

Tidying data

Tidying data

- ▶ Data rarely come to us as we want to use them.
- ▶ Before we can do analysis, typically have organizing to do.
- ▶ This is typical of ANOVA-type data, “wide format”:

pig	feed1	feed2	feed3	feed4
1	60.8	68.7	92.6	87.9
2	57.0	67.7	92.1	84.2
3	65.0	74.0	90.2	83.1
4	58.6	66.3	96.5	85.7
5	61.7	69.8	99.1	90.3

- ▶ 20 pigs randomly allocated to one of four feeds. At end of study, weight of each pig is recorded.
- ▶ Are any differences in mean weights among the feeds?
- ▶ Problem: want all weights in one column, with 2nd column labelling which feed. Untidy!

Tidy and untidy data (Wickham)

- ▶ Data set easier to deal with if:
 - ▶ each observation is one row
 - ▶ each variable is one column
 - ▶ each type of observation unit is one table
- ▶ Data arranged this way called “tidy”; otherwise called “untidy”.
- ▶ For the pig data:
 - ▶ response variable is weight, but scattered over 4 columns, which are levels of a factor feed.
 - ▶ Want all the weights in one column, with a second column feed saying which feed that weight goes with.
 - ▶ Then we can run aov.

Packages for this section

```
library(tidyverse)
```

Reading in the pig data

```
my_url <- "http://ritsokiguess.site/datafiles/pigs1.txt"  
pigs1 <- read_delim(my_url, " ")  
pigs1
```

```
# A tibble: 5 x 5  
#>   pig feed1 feed2 feed3 feed4  
#>   <dbl> <dbl> <dbl> <dbl> <dbl>  
#> 1     1   60.8  68.7  92.6  87.9  
#> 2     2    57    67.7  92.1  84.2  
#> 3     3    65    74    90.2  83.1  
#> 4     4   58.6  66.3  96.5  85.7  
#> 5     5   61.7  69.8  99.1  90.3
```

Making it longer

- ▶ We wanted all the weights in one column, labelled by which feed they went with.
- ▶ This is a very common reorganization, and the magic “verb” is `pivot_longer`:

```
pigs1 %>% pivot_longer(feed1:feed4, names_to="feed",
                           values_to="weight") -> pigs2
pigs2
```

```
# A tibble: 20 x 3
  pig   feed  weight
  <dbl> <chr> <dbl>
1     1 feed1  60.8
2     1 feed2  68.7
3     1 feed3  92.6
4     1 feed4  87.9
5     2 feed1  57
6     2 feed2  67.7
7     2 feed3  92.1
```

Alternatives

Any way of choosing the columns to pivot longer is good, eg:

```
pigs1 %>% pivot_longer(-pig, names_to="feed",
                           values_to="weight")
```

```
# A tibble: 20 x 3
  pig   feed  weight
  <dbl> <chr> <dbl>
1 1     feed1  60.8
2 1     feed2  68.7
3 1     feed3  92.6
4 1     feed4  87.9
5 2     feed1  57
6 2     feed2  67.7
7 2     feed3  92.1
8 2     feed4  84.2
9 3     feed1  65
10 3    feed2  74
11 3    feed3  90.2
```

Comments

- ▶ pigs2 now in “long” format, ready for analysis.
- ▶ Anatomy of `pivot_longer`:
 - ▶ columns to combine
 - ▶ a name for column that will contain groups (“names”)
 - ▶ a name for column that will contain measurements (“values”)

Identifying the pigs

- ▶ Values in pig identify pigs *within each group*: pig 1 is four different pigs!
- ▶ Create unique pig IDs by gluing pig number onto feed:

```
pigs2 %>% mutate(pig_id=str_c(feed, "_", pig)) -> pigs2  
pigs2
```

```
# A tibble: 20 x 4  
  pig feed  weight pig_id  
  <dbl> <chr>  <dbl> <chr>  
1     1 feed1   60.8 feed1_1  
2     1 feed2   68.7 feed2_1  
3     1 feed3   92.6 feed3_1  
4     1 feed4   87.9 feed4_1  
5     2 feed1    57    feed1_2  
6     2 feed2   67.7 feed2_2  
7     2 feed3   92.1 feed3_2  
8     2 feed4   84.2 feed4_2  
9     3 feed1   65    feed1_3
```

