# Environmental Health Big Data Analysis – R4ds (2) Data Wrangling and Programing

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## What we will learn

- 1. Data transformation
- 2. Data visualization
- 3. Data wrangling
- 4. Functional programming with R
- 5. Something useful

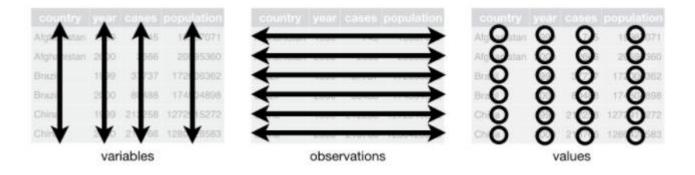


# **Data Wrangling - tidyr**

"Tidy datasets are all alike, but every messy dataset is messy in its own way." — Hadley Wickham

### Tidy data:

- Every column is variable
- Every row is an observation.
- Every cell is a single value.





# Why tidyr?

- If you have a consistent data structure, it's easier to learn the tools that work with it.
- Most built-in R functions work with vectors of values. That makes transforming tidy data feel particularly natural.
- dplyr, ggplot2, and all the other packages in the tidyverse are designed to work with tidy data.



# **Example Data**

### Representation of the same data in multiple ways

```
table1
#> # A tibble: 6 x 4
                 year cases population
     country
     <chr>>
                 <int> <int>
                                   <int>
#> 1 Afghanistan 1999
                                19987071
#> 2 Afghanistan
                  2000
                                20595360
                         2666
#> 3 Brazil
                  1999
                               172006362
#> 4 Brazil
                  2000
                               174504898
#> 5 China
                  1999 212258 1272915272
#> 6 China
                  2000 213766 1280428583
```

```
table2
#> # A tibble: 12 x 4
    country
                 year type
                                     count
                <int> <chr>
                                    <int>
    <chr>>
#> 1 Afghanistan 1999 cases
                                      745
#> 2 Afghanistan 1999 population 19987071
#> 3 Afghanistan 2000 cases
                                     2666
#> 4 Afghanistan 2000 population 20595360
#> 5 Brazil
                                     37737
                 1999 cases
#> 6 Brazil 1999 population 172006362
#> # ... with 6 more rows
```

```
table3
#> # A tibble: 6 x 3
    country
                 vear rate
#> * <chr>
                <int> <chr>
#> 1 Afghanistan 1999 745/19987071
#> 2 Afghanistan 2000 2666/20595360
#> 3 Brazil
                 1999 37737/172006362
#> 4 Brazil
                 2000 80488/174504898
#> 5 China
                 1999 212258/1272915272
#> 6 China
                 2000 213766/1280428583
```

```
table4b # population
#> # A tibble: 3 x 3
     country
                      `1999`
                                 `2000`
#> * <chr>
                       <int>
                                  <int>
#> 1 Afahanistan
                   19987071
                               20595360
#> 2 Brazil
                  172006362
                              174504898
#> 3 China
                 1272915272 1280428583
```



# **Pivoting**

 One variable might be spread across multiple columns. (Use Pivot\_longer())

 One observation might be scattered across multiple rows. (Use Pivot\_wider())



# Apply Pivot\_longer() to table4a

```
table4a %>%

pivot_longer(c(`1999`, `2000`), names_to = "year", values_to = "cases")

#> # A tibble: 6 x 3

#> country year cases

#> <chr> <chr> <int>
#> 1 Afghanistan 1999 745

#> 2 Afghanistan 2000 2666

#> 3 Brazil 1999 37737

#> 4 Brazil 2000 80488

#> 5 China 1999 212258

#> 6 China 2000 213766
```

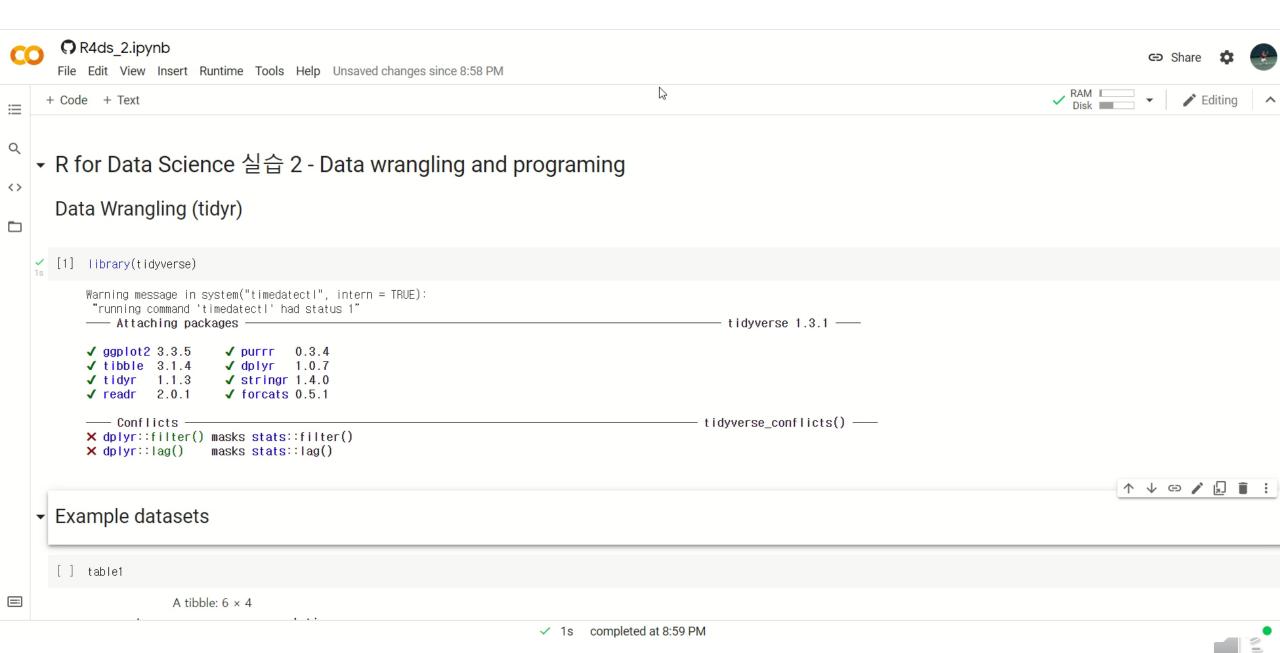
country	year	cases		country	1999	2000
Afghanistan	1999	745	<b>←</b>	Algivariatan	745	2666
Afghanistan	2000	2666	$\leftarrow$	Brazil	37737	80488
Brazil	1999	37737	←	China	212258	213766
Brazil	2000	80488	<b>←</b>		_	
China	1999	212258	<b>—</b>			
China	2000	213766	$\leftarrow$		table4	



