When pivot-wider goes wrong

Packages

The inevitable:

library(tidyverse)

Some long data that should be wide

d

- Six observations of variable y, but three measured before some treatment and three measured after.
- Really matched pairs, so want column of y-values for pre and for post.
- pivot_wider.

What happens here?

```
d %>% pivot_wider(names_from = time, values_from = y)
```

```
# A tibble: 6 x 3
   obs pre post
 <dbl> <dbl> <dbl>
         19
              NA
    2 NA
             18
3
      17
              NΑ
      NA
              16
5
    5 15
              NA
6
         NA
              14
```

- Should be three pre values and three post. Why did this happen?
- pivot_wider needs to know which row to put each observation in.
- Uses combo of columns not named in pivot_wider, here obs (only).

The problem

```
d %>% pivot_wider(names_from = time, values_from = y)
```

```
# A tibble: 6 x 3
   obs pre post
 <dbl> <dbl> <dbl>
        19
             NA
    2 NA 18
3
   3 17 NA
  4
      NA 16
5
      15 NA
6
    6
        NΑ
             14
```

- There are 6 different obs values, so 6 different rows.
- No data for obs 2 and pre, so that cell missing (NA).
- Not enough data (6 obs) to fill $12 (= 2 \times 6)$ cells.
- obs needs to say which subject provided which 2 observations.

Fixing it up

d2

```
# A tibble: 6 x 3
 subject time
   <dbl> <chr> <dbl>
       1 pre
                19
       1 post 18
3
             17
      2 pre
4
     2 post 16
      3 pre
5
             15
6
      3 post
                14
```

- column subject shows which subject provided each pre and post.
- when we do pivot_wider, now only 3 rows, one per subject.

Coming out right

```
d2 %>% pivot_wider(names_from = time, values_from = y)
```

- row each observation goes to determined by other column subject,
 and now a pre and post for each subject.
- right layout for matched pairs t or to make differences for sign test or normal quantile plot.

Another example

• Two independent samples this time

```
A tibble: 8 \times 2
 group
 <chr>
           <dbl>
1 control
              11
2 control
         13
3 control
4 control
         14
5 treatment 12
          15
6 treatment
7 treatment
           16
8 treatment
              17
```

- These should be arranged like this
- but what if we make them wider?

Wider

```
d3 %>% pivot_wider(names_from = group, values_from = y)
```

```
control treatment
    list> < list>
1 < dbl [4] > < dbl [4] >
```

A tibble: 1×2

- row determined by what not used for pivot_wider: nothing!
- everything smooshed into one row!
- this time, too much data for the layout.
- Four data values squeezed into each of the two cells: "list-columns".

Get the data out

 To expand list-columns out into the data values they contain, can use unnest:

```
d3 %>% pivot_wider(names_from = group, values_from = y) %>%
  unnest(c(control, treatment))
```

• in this case, wrong layout, because data values not paired.