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.

country	year	column	cases
AD	2000	m014	0
$^{\mathrm{AD}}$	2000	m1524	0
$^{\mathrm{AD}}$	2000	m2534	1
$^{\mathrm{AD}}$	2000	m3544	0
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	m	0-14	0
AD	2000	m	15-24	0
AD	2000	\mathbf{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	\mathbf{m}	0-14	2
AE	2000	m	15-24	4
\mathbf{AE}	2000	m	25 - 34	4
\mathbf{AE}	2000	\mathbf{m}	35-44	6
\mathbf{AE}	2000	\mathbf{m}	45-54	5
\mathbf{AE}	2000	m	55-64	12
\mathbf{AE}	2000	\mathbf{m}	65 +	10
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	$_{\mathrm{tmax}}$	_	_	_	_	_	_	_	_
MX17004	2010	4	$_{ m tmin}$	_	_	_	_	_	_	_	_
MX17004	2010	5	tmax	_	_	_	_	_	_	_	_
MX17004	2010	5	$_{ m tmin}$	_	_	_	_	_	_	_	_

Solution:

id	date	element	value	id	date	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 1 of the In-course Exercise for Chapter 6).

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.

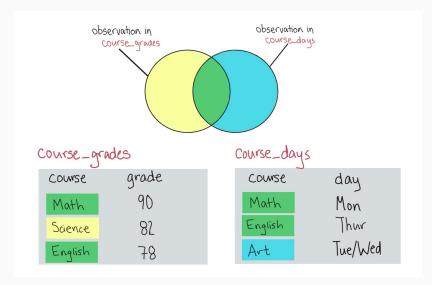
For example, for air pollution epidemiology, you will often have to join several datasets:

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

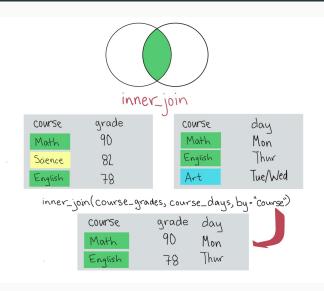
The dplyr package has a family of different functions to join two dataframes together, the *_join family of functions. These include:

- inner_join
- full_join
- left_join
- right_join

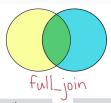
All combine two dataframes, which I'll call course_grades and course_days here.



inner_join



full_join



course	grade	ľ
Math	90	
Science	82	
English	78	

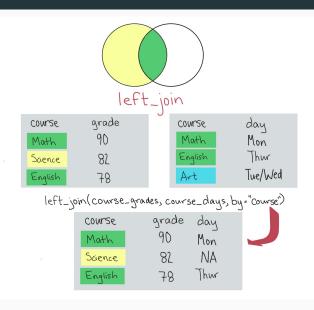
course	
Math	
English	
V 7	

day Mon Thur Tue/Wed

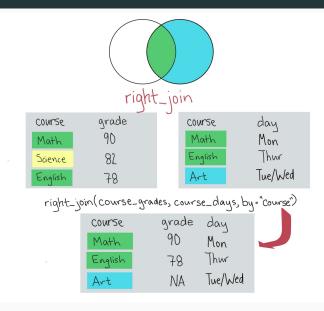
full_join(course_grades, course_days, by="course")

course	grade	day	
Math	90	Mon	
Science	82	NA	
English	78	Thur	
Art	NA	Tue/Wed	

left_join



right_join



For some more complex examples of using join, I'll use these example datasets (x and y):

```
## # A tibble: 4 x 3
## course grade student
## <chr> <dbl> <chr>
## 1 x
            92 a
## 2 x
            90 b
## 3 y
          82 a
## 4 z 78 b
## # 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
        Tue
## 4 y
                 а
```

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> <chr>
## 1 x
            92 a
                  Mon / Fri a
## 2 x
           92 a
                  Mon / Fri b
            90 b
## 3 x
                      Mon / Fri a
## 4 x
           90 b
                      Mon / Fri b
## 5 y
           82 a
                      Tue
                              а
## 6 z
           78 b
                      <NA> <NA>
           NA <NA>
## 7 w
                      Tues
                              а
```

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
## <chr> <dbl> <chr>
                       <chr>>
## 1 x
             92 a
                    Mon / Fri
## 2 x
             90 b
                       Mon / Fri
## 3 v
             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 2 of the In-course Exercise for Chapter 6).

Longer and wider data

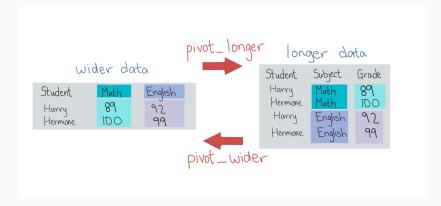
There are two functions from the tidyr package (another member of the tidyverse) that you can use to change between wide and long data: pivot_longer and pivot_wider.

These are brand new, and they replace the older gather and spread functions. To use the new functions, you many need to install the development version of the tidyr package. You can do that with:

```
devtools::install_github("tidyverse/tidyr")
```

Here is a description of these two functions:

- pivot_longer: Takes several columns and pivots them down into two columns. One of the new columns contains the former column names and the other contains the former cell values.
- pivot_wider: Takes two columns and pivots them up into multiple columns. Column names for the new columns will come from one column and the cell values from the other.



The following examples show the effects of making a dataset longer or wider.

Here is some example wide data:

hogwarts_wide

```
## # A tibble: 2 x 4
## student math english science
## <chr> <dbl> <dbl> <dbl> <dbl> > 93
## 1 Harry 89 92 93
## 2 Hermione 100 99 98
```

In the hogwarts_wide dataset, there are separate columns for three different courses (math, english, and science). Each cell gives the value for a certain stock on a certain day.

hogwarts_wide

```
## # A tibble: 2 x 4
## student math english science
## <chr> <dbl> <dbl> <dbl> <dbl> ## 1 Harry 89 92 93
## 2 Hermione 100 99 98
```

This data isn't "tidy", because the identify of the course (math, english, or science) is a variable, and you'll probably want to include it as a variable in modeling.

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

In this "longer" dataframe, there is now one column that gives the identify of the course (subject) and another column that gives the grade a student got for that course (grade):

```
hogwarts_long
```

```
## # A tibble: 6 x 3
##
    student subject grade
   <chr> <chr> <chr> <dbl>
##
## 1 Harry math
                        89
                        92
## 2 Harry english
  3 Harry science
                        93
##
## 4 Hermione math
                       100
## 5 Hermione english
                        99
## 6 Hermione science
                        98
```

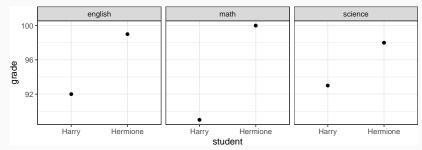
The format for a pivots_longer call is:

Three important notes:

- Everything is pivoted into one of two columns—one column with the old column names, and one column with the old cell values
- With the names_to and values_to arguments, you are just providing column names for the two columns that everything's pivoted into. When you are pivoting from "wide" to "long", you get to pick these names.
- If there is a column you don't want to include in the pivot (date in the example), use - to exclude it in the cols argument.

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

```
ggplot(hogwarts_long, aes(x = student, y = grade)) +
  geom_point() +
  facet_wrap(~ subject) +
  theme_bw()
```



If you have data in a "longer" format and would like to make it "wider", you can use pivot_wider to do that:

Notice that this reverses the action of pivot_longer.

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, pivoting, and joining different datasets.

Note: At this time "R for Data Science" uses the gather and spread instead of pivot_*. These are older functions, you should use pivot_*.

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

separate

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 \times 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 x 3
## ebola_key country outcome
## <chr> <chr> <chr>
## 1 Liberia Cases Liberia Cases
## 2 Liberia_Deaths Liberia Deaths
                  Spain Cases
## 3 Spain Cases
## 4 Spain Deaths Spain Deaths
```

separate

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 3 of the In-course Exercise for Chapter 6).

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	tibl	ole:	4	X	3
##		2	/ear	mont	th		day
##		<0	lbl>	<db1< td=""><td>1></td><td><j< td=""><td>nt></td></j<></td></db1<>	1>	<j< td=""><td>nt></td></j<>	nt>
##	1	2	2016	:	10		1
##	2	2	2016	:	10		2
##	3	2	2016	:	10		3
##	4	2	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

```
worldcup %>%
 group by(Position) %>%
 mutate(mean time = mean(Time)) %>%
 slice(1:2) %>% select(Team:Time, mean time)
## # A tibble: 8 x 4
## # Groups: Position [4]
   Team
             Position
##
                       Time mean time
## <fct>
             <fct> <int>
                               <dbl>
             Defender
## 1 France
                        180
                                242.
## 2 Ghana
             Defender
                        138
                                242.
## 3 Cameroon Forward
                         46
                                167.
## 4 Uruguay
              Forward 72
                                167.
## 5 Ivory Coast Goalkeeper
                        270
                                315.
                        270
## 6 Switzerland Goalkeeper
                                315.
             Midfielder 16
## 7 Algeria
                                192.
## 8 Japan
             Midfielder
                        351
                                192.
```