# Apply Pivot\_longer() and combine data

```
tidy4a <- table4a %>%
 pivot longer(c(1999), 2000), names to = "year", values to = "cases")
tidy4b <- table4b %>%
 pivot longer(c(`1999`, `2000`), names to = "year", values to = "population")
left join(tidy4a, tidy4b)
#> Joining, by = c("country", "year")
#> # A tibble: 6 x 4
#> country year cases population
#> <chr> <int> <int>
#> 1 Afghanistan 1999 745 19987071
#> 2 Afghanistan 2000 2666 20595360
#> 3 Brazil
            1999 37737 172006362
#> 4 Brazil
            2000 80488 174504898
#> 5 China
           1999 212258 1272915272
           2000 213766 1280428583
#> 6 China
```





# **Pivoting**

- One variable might be spread across multiple columns. (Use Pivot\_longer())
- One observation might be scattered across multiple rows. (Use Pivot\_wider())

```
#> # A tibble: 12 x 4

#> country year type count

#> <chr> <int> <chr> <int> <int>
#> 1 Afghanistan 1999 cases 745

#> 2 Afghanistan 1999 population 19987071

#> 3 Afghanistan 2000 cases 2666

#> 4 Afghanistan 2000 population 20595360

#> 5 Brazil 1999 cases 37737

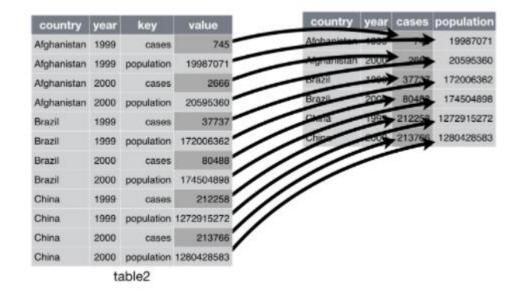
#> 6 Brazil 1999 population 172006362

#> # ... with 6 more rows
```

