# Getting / cleaning data 2

# Tidy data

### Tidy data

All of the material in this section comes directly from Hadley Wickham's paper on tidy data. You will need to read this paper to prepare for the quiz on this section.

### Characteristics of tidy data

### Characteristics of tidy data are:

- 1. Each variable forms a column.
- 2. Each observation forms a row.
- 3. Each type of observational unit forms a table.

Getting your data into a "tidy" format makes it easier to model and plot. By taking the time to tidy your data at the start of an analysis, you will save yourself time, and make it easier to plan out, later steps.

Here are five common problems that Hadley Wickham has identified that keep data from being tidy:

- 1. Column headers are values, not variable names.
- 2. Multiple variables are stored in one column.
- 3. Variables are stored in both rows and columns.
- 4. Multiple types of observational units are stored in the same table.
- 5. A single observational unit is stored in multiple tables.

In the following slides, I'll give examples of each of these problems.

### (1.) Column headers are values, not variable names.

religion	<\$10k	10-20k	\$20-30k	\$30-40k	\$40-50k	\$50-75k
Agnostic	27	34	60	81	76	137
Atheist	12	27	37	52	35	70
Buddhist	27	21	30	34	33	58
Catholic	418	617	732	670	638	1116
Don't know/refused	15	14	15	11	10	38
Evangelical Prot	575	869	1064	982	881	1486
Hindu	1	9	7	9	11	34
Historically Black Prot	228	244	236	238	197	223
Jehovah's Witness	20	27	24	24	21	30
Jewish	19	19	25	25	30	98

### Solution:

religion	income	freq
Agnostic	<\$10k	27
Agnostic	\$10-20k	34
Agnostic	\$20-30k	60
Agnostic	\$30-40k	81
Agnostic	\$40-50k	76
Agnostic	\$50-75k	137
Agnostic	\$75-100k	122
Agnostic	\$100-150k	109
Agnostic	>150k	84
Agnostic	Don't know/refused	96

(2.) Multiple variables are stored in one column.

		column	
country	year	corumn	cases
AD	2000	m014	0
AD	2000	m1524	0
AD	2000	m2534	1
$^{\mathrm{AD}}$	2000	m3544	0
$^{\mathrm{AD}}$	2000	m4554	0
$^{\mathrm{AD}}$	2000	m5564	0
$^{\mathrm{AD}}$	2000	m65	0
$\mathbf{AE}$	2000	m014	2
$\mathbf{AE}$	2000	m1524	4
$\mathbf{AE}$	2000	m2534	4
$\mathbf{AE}$	2000	m3544	6
$\mathbf{AE}$	2000	m4554	5
$\mathbf{AE}$	2000	m5564	12
$\mathbf{AE}$	2000	m65	10
$\mathbf{AE}$	2000	f014	3

### Solution:

country	year	sex	age	cases
AD	2000	$\mathbf{m}$	0-14	0
AD	2000	m	15-24	0
AD	2000	m	25 - 34	1
AD	2000	$\mathbf{m}$	35-44	0
AD	2000	m	45-54	0
AD	2000	m	55-64	0
AD	2000	$\mathbf{m}$	65 +	0
$\mathbf{AE}$	2000	m	0-14	2
$\mathbf{AE}$	2000	m	15-24	4
$\mathbf{AE}$	2000	$\mathbf{m}$	25 - 34	4
$\mathbf{AE}$	2000	$\mathbf{m}$	35-44	6
$\mathbf{AE}$	2000	m	45-54	5
$\mathbf{AE}$	2000	m	55-64	12
$\mathbf{AE}$	2000	$\mathbf{m}$	65 +	10
$\mathbf{AE}$	2000	$\mathbf{f}$	0-14	3

(3.) Variables are stored in both rows and columns.

id	year	month	element	d1	d2	d3	d4	d5	d6	d7	d8
MX17004	2010	1	tmax	_	_	_	_	_	_	_	_
MX17004	2010	1	$_{ m tmin}$	_	_	_	_	_	_	_	_
MX17004	2010	2	tmax	_	27.3	24.1	_	_	_	_	_
MX17004	2010	2	$_{ m tmin}$	_	14.4	14.4	_	_	_	_	_
MX17004	2010	3	tmax	_	_	_	_	32.1	_	_	_
MX17004	2010	3	$_{ m tmin}$	_	_	_	_	14.2	_	_	_
MX17004	2010	4	$_{\text{tmax}}$	_	_	_	_	_	_	_	_
MX17004	2010	4	$_{ m tmin}$		_	_	_	_	_	_	_
MX17004	2010	5	tmax	_	_	_	_	_	_	_	_
MX17004	2010	5	$_{ m tmin}$	_	_	_	_	_	_	_	_