...and finally, the analysis

- ▶ which is just what we saw before:

```
weight.1 <- aov(weight ~ feed, data = pigs2)
summary(weight.1)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)					
feed	3	3521	1173.5	119.1	3.72e-11 ***					
Residuals	16	158	9.8							

Signif. codes:	0	'***'	0.001	'**'	0.01	'*'	0.05	'..'	0.1	'

- ▶ The mean weights of pigs on the different feeds are definitely not all equal.
- ▶ So we run Tukey to see which ones differ (over).

Tukey

```
TukeyHSD(weight.1)
```

Tukey multiple comparisons of means
95% family-wise confidence level

Fit: aov(formula = weight ~ feed, data = pigs2)

\$feed

	diff	lwr	upr	p adj
feed2-feed1	8.68	3.001038	14.358962	0.0024000
feed3-feed1	33.48	27.801038	39.158962	0.0000000
feed4-feed1	25.62	19.941038	31.298962	0.0000000
feed3-feed2	24.80	19.121038	30.478962	0.0000000
feed4-feed2	16.94	11.261038	22.618962	0.0000013
feed4-feed3	-7.86	-13.538962	-2.181038	0.0055599

All of the feeds differ!

Mean weights by feed

To find the best and worst, get mean weight by feed group. I borrowed an idea from earlier to put the means in descending order:

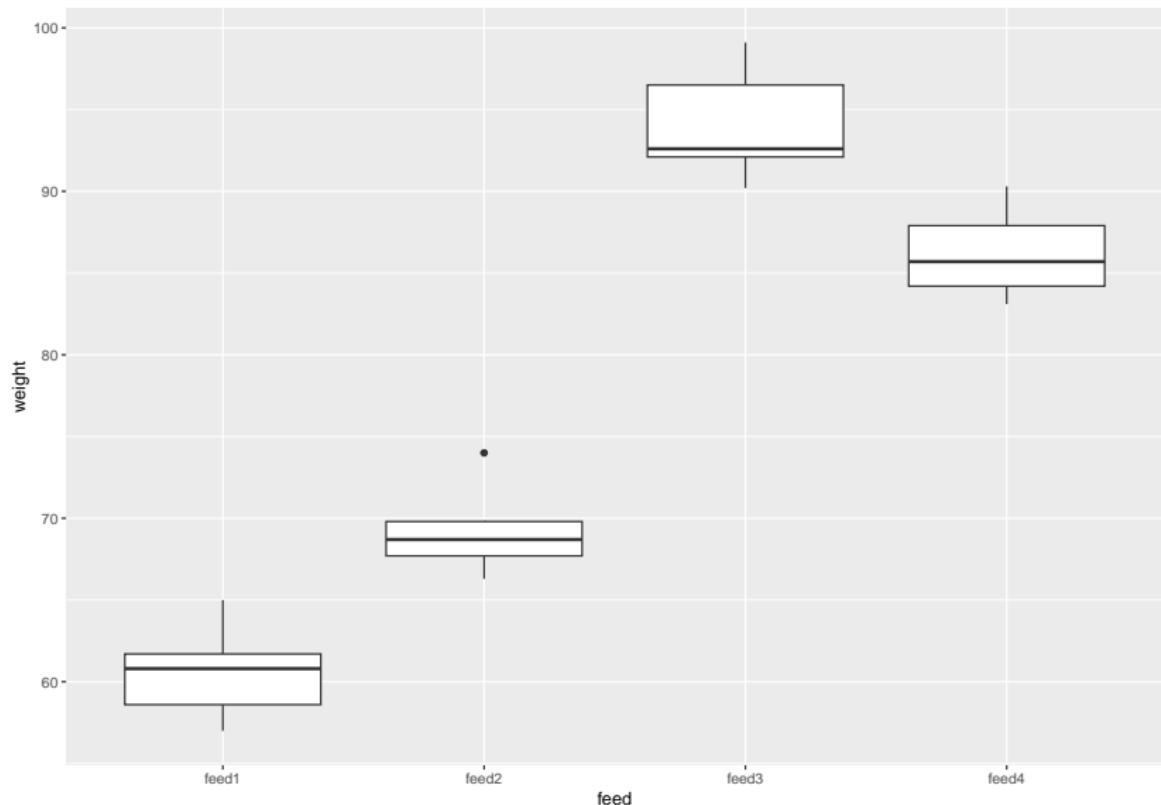
```
pigs2 %>%
  group_by(feed) %>%
  summarize(mean_weight = mean(weight))%>%
  arrange(desc(mean_weight))
```

```
# A tibble: 4 x 2
  feed   mean_weight
  <chr>     <dbl>
1 feed3      94.1
2 feed4      86.2
3 feed2      69.3
4 feed1      60.6
```

Feed 3 is best, feed 1 worst.

Should we have any concerns about the ANOVA?