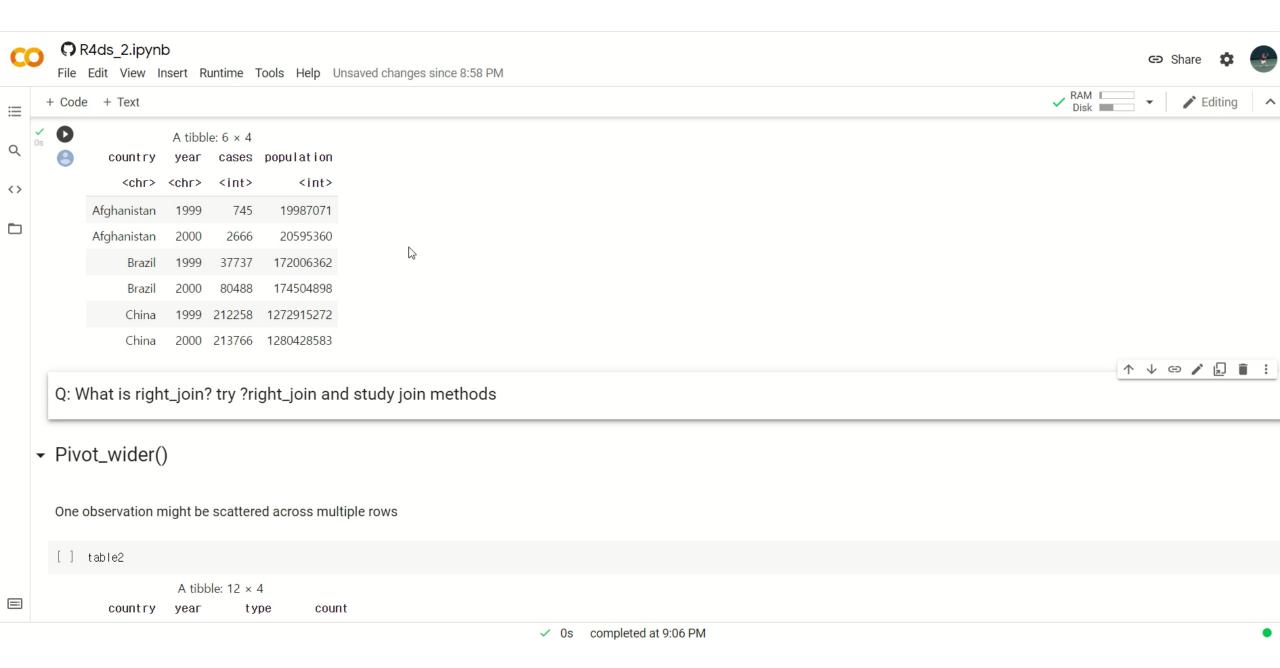


# Apply Pivot\_wider()

```
table2 %>%
  pivot wider(names from = type, values from = count)
#> # A tibble: 6 x 4
   country year cases population
#> <chr>
            <int> <int> <int>
#> 1 Afghanistan 1999 745 19987071
#> 2 Afghanistan 2000 2666 20595360
            1999 37737 172006362
#> 3 Brazil
#> 4 Brazil
            2000 80488 174504898
            1999 212258 1272915272
#> 5 China
#> 6 China
            2000 213766 1280428583
```







# Separating and uniting

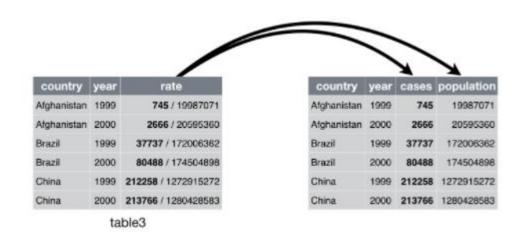
- One column contains two variables (Use separate())

Single variable is spread across multiple columns.
 (Use unite())



# Apply separate()

```
table3 %>%
separate(rate, into = c("cases", "population"))
#> # A tibble: 6 x 4
   country year cases population
#> <chr>
            <int> <chr> <chr>
#> 1 Afghanistan 1999 745 19987071
#> 2 Afghanistan 2000 2666 20595360
#> 3 Brazil
            1999 37737 172006362
#> 4 Brazil
            2000 80488 174504898
#> 5 China
            1999 212258 1272915272
#> 6 China
             2000 213766 1280428583
```





# Separating and uniting

- One column contains two variables (Use separate())
- Single variable is spread across multiple columns.
   (Use unite())

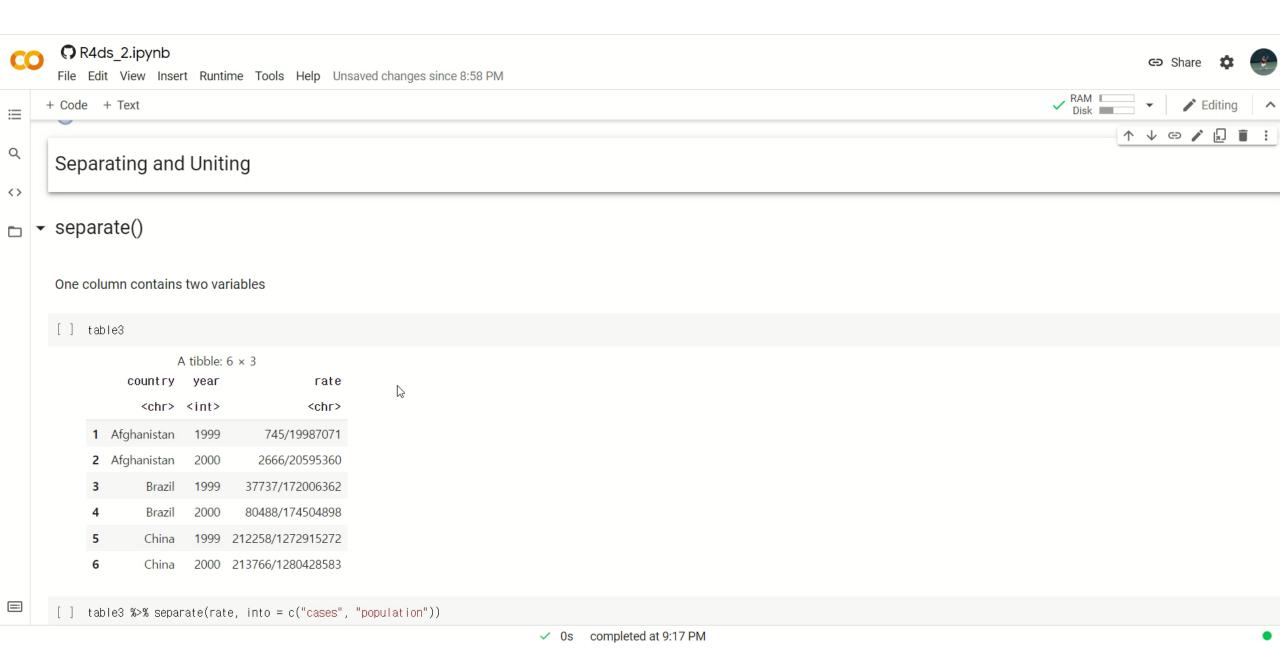
	country	century	year	rate
	<chr></chr>	<chr></chr>	<chr></chr>	<chr></chr>
1	Afghanistan	19	99	745/19987071
2	Afghanistan	20	00	2666/20595360
3	Brazil	19	99	37737/172006362
4	Brazil	20	00	80488/174504898
5	China	19	99	212258/1272915272
6	China	20	00	213766/1280428583