### Solution:

id	date	element	value	id	date	$\operatorname{tmax}$	$_{ m tmin}$
MX17004	2010-01-30	tmax	27.8	MX17004	2010-01-30	27.8	14.5
MX17004	2010-01-30	$_{ m tmin}$	14.5	MX17004	2010-02-02	27.3	14.4
MX17004	2010-02-02	tmax	27.3	MX17004	2010-02-03	24.1	14.4
MX17004	2010-02-02	$_{ m tmin}$	14.4	MX17004	2010-02-11	29.7	13.4
MX17004	2010-02-03	tmax	24.1	MX17004	2010-02-23	29.9	10.7
MX17004	2010-02-03	$_{ m tmin}$	14.4	MX17004	2010-03-05	32.1	14.2
MX17004	2010-02-11	tmax	29.7	MX17004	2010-03-10	34.5	16.8
MX17004	2010-02-11	$_{ m tmin}$	13.4	MX17004	2010-03-16	31.1	17.6
MX17004	2010-02-23	tmax	29.9	MX17004	2010-04-27	36.3	16.7
MX17004	2010-02-23	$_{ m tmin}$	10.7	MX17004	2010-05-27	33.2	18.2

(4.) Multiple types of observational units are stored in the same table.

year	artist	time	track	date	week	rank
2000	2 Pac	4:22	Baby Don't Cry	2000-02-26	1	87
2000	2 Pac	4:22	Baby Don't Cry	2000-03-04	2	82
2000	2 Pac	4:22	Baby Don't Cry	2000-03-11	3	72
2000	2 Pac	4:22	Baby Don't Cry	2000-03-18	4	77
2000	2 Pac	4:22	Baby Don't Cry	2000-03-25	5	87
2000	2 Pac	4:22	Baby Don't Cry	2000-04-01	6	94
2000	2 Pac	4:22	Baby Don't Cry	2000-04-08	7	99
2000	2Ge+her	3:15	The Hardest Part Of	2000-09-02	1	91
2000	2Ge+her	3:15	The Hardest Part Of	2000-09-09	2	87
2000	2Ge+her	3:15	The Hardest Part Of	2000-09-16	3	92
2000	3 Doors Down	3:53	Kryptonite	2000-04-08	1	81
2000	3 Doors Down	3:53	Kryptonite	2000-04-15	2	70
2000	3 Doors Down	3:53	Kryptonite	2000-04-22	3	68
2000	3 Doors Down	3:53	Kryptonite	2000-04-29	4	67
2000	3 Doors Down	3:53	Kryptonite	2000-05-06	5	66

### Solution:

id	artist	track	time	id	date	rank
1	2 Pac	Baby Don't Cry	4:22	1	2000-02-26	87
2	2Ge+her	The Hardest Part Of	3:15	1	2000-03-04	82
3	3 Doors Down	Kryptonite	3:53	1	2000-03-11	72
4	3 Doors Down	Loser	4:24	1	2000-03-18	77
5	504 Boyz	Wobble Wobble	3:35	1	2000-03-25	87
6	98^0	Give Me Just One Nig	3:24	1	2000-04-01	94
7	A*Teens	Dancing Queen	3:44	1	2000-04-08	99
8	Aaliyah	I Don't Wanna	4:15	2	2000-09-02	91
9	Aaliyah	Try Again	4:03	2	2000-09-09	87
10	Adams, Yolanda	Open My Heart	5:30	2	2000-09-16	92
11	Adkins, Trace	More	3:05	3	2000-04-08	81
12	Aguilera, Christina	Come On Over Baby	3:38	3	2000-04-15	70
13	Aguilera, Christina	I Turn To You	4:00	3	2000-04-22	68
14	Aguilera, Christina	What A Girl Wants	3:18	3	2000-04-29	67
15	Alice Deejay	Better Off Alone	6:50	3	2000-05-06	66

(5.) A single observational unit is stored in multiple tables.

Example: exposure and outcome data stored in different files:

- File 1: Daily mortality counts
- File 2: Daily air pollution measurements

### In-course exercise

We'll take a break now to do the In-Course Exercise (Section 6.4.1).

# Joining datasets

### Joining datasets

So far, you have only worked with a single data source at a time. When you work on your own projects, however, you typically will need to merge together two or more datasets to create the a data frame to answer your research question.

For example, for air pollution epidemiology, you will often have to join several datasets:

- Health outcome data (e.g., number of deaths per day)
- Air pollution concentrations
- Weather measurements (since weather can be a confounder)
- Demographic data

The dplyr package has a family of different functions to join two dataframes together, the \*\_join family of functions. All combine two dataframes, which I'll call x and y here.

#### The functions include:

- inner\_join(x, y): Keep only rows where there are observations in both x and y.
- left\_join(x, y): Keep all rows from x, whether they have a match in y or not.
- right\_join(x, y): Keep all rows from y, whether they have a match in x or not.
- full\_join(x, y): Keep all rows from both x and y, whether they
  have a match in the other dataset or not.

In the examples, I'll use two datasets, x and y. Both datasets include the column course. The other column in x is grade, while the other column in y is day. Observations exist for courses x and y in both datasets, but for w and z in only one dataset.