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.

```
worldcup %>%
  group_by(Position) %>%
  slice(1:2)
## # A tibble: 8 x 7
##
  # Groups:
               Position [4]
                 Position
                             Time Shots Passes Tackles Saves
##
    Team
##
     <fct>
                 <fct>
                            <int> <int> <int>
                                                 <int> <int>
##
  1 France
                 Defender
                              180
                                            91
                                                      6
  2 Ghana
                 Defender
                              138
                                            51
##
                               46
                                            16
  3 Cameroon
                 Forward
                                                      0
  4 Uruguay
                 Forward
                               72
                                            15
## 5 Ivory Coast Goalkeeper
                                            23
                                                            8
                              270
                                                      0
## 6 Switzerland Goalkeeper
                                            75
                                                           11
                              270
                                                      0
## 7 Algeria
                 Midfielder
                               16
                                      0
                                             6
                                                      0
                                                              52
```

arrange with group_by

worldcup %>%

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.

```
group by (Team) %>%
 arrange(desc(Saves)) %>%
 slice(1) %>%
 head(n = 4)
## # A tibble: 4 x 7
## # Groups: Team [4]
          Position
##
    Team
                          Time Shots Passes Tackles Saves
##
    <fct> <fct>
                         <int> <int> <int> <int> <int>
  1 Algeria Goalkeeper
                           180
                                  0
                                        30
                                                 0
                                                      12
                        450
                                        47
                                                      10
## 2 Argentina Goalkeeper
                                  0
                                                 0
## 3 Australia Goalkeeper
                           270
                                        51
                                                      13
                                  0
  4 Brazil
              Goalkeeper
                           450
                                  0
                                        69
                                                      10
```

53

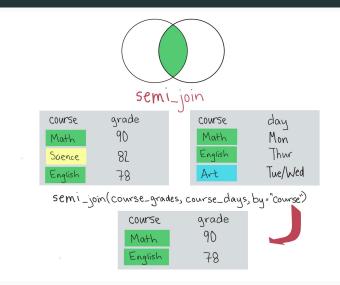
semi_join and anti_join

There are two more *_join functions we'll look at.

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).

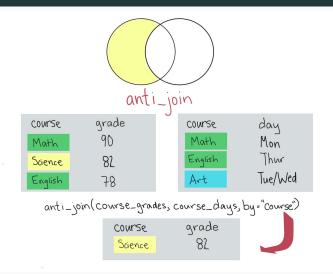
These functions do **not** bring in columns from the second dataset. Instead, they check the second dataset to decide whether or not to keep certain rows in the first dataset.

semi_join



The semi_join function filters to observations that **do** have a match in a second dataframe.

anti_join



The anti_join function filters to observations that **do not** have a match in a second dataframe.

In-course exercise

We'll take a break now to do the In-Course Exercise (Section 4 of the In-course Exercise for Chapter 6).

Tidying with dplyr

VADeaths data

For this example, I'll use the VADeaths dataset that comes with 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 Mal	e Rural	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.

A tibble: 5 x 5

41

66

(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_tibble() %>% ## Convert from matrix to dataframe
  rownames_to_column(var = "age")
```

##		age	`Rural Male`	`Rural Female`	`Urban Male`	`Urban Femal
##		<chr></chr>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<db< td=""></db<>
##	1	1	11.7	8.7	15.4	8
##	2	2	18.1	11.7	24.3	13
##	3	3	26.9	20.3	37	19

30.9

54.3

54.6

71.1

650

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

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

```
VADeaths %>%
  as_tibble() %>%
  rownames_to_column(var = "age") %>%
  gather(key = gender_loc, value = mort_rate, - age) %>%
  slice(1:4)
```

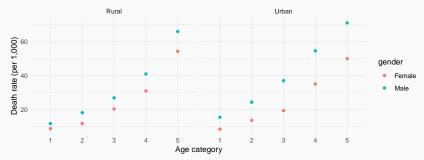
```
## # A tibble: 4 x 3
## age gender_loc mort_rate
## <chr> <chr> <chr> ## 1 1 Rural Male 11.7
## 2 2 Rural Male 18.1
## 3 3 Rural Male 26.9
## 4 4 Rural Male 41
```

(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 1 Rural Male 11.7
```

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

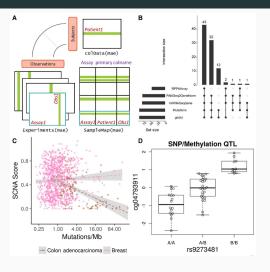


Bioconductor software

Biodiversity data

Bioconductor provides opensource software for bioinfomatics. Bioconductor provides an R package called microbiome that includes tools for exploring and analysing microbiome profiling data. The package provides sample datasets.