# Apply unite()

```
table5 %>%
unite(new, century, year, sep = "")
#> # A tibble: 6 x 3
   country
           new rate
#> <chr>
            <chr> <chr>
#> 1 Afghanistan 1999 745/19987071
#> 2 Afghanistan 2000 2666/20595360
            1999 37737/172006362
#> 3 Brazil
            2000 80488/174504898
#> 4 Brazil
#> 5 China
             1999 212258/1272915272
#> 6 China
             2000 213766/1280428583
```





# **Functional Programming**

- For loops are quite verbose, and require quite a bit of bookkeeping code that is duplicated for every for loop.
- Functional programming offers tools to extract out this duplicated code, so each common for loop pattern gets its own function.



# For loops

### Random data

```
df <- tibble(
    a = rnorm(10),
    b = rnorm(10),
    c = rnorm(10),
    d = rnorm(10)
)</pre>
```

## With for loops

```
output <- vector("double", ncol(df)) # 1. output
for (i in seq_along(df)) { # 2. sequence
  output[[i]] <- median(df[[i]]) # 3. body
}
output
#> [1] -0.24576245 -0.28730721 -0.05669771 0.14426335
```

### To calculate median

```
median(df$a)
#> [1] -0.2457625
median(df$b)
#> [1] -0.2873072
median(df$c)
#> [1] -0.05669771
median(df$d)
#> [1] 0.1442633
```



### Three components

The output: output <- vector("double", length(x)).
Allocate sufficient space for the output.
(If you grow the for loop at each iteration using c() (for example), your for loop will be very slow)

The sequence: i in seq\_along(df). what to loop over: each run of the for loop will assign i to a different value from seq\_along(df)

The body: output[[i]] <- median(df[[i]]). This is the code that does the work.

```
output <- vector("double", ncol(df)) # 1. output
for (i in seq_along(df)) { # 2. sequence
  output[[i]] <- median(df[[i]]) # 3. body
}
output
#> [1] -0.24576245 -0.28730721 -0.05669771 0.14426335
```



### For loops vs. functionals

#### Possible to wrap up for loops in a function

```
col_mean <- function(df) {
    output <- vector("double", ncol(df))
    for (i in seq_along(df)) {
        output[[i]] <- mean(df[[i]])
    }
    output
}</pre>
```

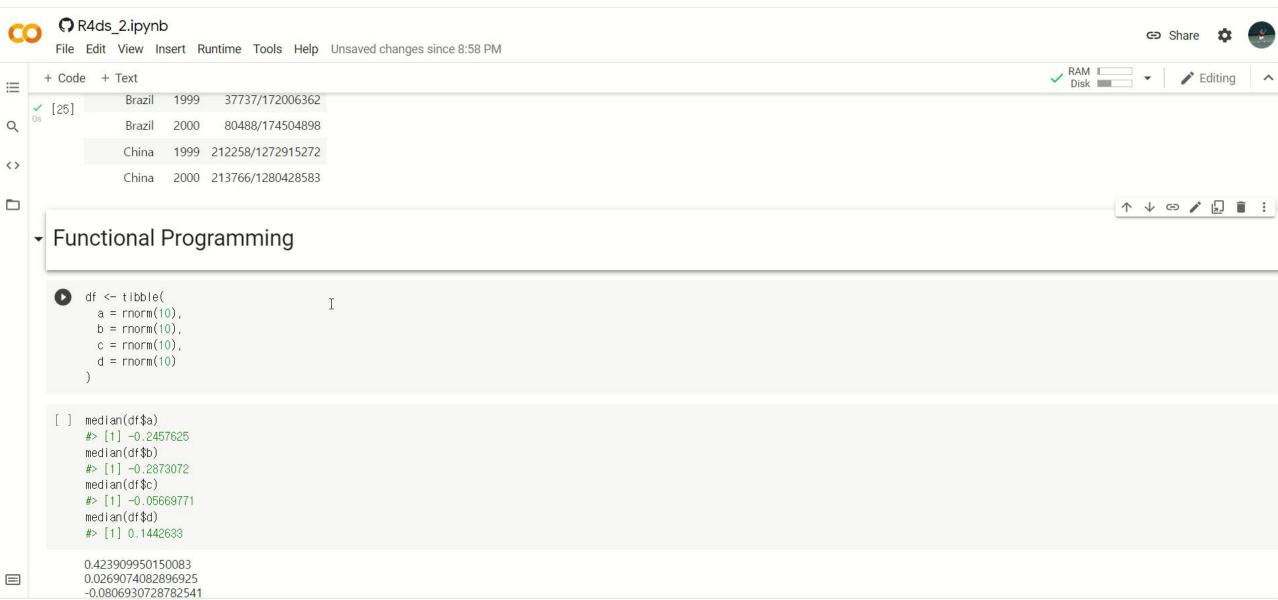
```
col_median <- function(df) {
    output <- vector("double", ncol(df))
    for (i in seq_along(df)) {
        output[[i]] <- median(df[[i]])
    }
    output
}</pre>
```

```
col_sd <- function(df) {
    output <- vector("double", ncol(df))
    for (i in seq_along(df)) {
        output[[i]] <- sd(df[[i]])
    }
    output
}</pre>
```

```
col_summary <- function(df, fun) {
  out <- vector("double", length(df))
  for (i in seq_along(df)) {
    out[i] <- fun(df[[i]])
  }
  out
}

col_summary(df, median)
#> [1] -0.51850298  0.02779864  0.17295591 -0.61163819
col_summary(df, mean)
#> [1] -0.3260369  0.1356639  0.4291403 -0.2498034
```