## 2 x Mon / Fri

## 3

Here is what these two example datasets look like:

Tue

```
х
##
    course grade
## 1
              90
         Х
         y 82
## 2
## 3
         z 78
У
##
                 day
    course
## 1
               Tues
```

With inner\_join, you'll only get the observations that show up in both datasets. That means you'll lose data on z (only in the first dataset) and w (only in the second dataset).

```
inner_join(x, y)

## Joining, by = "course"

## course grade day
## 1 x 90 Mon / Fri
## 2 y 82 Tue
```

## 3

Z

78

With left\_join, you'll keep everything in x (the "left" dataset), but not keep things in y that don't match something in x. That means that, here, you'll lose w:

```
left_join(x, y)

## Joining, by = "course"

## course grade day
## 1 x 90 Mon / Fri
## 2 y 82 Tue
```

<NA>

## 3

У

right\_join is the opposite— you keep all observations in the "right" dataframe, but only matching ones in the "left" dataframe:

```
right_join(x, y)

## Joining, by = "course"

## course grade day
## 1 w NA Tues
## 2 x 90 Mon / Fri
```

82 Tue

full\_join keeps everything from both datasets:

For some more complex examples of using join, I'll use these example datasets:

```
## # A tibble: 4 x 3
## course grade student
## <chr> <dbl> <chr>
         92. a
## 1 x
## 2 x
        90. b
## 3 v 82. a
         78. b
## 4 z
## # A tibble: 4 x 3
## class day student
## <chr> <chr> <chr>
## 1 w Tues
                 а
## 2 x Mon / Fri a
## 3 x Mon / Fri b
## 4 v
         Tue
                 a
```

If you have two datasets you want to join, but the column names for the joining column are different, you can use the by argument:

```
full_join(x, y, by = list(x = "course", y = "class"))
```

```
## # A tibble: 7 x 5
## course grade student.x day student.y
## <chr> <dbl> <chr> <chr> <dbl> <chr> <chr> <dbl> chr> <chr> <chr> ## 1 x 92. a Mon / Fri a
## 2 x 92. a Mon / Fri b
## 3 x 90. b Mon / Fri a
## 4 x 90. b Mon / Fri a
## 5 y 82. a Tue a
## 6 z 78. b <NA> Tues a
```

#### A few things to note about this example:

- The joining column name for the "left" dataframe (x in this case) is used as the column name for the joined data
- student was a column name in both x and y. If we're not using it to join the data, the column names are changed in the joined data to student.x and student.y.
- Values are recycled for rows where there were multiple matches across the dataframe (e.g., rows for course "x")

Sometimes, you will want to join by more than one column. In this example data, it would make sense to join the data by matching both course and student. You can do this by using a vector of all columns to join on:

```
## # A tibble: 5 x 4
## course grade student day
## 
## 1 x 92. a Mon / Fri
## 2 x 90. b Mon / Fri
## 3 y 82. a Tue
## 4 z 78. b <NA>
## 5 w NA a Tues
```

### In-course exercise

We'll take a break now to do the In-Course Exercise (Section 6.4.2).

# **Gathering**

There are two functions from the tidyr package (another member of the tidyverse) that you can use to change between wide and long data: gather and spread.

Here is a description of these two functions:

- gather: Take several columns and gather them into two columns, one with the former column names, and one with the former cell values
- spread: Take two columns and spread them into multiple columns. Column names for the new columns will come from one of the two original columns, while cell values will come from the other of the original columns.

The following examples are from tidyr help files and show the effects of gathering and spreading a dataset.

Here is some wide data:

```
wide_stocks[1:3, ]
```

```
## time X Y Z
## 1 2009-01-01 1.26418181 -1.4808143 -2.97040854
## 2 2009-01-02 -0.06024213 -0.0122563 -1.95975963
## 3 2009-01-03 0.23477856 1.4407623 0.09714659
```

In the wide\_stocks dataset, there are separate columns for three different stocks (X, Y, and Z). Each cell gives the value for a certain stock on a certain day.

This data isn't "tidy", because the identify of the stock (X, Y, or Z) is a variable, and you'll probably want to include it as a variable in modeling.

```
wide_stocks[1:3, ]
```

```
## time X Y Z
## 1 2009-01-01 1.26418181 -1.4808143 -2.97040854
## 2 2009-01-02 -0.06024213 -0.0122563 -1.95975963
## 3 2009-01-03 0.23477856 1.4407623 0.09714659
```

If you want to convert the dataframe to have all stock values in a single column, you can use gather to convert wide data to long data:

In this "long" dataframe, there is now one column that gives the identify of the stock (stock) and another column that gives the price of that stock that day (price):

```
long_stocks[1:5, ]
```

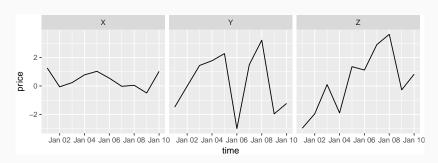
The format for a gather call is:

#### Three important notes:

- Everything is gathered into one of two columns— one column with the old column names, and one column with the old cell values
- With the key and value arguments, you are just providing column names for the two columns that everything's gathered into.
- If there is a column you don't want to gather (date in the example),
   use to exclude it in the gather call.