```
ggplot(pigs2, aes(x = feed, y = weight)) + geom_boxplot()
```



Comments

- ▶ Feed 2 has an outlier
- ▶ But there are only 5 pigs in each group
- ▶ The conclusion is so clear that I am OK with this.

Tuberculosis

- ▶ The World Health Organization keeps track of number of cases of various diseases, eg. tuberculosis.
- ▶ Some data:

```
my_url <- "http://ritsokiguess.site/datafiles/tb.csv"  
tb <- read_csv(my_url)
```

The data (randomly chosen rows)

```
tb %>% slice_sample(n = 10)
```

```
# A tibble: 10 x 22
```

	iso2	year	m04	m514	m014	m1524	m2534	m3544	m4554	m5
	<chr>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>
1	ZM	1995	NA	NA	91	659	1668	1124	487	
2	NZ	1982	NA	NA	NA	NA	NA	NA	NA	
3	BR	1982	NA	NA	NA	NA	NA	NA	NA	
4	NL	2002	NA	NA	1	40	54	39	33	
5	CO	1988	NA	NA	NA	NA	NA	NA	NA	
6	RW	1995	NA	NA	NA	NA	NA	NA	NA	
7	PK	1987	NA	NA	NA	NA	NA	NA	NA	
8	SL	1998	NA	NA	14	226	445	338	191	
9	PE	1981	NA	NA	NA	NA	NA	NA	NA	
10	ZM	1986	NA	NA	NA	NA	NA	NA	NA	
					mu <dbl>, f04 <dbl>, f514 <dbl>, f01					
					f1524 <dbl>, f2534 <dbl>, f3544 <dbl>, f4554 <dbl>, f55					
					f65 <dbl>, fu <dbl>					

What we have

- ▶ Variables: country (abbreviated), year. Then number of cases for each gender and age group, eg. m1524 is males aged 15–24. Also mu and fu, where age is unknown.
- ▶ Lots of missings. Want to get rid of.
- ▶ Abbreviations here.

```
tb %>%  
  pivot_longer(m04:fu, names_to = "genage",  
               values_to = "freq", values_drop_na = TRUE)
```

- ▶ Code for `pivot_longer`:
 - ▶ columns to make longer
 - ▶ column to contain the names (categorical)
 - ▶ column to contain the values (quantitative)
 - ▶ drop missings in the values

Results (some)

tb2

```
# A tibble: 35,750 x 4
  iso2    year genage   freq
  <chr> <dbl> <chr>   <dbl>
1 AD      1996 m014     0
2 AD      1996 m1524    0
3 AD      1996 m2534    0
4 AD      1996 m3544    4
5 AD      1996 m4554    1
6 AD      1996 m5564    0
7 AD      1996 m65      0
8 AD      1996 f014     0
9 AD      1996 f1524    1
10 AD     1996 f2534    1
# i 35,740 more rows
```

Separating

- ▶ 4 columns, but 5 variables, since genage contains both gender and age group. Split that up using separate.
- ▶ separate needs 3 things:
 - ▶ what to separate (no quotes needed),
 - ▶ what to separate into (here you do need quotes),
 - ▶ how to split.
- ▶ For “how to split”, here “after first character”:

```
tb2 %>% separate_wider_position(genage,
                                    widths = c("gender" = 1, "a
                                    too_few = "align_start") ->
tb3
```

```
# A tibble: 35,750 x 5
  iso2    year gender age     freq
  <chr> <dbl> <chr> <chr> <dbl>
1 AD      1996 m     014     0
2 AD      1996 m     1524    0
3 AD      1996 m     2534    0
4 AD      1996 m     3544    0
```

Tidied tuberculosis data (some)

tb3

```
# A tibble: 35,750 x 5
  iso2    year gender age     freq
  <chr> <dbl> <chr> <chr> <dbl>
1 AD      1996 m     014     0
2 AD      1996 m     1524    0
3 AD      1996 m     2534    0
4 AD      1996 m     3544    4
5 AD      1996 m     4554    1
6 AD      1996 m     5564    0
7 AD      1996 m     65      0
8 AD      1996 f     014     0
9 AD      1996 f     1524    1
10 AD     1996 f     2534   1
# i 35,740 more rows
```

In practice...

- ▶ instead of doing the pipe one step at a time, you *debug* it one step at a time, and when you have each step working, you use that step's output as input to the next step, thus:

```
tb %>%  
  pivot_longer(m04:fu, names_to = "genage",  
               values_to = "freq", values_drop_na = TRUE) %>%  
  separate_wider_position(genage,  
                           widths = c("gender" = 1, "age" =  
                           too_few = "align_start")
```

```
# A tibble: 35,750 x 5  
  iso2    year gender age     freq  
  <chr>   <dbl> <chr>  <chr>   <dbl>  
1 AD      1996 m    014     0  
2 AD      1996 m    1524    0  
3 AD      1996 m    2534    0  
4 AD      1996 m    3544    4  
5 AD      1996 m    4554    1
```

Total tuberculosis cases by year (some of the years)

```
tb3 %>%
  filter(between(year, 1991, 1998)) %>%
  group_by(year) %>% summarize(total=sum(freq))
```

```
# A tibble: 8 x 2
  year   total
  <dbl>   <dbl>
1 1991     544
2 1992     512
3 1993     492
4 1994     750
5 1995 513971
6 1996 635705
7 1997 733204
8 1998 840389
```

- ▶ Something very interesting happened between 1994 and 1995.

To find out what

- ▶ try counting up total cases by country:

```
tb3 %>% group_by(iso2) %>%
  summarize(total=sum(freq)) %>%
  arrange(desc(total))
```

```
# A tibble: 213 x 2
  iso2     total
  <chr>    <dbl>
1 CN      4065174
2 IN      3966169
3 ID      1129015
4 ZA      900349
5 BD      758008
6 VN      709695
7 CD      603095
8 PH      490040
9 BR      440609
10 KE     431523
```

What years do I have for China?

China started recording in 1995, which is at least part of the problem:

```
tb3 %>% filter(iso2=="CN") %>%
  group_by(year) %>%
  summarize(total=sum(freq))
```

```
# A tibble: 14 x 2
```

	year	total
	<dbl>	<dbl>
1	1995	131194
2	1996	168270
3	1997	195895
4	1998	214404
5	1999	212258
6	2000	213766
7	2001	212766
8	2002	194972
9	2003	267280

First year of recording by country?

- ▶ A lot of countries started recording in about 1995, in fact:

```
tb3 %>% group_by(iso2) %>%
  summarize(first_year=min(year)) %>%
  count(first_year)
```

```
# A tibble: 14 x 2
  first_year     n
  <dbl> <int>
1 1980          2
2 1994          2
3 1995         130
4 1996          31
5 1997          17
6 1998           6
7 1999          10
8 2000           4
9 2001           1
10 2002          3
```

Some Toronto weather data

```
my_url <-  
  "http://ritsokiguess.site/STAC32/toronto_weather.csv"  
weather <- read_csv(my_url)  
weather
```

```
# A tibble: 24 x 35
```

The columns

- ▶ Daily weather records for “Toronto City” weather station in 2018:
 - ▶ station: identifier for this weather station (always same here)
 - ▶ Year, Month
 - ▶ element: whether temperature given was daily max or daily min
 - ▶ d01, d02,... d31: day of the month from 1st to 31st.

Off we go

Numbers in data frame all temperatures (for different days of the month), so first step is

```
weather %>%
  pivot_longer(d01:d31, names_to="day",
               values_to="temperature",
               values_drop_na = TRUE)
```

	# A tibble: 703 x 6	station	Year	Month	element	day	temperature
		<chr>	<dbl>	<chr>	<chr>	<chr>	<dbl>
1	TORONTO CITY	2018	01	tmax	d01		-7.9
2	TORONTO CITY	2018	01	tmax	d02		-7.1
3	TORONTO CITY	2018	01	tmax	d03		-5.3
4	TORONTO CITY	2018	01	tmax	d04		-7.7
5	TORONTO CITY	2018	01	tmax	d05		-14.7
6	TORONTO CITY	2018	01	tmax	d06		-15.4
7	TORONTO CITY	2018	01	tmax	d07		-1
8	TORONTO CITY	2018	01	tmax	d08		3

Element

- ▶ Column element contains names of two different variables, that should each be in separate column.
- ▶ Distinct from eg. `m1524` in tuberculosis data, that contained levels of two different factors, handled by `separate`.
- ▶ Untangling names of variables handled by `pivot_wider`.

Handling element