Biodiversity data



Source: Ramos et al. Cancer Research 2017.

https://cancerres.aacrjournals.org/content/77/21/e39

In-course exercise

We'll take a break now to do the In-Course Exercise (Section 5 of the In-course Exercise for Chapter 6).

forcats

forcats

Hadley Wickham has developed a package called forcats that helps you work with factors.

library(forcats)

The fct_recode function can be used to change the labels of a function (along the lines of using factor with levels and labels to reset factor labels).

One big advantage is that fct_recode lets you change labels for some, but not all, levels. For example, here are the team names:

```
worldcup %>%
filter(Team == "USA") %>%
slice(1:3) %>% select(Team, Position, Time)

## Team Position Time
## 1 USA Midfielder 10
## 2 USA Defender 390
## 3 USA Defender 200
```

```
If you just want to change "USA" to "United States", you can run:
worldcup <- worldcup %>%
  mutate(Team = fct_recode(Team, `United States` = "USA"))
worldcup %>%
  filter(Team == "United States") %>%
  slice(1:3) %>% select(Team, Position, Time)
##
              Team
                    Position Time
## 1 United States Midfielder
                              10
## 2 United States Defender 390
## 3 United States Defender 200
```

forcats

You can use the fct_lump function to lump uncommon factors into an "Other" category. For example, to lump the two least common positions together, you can run (n specifies how many categories to keep outside of "Other"):

forcats

You can use the fct_infreq function to reorder the levels of a factor from most common to least common:

```
levels(worldcup$Position)

## [1] "Defender" "Forward" "Goalkeeper" "Midfielder"

worldcup <- worldcup %>%
   mutate(Position = fct_infreq(Position))
levels(worldcup$Position)

## [1] "Midfielder" "Defender" "Forward" "Goalkeeper"
```

If you want to reorder one factor by another variable (ascending order), you can use fct_reorder (e.g., homework 3). For example, to re-level Position by the average shots on goals for each position, you can run:

```
levels(worldcup$Position)
## [1] "Midfielder" "Defender" "Forward"
                                              "Goalkeeper"
worldcup <- worldcup %>%
  group_by(Position) %>%
  mutate(ave shots = mean(Shots)) %>%
  ungroup() %>%
  mutate(Position = fct_reorder(Position, ave_shots))
levels(worldcup$Position)
## [1] "Goalkeeper" "Defender" "Midfielder" "Forward"
```

In-course exercise

We'll now take a break to do Section 6 of the In-course Exercise for Chapter 6.

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":

```
## Name
## 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:

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
```

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.":

```
titanic_train %>%
  filter(str_detect(Name, "Mrs\\.")) %>%
  select(Name) %>%
  slice(1:3)
```

```
## Name
## 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)
```

```
## Name
## 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)
```

```
## Name title
## 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 ($\backslash \backslash$) **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)
```

```
## Name title
## 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)
```

```
## Name title
## 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)

## title
## 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.

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

```
## Name title
## 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
```

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":

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

```
## Name
## 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)

## Name
## 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)
```

```
## Name
## 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

Tidy select

There are tidyverse functions to make selecting variables more straightforwards. You can call these functions as arguments of the select function to streamline variable selection. Examples include: starts_with(), ends_with(), and contains().

Tidy select (helpers)

Here we use starts_with("t") to select all variables that begin with t.

```
titanic_train %>%
  select(starts_with("t")) %>%
  slice(1:3)
```

```
## Ticket
## 1 A/5 21171
## 2 PC 17599
## 3 STON/02. 3101282
```

Tidy select

The are also tidyverse functions that allow us to easily operate on a selection of variables. These functions are called scoped varients. You can identify these functions by these _all, _at, and _if suffixes.

Tidy select (*_if)

Here we use select_if to select all the numeric variables in a dataframe and covert their names to lower case (a handy function to tidy the variable names).

```
titanic_train %>%
  select_if(is.numeric, tolower) %>%
  slice(1:3)
```

##		passengerid	survived	pclass	age	sibsp	parch	fare
##	1	1	0	3	22	1	0	7.2500
##	2	2	1	1	38	1	0	71.2833
##	3	3	1	3	26	0	0	7.9250

Tidy select (*_if)

The select_if function takes the following form.

Tidy select (*_at)

Here we use select_at to select all the variables that contain ss in their name and then covert their names to lower case (a handy function to tidy the variable names).

```
titanic_train %>%
  select_at(vars(contains("ss")), tolower) %>%
  slice(1:3)
```

```
## passengerid pclass
## 1 1 3
## 2 2 1
## 3 3 3
```

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

In-course exercise

We'll now take a break to do Section 7 of the In-course Exercise for Chapter 6.