Notice how easy it is, now that the data is gathered, to use stock for aesthetics of faceting in a ggplot2 call:

```
ggplot(long_stocks, aes(x = time, y = price)) +
  geom_line() +
  facet_grid(. ~ stock)
```



If you have data in a "long" format and would like to spread it out, you can use spread to do that:

```
stocks <- spread(long_stocks, key = stock, value = price)
stocks[1:5, ]</pre>
```

```
## time X Y Z
## 1 2009-01-01 1.26418181 -1.4808143 -2.97040854
## 2 2009-01-02 -0.06024213 -0.0122563 -1.95975963
## 3 2009-01-03 0.23477856 1.4407623 0.09714659
## 4 2009-01-04 0.77918916 1.7725292 -1.89164625
## 5 2009-01-05 1.02970742 2.2753612 1.35743073
```

Notice that this reverses the action of gather.

"Spread" data is typically not tidy, so you often won't want to use spread when you are preparing data for analysis. However, spread can be very helpful in creating clean tables for final reports and presentations.

For example, if you wanted to create a table with means and standard deviations for each of the three stocks, you could use spread to rearrange the final summary to create an attractive table.

```
stock_summary <- long_stocks %>%
  group_by(stock) %>%
  summarize(N = n(), mean = mean(price), sd = sd(price))
stock_summary
```

```
## # A tibble: 3 x 4
## stock N mean sd
## <chr> <int> <dbl> <dbl> <dbl> <dbl> ## 1 X 10 0.434 0.580
## 2 Y 10 0.253 2.08
## 3 Z 10 0.284 2.13
```

Statistic	Χ	Υ	Z
Mean (Std.dev.)	0.43 (0.58)	0.25 (2.08)	0.28 (2.13)
N	10	10	10

## **Further reading**

Chapters 12 and 13 of "R for Data Science" are an excellent supplemental resource if you'd like extra practice on tidy data, gathering / spreading, and joining different datasets.

# More with dplyr

### dplyr

So far, you've used several dplyr functions:

- rename
- filter
- select
- mutate
- group\_by
- summarize

Some other useful dplyr functions to add to your toolbox are:

- separate and unite
- mutate and other dplyr functions with group\_by
- anti\_join and semi\_join

Sometimes, you want to take one column and split it into two columns. For example, you may have information for two variables in one column:

ebola

```
## # A tibble: 4 x 1
## ebola_key
## <chr>
## 1 Liberia_Cases
## 2 Liberia_Deaths
## 3 Spain_Cases
## 4 Spain Deaths
```

If you have a consistent "split" character, you can use the separate function to split one column into two:

```
ebola %>%
 separate(col = ebola_key, into = c("country", "outcome"),
          sep = ""
## # A tibble: 4 x 2
## country outcome
## <chr> <chr>
## 1 Liberia Cases
## 2 Liberia Deaths
## 3 Spain Cases
## 4 Spain Deaths
```

Here is the generic code for separate:

The default is to drop the original column and only keep the columns into which it was split. However, you can use the argument remove = FALSE to keep the first column, as well:

```
ebola %>%
 separate(col = ebola_key, into = c("country", "outcome"),
          sep = " ", remove = FALSE)
## # A tibble: 4 \times 3
## ebola_key country outcome
## <chr>
            <chr> <chr>
## 1 Liberia Cases Liberia Cases
## 2 Liberia Deaths Liberia Deaths
## 3 Spain_Cases
                   Spain Cases
## 4 Spain Deaths
                   Spain Deaths
```

You can use the fill argument (fill = "right" or fill = "left") to control what happens when there are some observations that do not have the split character.

For example, say your original column looked like this:

```
## # A tibble: 4 x 1
## ebola_key
## <chr>
## 1 Liberia_Cases
## 2 Liberia
## 3 Spain_Cases
## 4 Spain_Deaths
```

You can use fill = "right" to set how to split observations like the second one, where there is no separator character ("\_"):

```
ebola %>%
 separate(col = ebola_key, into = c("country", "outcome"),
          sep = "_", fill = "right")
## # A tibble: 4 x 2
## country outcome
## <chr> <chr>
## 1 Liberia Cases
## 2 Liberia <NA>
## 3 Spain Cases
## 4 Spain Deaths
```

#### In-course exercise

We'll take a break now to do the In-Course Exercise (Section 6.4.3).

#### unite

The unite function does the reverse of the separate function: it lets you join several columns into a single column. For example, say you have data where year, month, and day are split into different columns:

```
## # A tibble: 4 x 3
## year month day
## <dbl> <dbl> <int>
## 1 2016. 10. 1
## 2 2016. 10. 2
## 3 2016. 10. 3
## 4 2016. 10. 4
```

#### unite

You can use unite to join these into a single column:

```
date_example %>%
  unite(col = date, year, month, day, sep = "-")
## # A tibble: 4 x 1
## date
## <chr>
## 1 2016-10-1
## 2 2016-10-2
## 3 2016-10-3
## 4 2016-10-4
```

#### unite

If the columns you want to unite are in a row (and in the right order), you can use the : syntax with unite:

```
date_example %>%
  unite(col = date, year:day, sep = "-")
## # A tibble: 4 x 1
## date
## <chr>
## 1 2016-10-1
## 2 2016-10-2
## 3 2016-10-3
## 4 2016-10-4
```

## Grouping with mutate versus summarize

So far, we have never used mutate with grouping.