```
weather %>%
  pivot_longer(d01:d31, names_to="day",
               values_to="temperature",
               values_drop_na = TRUE) %>%
  pivot_wider(names_from=element,
              values_from=temperature)
```

A tibble: 355 x 6

	station	Year	Month	day	tmax	tmin
	<chr>	<dbl>	<chr>	<chr>	<dbl>	<dbl>
1	TORONTO CITY	2018	01	d01	-7.9	-18.6
2	TORONTO CITY	2018	01	d02	-7.1	-12.5
3	TORONTO CITY	2018	01	d03	-5.3	-11.2
4	TORONTO CITY	2018	01	d04	-7.7	-19.7
5	TORONTO CITY	2018	01	d05	-14.7	-20.6
6	TORONTO CITY	2018	01	d06	-15.4	-22.3
7	TORONTO CITY	2018	01	d07	-1	-17.5
8	TORONTO CITY	2018	01	d08	3	-1.7
9	TORONTO CITY	2018	01	d09	1.6	0.6

Further improvements 1/2

- ▶ We have tidy data now, but can improve things further.
- ▶ `mutate` creates new columns from old (or assign back to change a variable).
- ▶ Would like numerical dates. `separate` works, or pull out number as below.
- ▶ `select` keeps columns (or drops, with minus). Station name has no value to us.

Further improvements 2/2

```
weather %>%
  pivot_longer(d01:d31, names_to="day",
               values_to="temperature", values_drop_na = TRUE)
  pivot_wider(names_from=element, values_from=temperature)
  mutate(Day = parse_number(day)) %>%
  select(-station)
```

A tibble: 355 x 6

	Year	Month	day	tmax	tmin	Day
1	2018	01	d01	-7.9	-18.6	1
2	2018	01	d02	-7.1	-12.5	2
3	2018	01	d03	-5.3	-11.2	3
4	2018	01	d04	-7.7	-19.7	4
5	2018	01	d05	-14.7	-20.6	5
6	2018	01	d06	-15.4	-22.3	6
7	2018	01	d07	-1	-17.5	7
8	2018	01	d08	3	-1.7	8
9	2018	01	d09	1.6	-0.6	9

Final step(s)

- ▶ Make year-month-day into proper date.
- ▶ Keep only date, tmax, tmin:

```
weather %>%
  pivot_longer(d01:d31, names_to="day",
               values_to="temperature", values_drop_na = T)
  pivot_wider(names_from=element, values_from=temperature)
  mutate(Day = parse_number(day)) %>%
  select(-station) %>%
  unite(datestr, c(Year, Month, Day), sep = "-") %>%
  mutate(date = as.Date(datestr)) %>%
  select(c(date, tmax, tmin)) -> weather_tidy
```

Our tidy data frame

```
weather_tidy
```

```
# A tibble: 355 x 3
  date       tmax  tmin
  <date>     <dbl> <dbl>
1 2018-01-01 -7.9 -18.6
2 2018-01-02 -7.1 -12.5
3 2018-01-03 -5.3 -11.2
4 2018-01-04 -7.7 -19.7
5 2018-01-05 -14.7 -20.6
6 2018-01-06 -15.4 -22.3
7 2018-01-07 -1      -17.5
8 2018-01-08 3      -1.7
9 2018-01-09 1.6     -0.6
10 2018-01-10 5.9     -1.3
# i 345 more rows
```

Plotting the temperatures

- ▶ Plot temperature against date joined by lines, but with separate lines for max and min. ggplot requires something like

```
ggplot(..., aes(x = date, y = temperature)) + geom_point() +  
  geom_line()
```

only we have two temperatures, one a max and one a min, that we want to keep separate.

- ▶ The trick: combine `tmax` and `tmin` together into one column, keeping track of what kind of temp they are. (This actually same format as untidy `weather`.) Are making `weather_tidy` untidy for purposes of drawing graph only.
- ▶ Then can do something like

```
ggplot(d, aes(x = date, y = temperature, colour = maxmin))  
  + geom_point() + geom_line()
```

to distinguish max and min on graph.

Setting up plot

- ▶ Since we only need data frame for plot, we can do the column-creation and plot in a pipeline.
- ▶ For a ggplot in a pipeline, the initial data frame is omitted, because it is whatever came out of the previous step.
- ▶ To make those “one column”s: pivot_longer. I save the graph to show overleaf:

```
weather_tidy %>%  
  pivot_longer(tmax:tmin, names_to="maxmin",  
              values_to="temperature") %>%  
  ggplot(aes(x = date, y = temperature, colour = maxmin))  
    geom_line() -> g
```

The plot

g



Summary of tidying “verbs”

Verb	Purpose
pivot_longer	Combine columns that measure same thing into one
pivot_wider	Take column that measures one thing under different conditions and put into multiple columns
separate	Turn a column that encodes several variables into several columns
unite	Combine several (related) variables into one “combination” variable

pivot_longer and pivot_wider are opposites; separate and unite are opposites.