## The map function (purrr)

the purrr package provides a family of functions for looping patterns over a vector

- map() makes a list
- map\_lgl() makes a logical vector
- map\_int() makes an integer vector
- map\_dbl() makes a double vector
- map\_chr() makes a character vector

Alternatives: apply, lapply, etc



## The map function (purrr)

```
df %>% map_dbl(mean)

#> a b c d

#> -0.3260369 0.1356639 0.4291403 -0.2498034

df %>% map_dbl(median)

#> a b c d

#> -0.51850298 0.02779864 0.17295591 -0.61163819

df %>% map_dbl(sd)

#> a b c d

#> 0.9214834 0.4848945 0.9816016 1.1563324
```

### You can define a function in a map function

```
models <- mtcars %>%
split(.$cyl) %>%
map(function(df) lm(mpg ~ wt, data = df))
```

```
models <- mtcars %>%
split(.$cyl) %>%
map(~lm(mpg ~ wt, data = .))
```



#### To extract a component

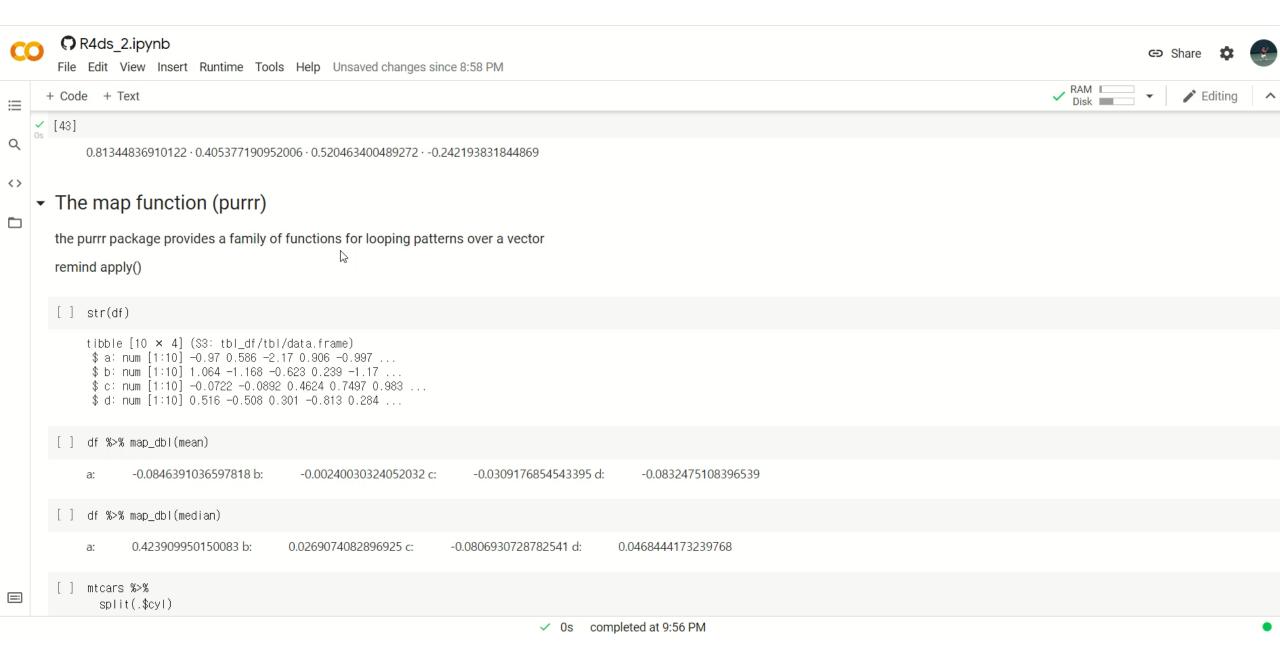
```
models %>%
   map(summary) %>%
   map_dbl(~.$r.squared)
#> 4 6 8
#> 0.5086326 0.4645102 0.4229655
```

```
models %>%
   map(summary) %>%
   map_dbl("r.squared")
#> 4 6 8
#> 0.5086326 0.4645102 0.4229655
```

### You can also use an integer to select elements by position

```
x <- list(list(1, 2, 3), list(4, 5, 6), list(7, 8, 9))
x %>% map_dbl(2)
#> [1] 2 5 8
```





# Something useful

- R markdown
- Boxplot vs Violinplot
- Do not use pie chart?
- Effective 2D visualization with t-SNE and UMAP



#### R for Data Science

Search

#### Table of contents

Welcome

1 Introduction

Explore

- 2 Introduction
- 3 Data visualisation
- 4 Workflow: basics
- 5 Data transformation
- 6 Workflow: scripts
- 7 Exploratory Data Analysis
- 8 Workflow: projects

Wrangle

- 9 Introduction
- 10 Tibbles
- 11 Data import
- 12 Tidy data
- 13 Relational data
- 14 Strings
- 15 Factors
- 16 Dates and times

#### 27 R Markdown

#### 27.1 Introduction &

R Markdown provides an unified authoring framework for data science, combining your code, its results, and your prose commentary. R Markdown documents are fully reproducible and support dozens of output formats, like PDFs, Word files, slideshows, and more.