You can use mutate after grouping—unlike summarize, the data will not be collapsed to fewer columns, but the summaries created by mutate will be added within each group.

For example, if you wanted to add the mean height and weight by sex to the nepali dataset, you could do that with group\_by and mutate (see next slide).

## Grouping with mutate versus summarize

```
nepali %>%
 group by(sex) %>%
 mutate(mean ht = mean(ht, na.rm = TRUE),
       mean wt = mean(wt, na.rm = TRUE)) %>%
 slice(1:3) %>% select(id, sex, wt, ht, mean ht, mean wt)
## # A tibble: 6 x 6
## # Groups: sex [2]
##
       id sex wt ht mean ht mean wt
## <int> <int> <dbl> <dbl> <dbl> <dbl>
## 1 120011 1 12.8 91.2 85.7 11.4
## 2 120011 1 12.8 93.9 85.7
                                  11.4
## 3 120011 1 13.1 95.2 85.7
                                  11.4
## 4 120012 2 14.9 104. 84.6
                                  10.9
## 5 120012 2 15.1 106. 84.6
                                  10.9
## 6 120012
             2 15.8 108. 84.6
                                   10.9
```

#### More on mutate

There are also some special functions that work well with mutate:

- lead: Measured value for following observation
- lag: Measured value for previous observation
- cumsum: Sum of all values up to this point
- cummax: Highest value up to this point
- cumany: For TRUE / FALSE, have any been TRUE up to this point

#### More on mutate

library(lubridate)

## # A tibble: 4 x 3

Here is an example of using lead and lag with mutate:

## 2 2016-10-02 2016-10-01 2016-10-03 ## 3 2016-10-03 2016-10-02 2016-10-04

## date yesterday tomorrow
## <date> <date> 
## 1 2016-10-01 NA 2016-10-02

#### slice

You can also group by a factor first using group\_by. Then, when you use slice, you will get the first few rows for each level of the group.

```
nepali %>%
  group_by(sex) %>%
  slice(1:2)
## # A tibble: 4 x 9
##
  # Groups:
            sex [2]
                             mage lit died alive
##
        id
             sex
                    wt
                         ht
                                                     age
     <int> <int> <dbl> <dbl> <int> <int> <int> <int> <int><</pre>
##
## 1 120011
                               35
               1 12.8
                       91.2
                                      0
                                           2
                                                 5
                                                      41
## 2 120011
                 12.8 93.9
                                                 5
                               35
                                                      45
## 3 120012
               2 14.9 104.
                               35
                                                 5
                                                      57
               2 15.1 106.
                               35
                                           2
                                                 5
## 4 120012
                                                      61
```

## arrange with group\_by

You can also group by a factor before arranging. In this case, all data for the first level of the factor will show up first, in the order given in arrange, then all data from the second level will show up in the specified order, etc.

```
nepali %>%
  group_by(sex) %>%
  arrange(desc(ht)) %>%
  slice(1:2)
```

```
## # A tibble: 4 x 9
## # Groups:
           sex [2]
##
       id
                     ht
                             lit died alive
           sex
                wt
                        mage
                                            age
##
    32
                                         5
                                             74
##
  1 360302
            1
              16.8
                   110.
##
  2 360392
            1 17.2
                  109.
                          35
                               0
                                             70
## 3 360791
            2 17.5 111.
                          32
                               0
                                    0
                                         5
                                             68
            2 17.0 111.
                                         8
                                             68
## 4 120112
                          52
                               0
                                    0
```

## semi\_join and anti\_join

These functions allow you to filter one dataframe on only values that **do** have a match in a second dataframe (semi\_join) or **do not** have a match in a second dataframe (anti\_join).

I'll show examples with the x and y dataframes defined before:

```
## 1 x 90
## 2 y 82
## 3 z 78
```

```
## course day
## 1 w Tues
## 2 x Mon / Fri
## 3 y Tue
```

## semi\_join and anti\_join

The semi\_join function filters: one dataframe on only values that **do** have a match in a second dataframe:

```
x %>%
semi_join(y)
```

```
## course grade
## 1 x 90
## 2 y 82
```

## semi\_join and anti\_join

The anti\_join function filters one dataframe on only values that **do not** have a match in a second dataframe:

```
y %>%
  anti_join(x)

## course day
## 1 w Tues
```

#### In-course exercise

We'll take a break now to do the In-Course Exercise (Section 6.4.4).

# Tidying with dplyr

#### VADeaths data

For this example, I'll use the VADeaths dataset that comes with  ${\sf R}.$ 

This dataset gives the death rates per 1,000 people in Virginia in 1940. It gives death rates by age, gender, and rural / urban:

```
data("VADeaths")
VADeaths
```

##		Rural	Male	Rural	${\tt Female}$	Urban	Male	Urban	Female
##	50-54		11.7		8.7		15.4		8.4
##	55-59		18.1		11.7		24.3		13.6
##	60-64		26.9		20.3		37.0		19.3
##	65-69		41.0		30.9		54.6		35.1
##	70-74		66.0		54.3		71.1		50.0

#### VADeaths data

There are a few things that make this data untidy:

- One variable (age category) is saved as row names, rather than a column.
- Other variables (gender, rural / urban) are in column names.
- Once you gather the data, you will have two variables (gender, rural / urban) in the same column.

