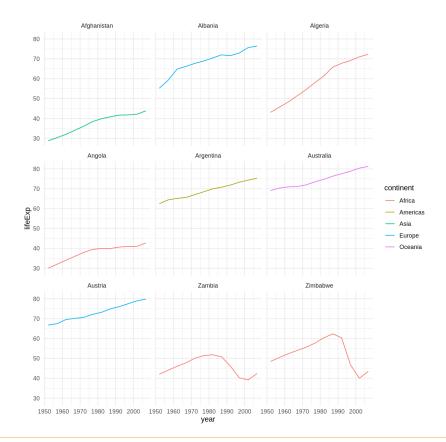
This code makes the right plot but it also creates an intermediate variable (americas) that we might not have any other uses for. Just as we used %>% to pipe data along a chain of dplyr functions we can use it to pass data to ggplot(). Because %>% replaces the first argument in a function we don't need to specify the data = argument in the ggplot() function. By combining dplyr and ggplot2 functions we can make the same figure without creating any new variables or modifying the

```
gapminder %>%
    # Filter countries located in the Americas
    filter(continent == "Americas") %>%
    # Make the plot
    ggplot(mapping = aes(x = year, y = lifeExp)) +
    geom_line() +
    facet_wrap( ~ country) +
    theme(axis.text.x = element_text(angle = 45))
```



More examples of using the function mutate() and the ggplot2 package.

```
gapminder %>%
    # extract first letter of country name into new column
    mutate(startsWith = substr(country, 1, 1)) %>%
    # only keep countries starting with A or Z
    filter(startsWith %in% c("A", "Z")) %>%
    # plot lifeExp into facets
    ggplot(aes(x = year, y = lifeExp, colour = continent)) +
    geom_line() +
    facet_wrap(vars(country)) +
    theme_minimal()
```



Advanced Challenge

Calculate the average life expectancy in 2002 of 2 randomly selected countries for each continent. Then arrange the continent names in reverse order. **Hint:** Use the dplyr functions arrange() and sample_n(), they have similar syntax to other dplyr functions.

Solution to Advanced Challenge

R

lifeExp_2countries_bycontinents <- gapminder %>%
 filter(year==2002) %>%
 group_by(continent) %>%
 sample_n(2) %>%
 summarize(mean_lifeExp=mean(lifeExp)) %>%
 arrange(desc(mean_lifeExp))

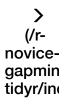
Other great resources

- R for Data Science (http://r4ds.had.co.nz/)
- Data Wrangling Cheat sheet (https://www.rstudio.com/wp-content/uploads/2015/02/data-wrangling-cheatsheet.pdf)
- Introduction to dplyr (https://dplyr.tidyverse.org/)
- Data wrangling with R and RStudio (https://www.rstudio.com/resources/webinars/data-wrangling-with-r-and-rstudio/)

Key Points

- Use the dplyr package to manipulate data frames.
- Use select() to choose variables from a data frame.
- Use filter() to choose data based on values.
- Use group_by() and summarize() to work with subsets of data.
- Use mutate() to create new variables.





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