R Markdown files are designed to be used in three ways:

- 1. For communicating to decision makers, who want to focus on the conclusions, not the code behind the analysis.
- 2. For collaborating with other data scientists (including future you!), who are interested in both your conclusions, and how you reached them (i.e. the code).
- 3. As an environment in which to *do* data science, as a modern day lab notebook where you can capture not only what you did, but also what you were thinking.

R Markdown integrates a number of R packages and external tools. This means that help is, by-and-large, not available through ? . Instead, as you work through this chapter, and use R Markdown in the future, keep these resources close to hand:

■ R Markdown Cheat Sheet: Help > Cheatsheets > R Markdown Cheat Sheet,

#### On this page

27 R Markdown

27.1 Introduction

27.2 R Markdown basics

27.3 Text formatting with Markdown

27.4 Code chunks

27.5 Troubleshooting

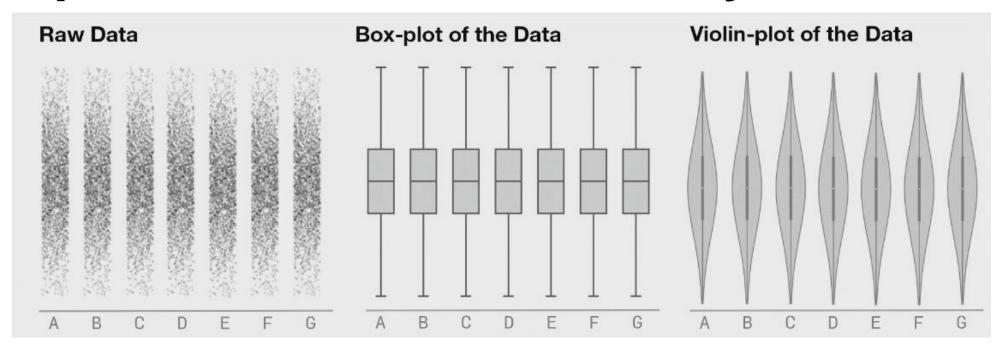
27.6 YAML header

27.7 Learning more

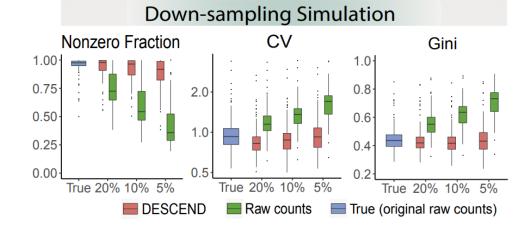
View source 😱

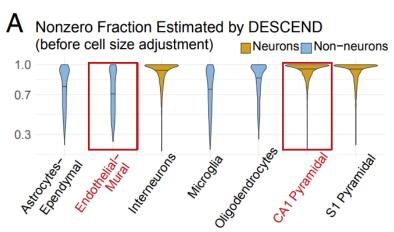


# Violinplot shows the data density

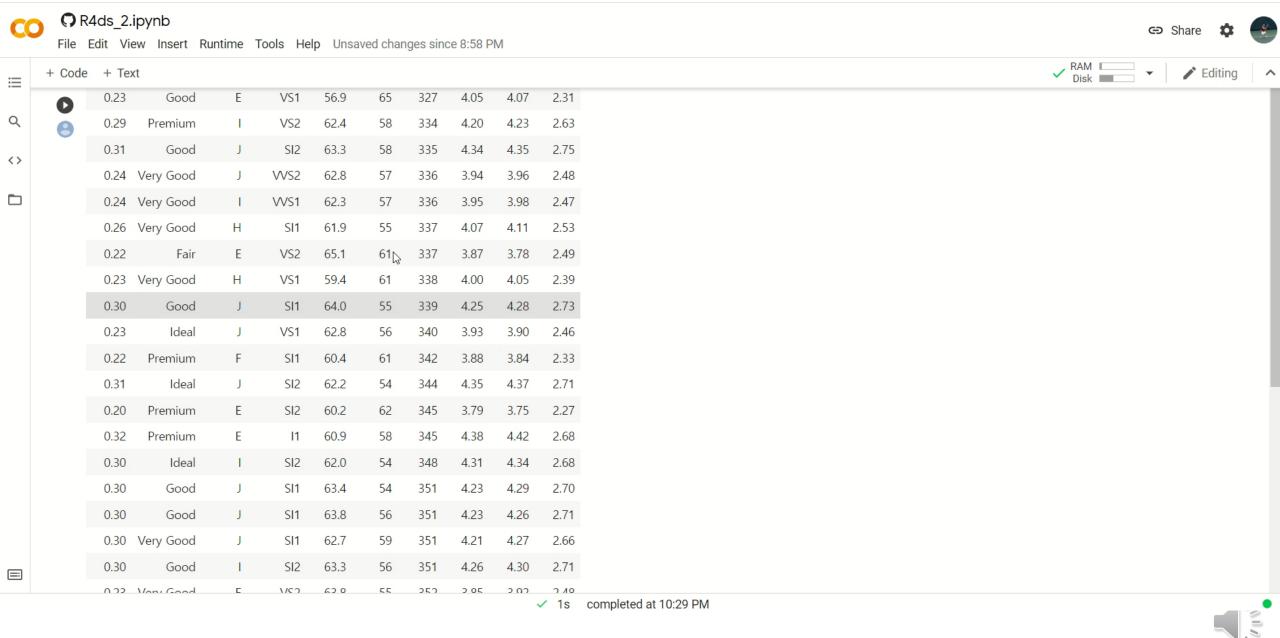




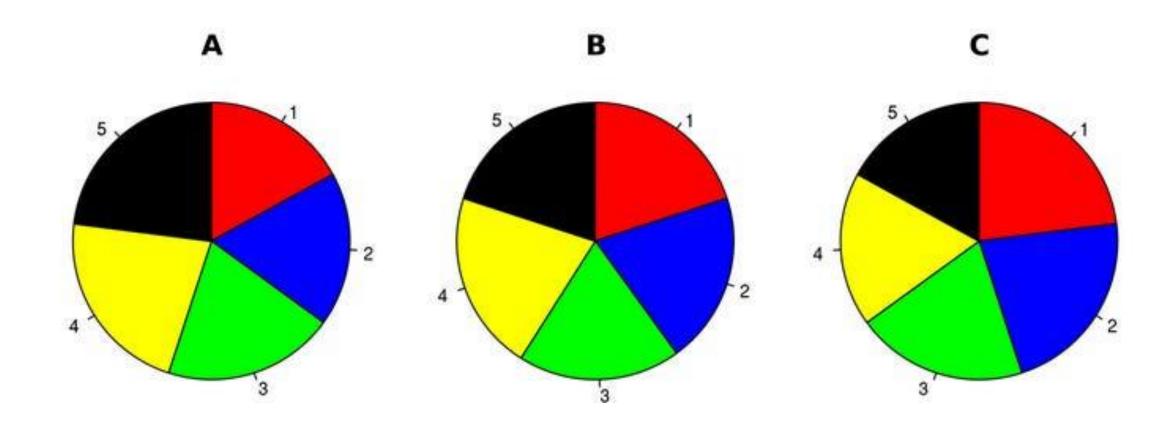




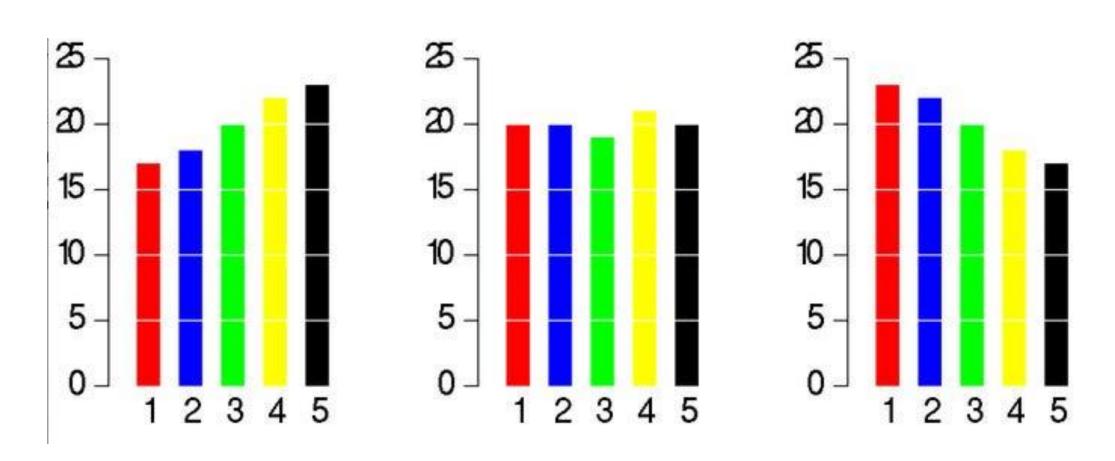




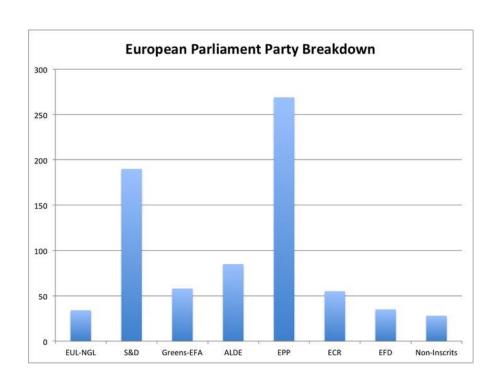
# Do not use pie chart?