In the following slides, we'll walk through how to tidy this data.

1. One variable (age category) is saved as row names, rather than a column.

To fix this, we need to convert the row names into a new column. We can do this using mutate (load tibble if needed):

```
VADeaths %>%
  as.data.frame() %>% ## Convert from matrix to dataframe
  rownames_to_column(var = "age")
```

##	age	Rural Mal	e Rural	Female	Urban	Male	Urban	Female
## 1	50-54	11.	7	8.7		15.4		8.4
## 2	55-59	18.	1	11.7		24.3		13.6
## 3	60-64	26.	9	20.3		37.0		19.3
## 4	65-69	41.	0	30.9		54.6		35.1
## 5	70-74	66.	0	54.3		71.1		50.0

2. Two variables (gender, rural / urban) are in column names.

Gather the data to convert column names to a new column:

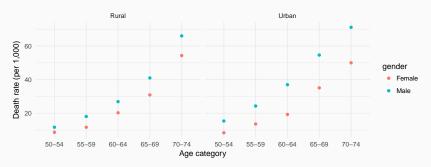
```
VADeaths %>%
 as.data.frame() %>%
 rownames to column(var = "age") %>%
 gather(key = gender loc, value = mort rate, - age) %>%
 slice(1:4)
## # A tibble: 4 x 3
##
  age gender_loc mort_rate
## <chr> <chr>
                        <dbl>
## 1 50-54 Rural Male 11.7
## 2 55-59 Rural Male 18.1
## 3 60-64 Rural Male 26.9
## 4 65-69 Rural Male 41.0
```

3. Two variables (gender, rural / urban) in the same column.

Separate the column into two separate columns for "gender" and "loc" (rural / urban):

```
## # A tibble: 1 x 4
## age gender loc mort_rate
## <chr> <chr> <chr> <chr> ## 1 50-54 Rural Male 11.7
```

Now that the data is tidy, it's much easier to plot:



## **FARS** data

#### FARS data

The US Department of Transportation runs the Fatality Analysis Reporting System (FARS), which gathers data on all fatal motor vehicle accidents. Here is a description from their documentation:

- Motor vehicle on a public road
- Resulted in a death within 30 days of the crash
- Includes crashes in the 50 states, DC, and Puerto Rico

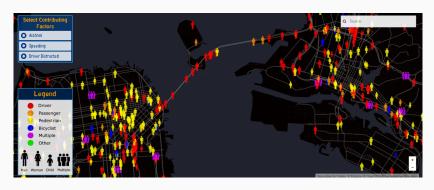
#### **FARS** data



Source: Max Galka

http://metrocosm.com/10-years-of-traffic-accidents-mapped.html

#### **FARS** data



Source: Max Galka

http://metrocosm.com/10-years-of-traffic-accidents-mapped.html

#### In-course exercise

We'll take a break now to do the In-Course Exercise (Section 6.4.5).

For these examples, we'll use some data on passengers of the Titanic. You can load this data using:

```
# install.packages("titanic")
library(titanic)
data("titanic_train")
```

We will be using the stringr package:

```
library(stringr)
```

This data includes a column called "Name" with passenger names. This column is somewhat messy and includes several elements that we might want to separate (last name, first name, title). Here are the first few values of "Name":

```
titanic_train %>% select(Name) %>% slice(1:3)

## # A tibble: 3 x 1

## Name

## <chr>
## 1 Braund, Mr. Owen Harris

## 2 Cumings, Mrs. John Bradley (Florence Briggs Thayer)

## 3 Heikkinen, Miss. Laina
```

The str\_trim function from the stringr package allows you to trim leading and trailing whitespace:

```
with_spaces <- c(" a ", " bob", " gamma")
with_spaces

## [1] " a " " bob" " gamma"

str_trim(with_spaces)

## [1] "a" "bob" "gamma"</pre>
```

This is rarer, but if you ever want to, you can add leading and / or trailing whitespace to elements of a character vector with str\_pad from the stringr package.

There are also functions to change a full character string to uppercase, lowercase, or title case:

```
titanic train$Name[1]
## [1] "Braund, Mr. Owen Harris"
str to upper(titanic train$Name[1])
## [1] "BRAUND, MR. OWEN HARRIS"
str_to_lower(titanic_train$Name[1])
## [1] "braund, mr. owen harris"
str to title(str to lower(titanic train$Name[1]))
## [1] "Braund, Mr. Owen Harris"
```

We've already done some things to manipulate strings. For example, if we wanted to separate "Name" into last name and first name (including title), we could actually do that with the separate function:

```
titanic train %>%
 select(Name) %>%
 slice(1:3) %>%
 separate(Name, c("last name", "first name"), sep = ", ")
## # A tibble: 3 x 2
##
    last_name first_name
## <chr> <chr>
## 1 Braund Mr. Owen Harris
## 2 Cumings Mrs. John Bradley (Florence Briggs Thayer)
## 3 Heikkinen Miss. Laina
```

Notice that separate is looking for a regular pattern (",") and then doing something based on the location of that pattern in each string (splitting the string).

There are a variety of functions in R that can perform manipulations based on finding regular patterns in character strings.

The str\_detect function will look through each element of a character vector for a designated pattern. If the pattern is there, it will return TRUE, and otherwise FALSE. The convention is:

For example, to create a logical vector specifying which of the Titanic passenger names include "Mrs.", you can call:

```
mrs <- str_detect(titanic_train$Name, "Mrs\\.")
head(mrs)</pre>
```

```
## [1] FALSE TRUE FALSE TRUE FALSE FALSE
```

titanic train %>%

The result is a logical vector, so str\_detect can be used in filter to subset data to only rows where the passenger's name includes "Mrs.":

```
filter(str detect(Name, "Mrs\\.")) %>%
  select(Name) %>%
  slice(1:3)
## # A tibble: 3 x 1
##
    Name
## <chr>
## 1 Cumings, Mrs. John Bradley (Florence Briggs Thayer)
## 2 Futrelle, Mrs. Jacques Heath (Lily May Peel)
## 3 Johnson, Mrs. Oscar W (Elisabeth Vilhelmina Berg)
```

As a note, in regular expressions, all of the following characters are special characters that need to be escaped with backslashes if you want to use them literally:

```
. * + ^ ? $ \ | ( ) [ ] { }
```

There is an older, base R function called grep1 that does something very similar (although note that the order of the arguments is reversed).

```
titanic train %>%
  filter(grepl("Mrs\\.", Name)) %>%
  select(Name) %>%
  slice(1:3)
## # A tibble: 3 x 1
##
    Name
##
   <chr>
## 1 Cumings, Mrs. John Bradley (Florence Briggs Thayer)
## 2 Futrelle, Mrs. Jacques Heath (Lily May Peel)
## 3 Johnson, Mrs. Oscar W (Elisabeth Vilhelmina Berg)
```

The str\_extract function can be used to extract a string (if it exists) from each value in a character vector. It follows similar conventions to str\_detect:

For example, you might want to extract "Mrs." if it exists in a passenger's name:

```
titanic_train %>%
  mutate(mrs = str_extract(Name, "Mrs\\.")) %>%
  select(Name, mrs) %>%
  slice(1:3)
```

Notice that now we're creating a new column (mrs) that either has "Mrs." (if there's a match) or is missing (NA) if there's not a match.

For this first example, we were looking for an exact string ("Mrs"). However, you can use patterns that match a particular pattern, but not an exact string. For example, we could expand the regular expression to find "Mr." or "Mrs.":

```
titanic_train %>%
 mutate(title = str_extract(Name, "Mr\\.|Mrs\\.")) %>%
  select(Name, title) %>%
  slice(1:3)
## # A tibble: 3 x 2
##
     Name
                                                           title
##
     <chr>>
                                                           <chr>>
## 1 Braund, Mr. Owen Harris
                                                           Mr.
## 2 Cumings, Mrs. John Bradley (Florence Briggs Thayer) Mrs.
## 3 Heikkinen, Miss. Laina
                                                           <NA>
```

Note that this pattern uses a special operator (|) to find one pattern **or** another. Double backslashs (\\) **escape** the special character ".".

Notice that "Mr." and "Mrs." both start with "Mr", end with ".", and may or may not have an "s" in between.

```
titanic_train %>%
  mutate(title = str_extract(Name, "Mr(s)*\\.")) %>%
  select(Name, title) %>%
  slice(1:3)
```

```
## # A tibble: 3 x 2
## Name title
## <chr>
## 1 Braund, Mr. Owen Harris Mr.
## 2 Cumings, Mrs. John Bradley (Florence Briggs Thayer) Mrs.
## 3 Heikkinen, Miss. Laina <NA>
```

This pattern uses (s)\* to match zero or more "s"s at this spot in the pattern.

In the previous code, we found "Mr." and "Mrs.", but missed "Miss.". We could tweak the pattern again to try to capture that, as well. For all three, we have the pattern that it starts with "M", has some lowercase letters, and then ends with ".".

```
titanic_train %>%
  mutate(title = str_extract(Name, "M[a-z]+\\.")) %>%
  select(Name, title) %>%
  slice(1:3)
```

```
## # A tibble: 3 x 2
## Name title
## <chr>
## 1 Braund, Mr. Owen Harris Mr.
## 2 Cumings, Mrs. John Bradley (Florence Briggs Thayer) Mrs.
## 3 Heikkinen, Miss. Laina Miss.
```

The last pattern used [a-z]+ to match one or more lowercase letters. The [a-z] is a **character class**.

You can also match digits ([0-9]), uppercase letters ([A-Z]), just some letters ([aeiou]), etc.

You can negate a character class by starting it with  $\hat{}$ . For example,  $[^0-9]$  will match anything that **isn't** a digit.