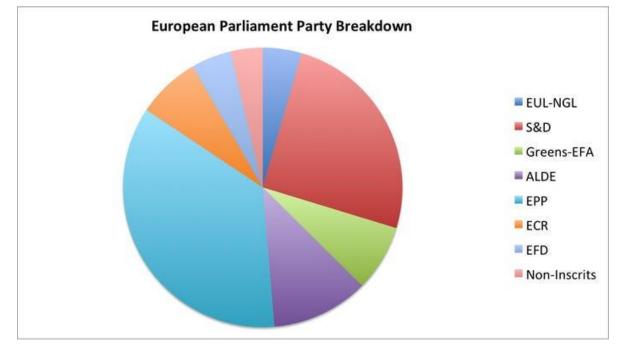


## Same Information from Bar Plot

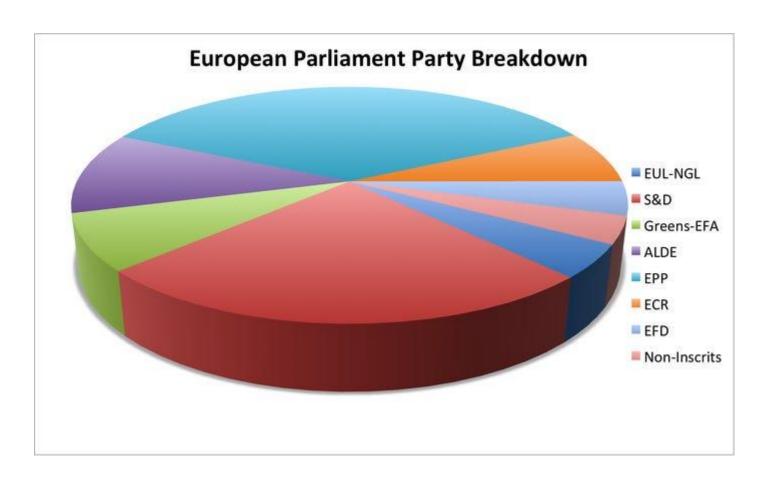


## Human Eyes are not Good at Evaluating Sizes

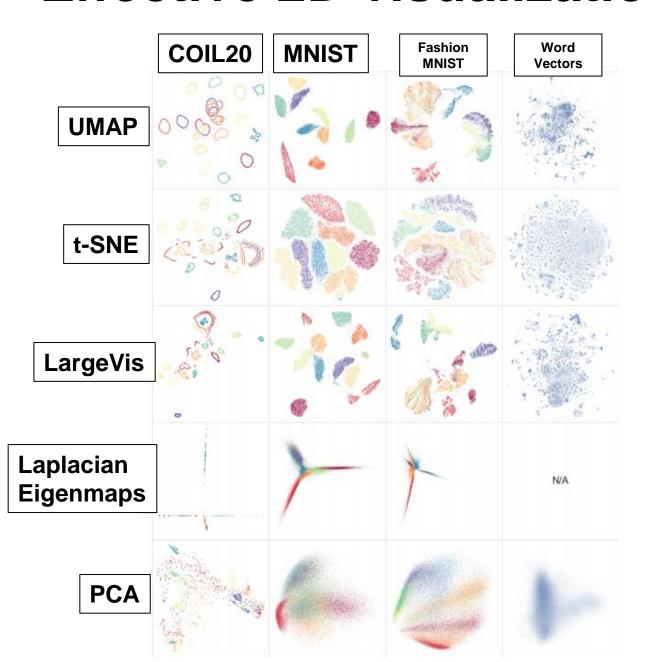




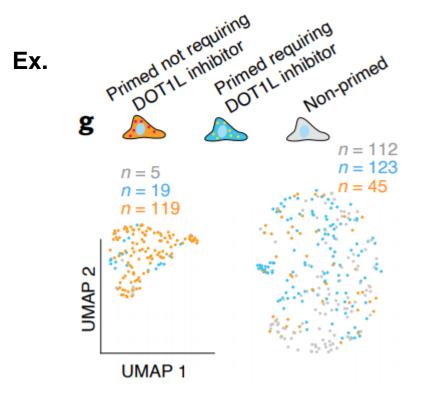
## 3d Plot even Worse



## Effective 2D visualization with t-SNE and UMAP



When visualizing multidimensional data in 2D, t-SNE and UMAP works well





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#### Visualizing Data using t-SNE

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Editor: Yoshua Bengio

#### Abstract

We present a new technique called "t-SNE" that visualizes high-dimensional data by giving each datapoint a location in a two or three-dimensional map. The technique is a variation of Stochastic Neighbor Embedding (Hinton and Roweis, 2002) that is much easier to optimize, and produces significantly better visualizations by reducing the tendency to crowd points together in the center of the map. t-SNE is better than existing techniques at creating a single map that reveals structure at many different scales. This is particularly important for high-dimensional data that lie on several different, but related, low-dimensional manifolds, such as images of objects from multiple classes seen from multiple viewpoints. For visualizing the structure of very large data sets, we show how t-SNE can use random walks on neighborhood graphs to allow the implicit structure of all of the data to influence the way in which a subset of the data is displayed. We illustrate the performance of t-SNE on a wide variety of data sets and compare it with many other non-parametric visualization techniques, including Sammon mapping, Isomap, and Locally Linear Embedding. The visualizations produced by t-SNE are significantly better than those produced by the other techniques on almost all of the data sets.

# Thank you! ©