Sometimes, you want to match a pattern, but then only subset a part of it. For example, each passenger seems to have a title ("Mr.", "Mrs.", etc.) that comes after "," and before ".". We can use this pattern to find the title, but then we get some extra stuff with the match:

```
titanic_train %>%
  mutate(title = str_extract(Name, ",\\s[A-Za-z]*\\.\\s")) %>%
  select(title) %>%
  slice(1:3)
## # A tibble: 3 x 1
## title
## <chr>
## 1 ", Mr. "
## 2 ", Mrs. "
## 3 ", Miss. "
```

As a note, in this pattern, \\s is used to match a space.

We are getting things like ", Mr.", when we really want "Mr". We can use the str\_match function to do this. We group what we want to extract from the pattern in parentheses, and then the function returns a matrix. The first column is the full pattern match, and each following column gives just what matches within the groups.

```
## [,1] [,2]
## [1,] ", Mr. " "Mr"
## [2,] ", Mrs. " "Mrs"
## [3,] ", Miss. " "Miss"
## [4,] ", Mrs. " "Mrs"
## [5,] ", Mr. " "Mr"
## [6,] ", Mr. " "Mr"
```

To get just the title, then, we can run:

```
titanic_train %>%
  mutate(title =
           str match(Name, ",\\s([A-Za-z]*)\\.\\s")[ , 2]) %>%
  select(Name, title) %>%
  slice(1:3)
## # A tibble: 3 x 2
##
    Name
                                                          title
## <chr>
                                                          <chr>>
## 1 Braund, Mr. Owen Harris
                                                          Mr
## 2 Cumings, Mrs. John Bradley (Florence Briggs Thayer) Mrs
## 3 Heikkinen, Miss. Laina
                                                          Miss
```

The [ , 2] pulls out just the second column from the matrix returned by str\_match.

Here are some of the most common titles:

```
titanic_train %>%
 mutate(title =
          str match(Name, ",\\s([A-Za-z]*)\\.\\s")[ , 2]) %>%
 group_by(title) %>% summarize(n = n()) %>%
 arrange(desc(n)) %>% slice(1:5)
## # A tibble: 5 x 2
## title
          n
## <chr> <int>
## 1 Mr 517
## 2 Miss 182
## 3 Mrs 125
## 4 Master 40
## 5 Dr
```

titanic\_train %>%

The following slides have a few other examples of regular expressions in action with this dataset.

Get just names that start with ("^") the letter "A":

```
filter(str_detect(Name, "^A")) %>%
  select(Name) %>%
  slice(1:3)

## # A tibble: 3 x 1

## Name

## <chr>
## 1 Allen, Mr. William Henry

## 2 Andersson, Mr. Anders Johan
```

## 3 Asplund, Mrs. Carl Oscar (Selma Augusta Emilia Johansson)

Get names with "II" or "III" ( $\{2,\}$  says to match at least two times):

```
titanic train %>%
  filter(str detect(Name, "I{2,}")) %>%
  select(Name) %>%
  slice(1:3)
## # A tibble: 2 x 1
##
    Name
## <chr>
## 1 Carter, Master. William Thornton II
## 2 Roebling, Mr. Washington Augustus II
```

Get names with "Andersen" or "Anderson" (alternatives in square brackets):

```
titanic_train %>%
  filter(str_detect(Name, "Anders[eo]n")) %>%
  select(Name)
```

```
## Name
## 1 Andersen-Jensen, Miss. Carla Christine Nielsine
## 2 Anderson, Mr. Harry
## 3 Walker, Mr. William Anderson
## 4 Olsvigen, Mr. Thor Anderson
## 5 Soholt, Mr. Peter Andreas Lauritz Andersen
```

Get names that start with (" $^{"}$ " outside of brackets) the letters "A" and "B":

```
titanic_train %>%
  filter(str detect(Name, "^[AB]")) %>%
  select(Name) %>%
  slice(1:3)
## # A tibble: 3 x 1
##
    Name
## <chr>
## 1 Braund, Mr. Owen Harris
## 2 Allen, Mr. William Henry
## 3 Bonnell, Miss. Elizabeth
```

Get names that end with ("\$") the letter "b" (either lowercase or uppercase):

```
titanic_train %>%
  filter(str_detect(Name, "[bB]$")) %>%
  select(Name)
```

```
## Name
## 1 Emir, Mr. Farred Chehab
## 2 Goldschmidt, Mr. George B
## 3 Cook, Mr. Jacob
## 4 Pasic, Mr. Jakob
```

Some useful regular expression operators include:

Operator	Meaning
	Any character
	Match 0 or more times (greedy)
?	Match 0 or more times (non-greedy)
+	Match 1 or more times (greedy)
+?	Match 1 or more times (non-greedy)
^	Starts with (in brackets, negates)
\$	Ends with
[]	Character classes

For more on these patterns, see:

- Help file for the stringi-search-regex function in the stringi package (which should install when you install stringr)
- Chapter 14 of R For Data Science
- http://gskinner.com/RegExr: Interactive tool for helping you build regular expression pattern strings