

Practical Approaches to Bioanalysis in R

Intro to the tidyverse & making data tidy with tidyr

Tidy data

“Tidy datasets are all alike but every messy dataset is messy in its own way” — Hadley Wickham

Tidy data

Three rules:

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

Example: Contingency table

	survived	died	
drug	15	3	not tidy
placebo	4	12	

Example: Contingency table

	survived	died	
drug	15	3	not tidy
placebo	4	12	

tidy	treatment	outcome	count
	drug	survived	15
	drug	died	3
	placebo	survived	4
	placebo	died	12

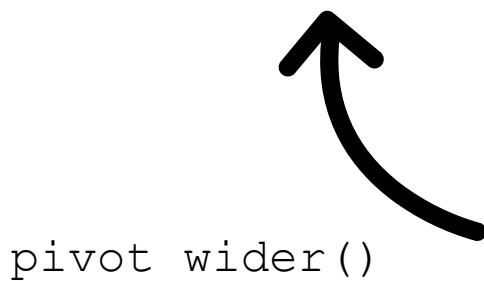
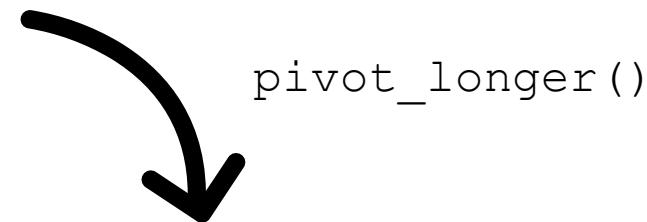
Example: Contingency table

	survived	died	
drug	15	3	not tidy
placebo	4	12	

	patient	treatment	outcome
tidy	1	drug	survived
	2	drug	died
	3	drug	survived
	4	placebo	died
		•	•
		•	•

tidyverse library provides functions for transforming tables

	survived	died
drug	15	3
placebo	4	12



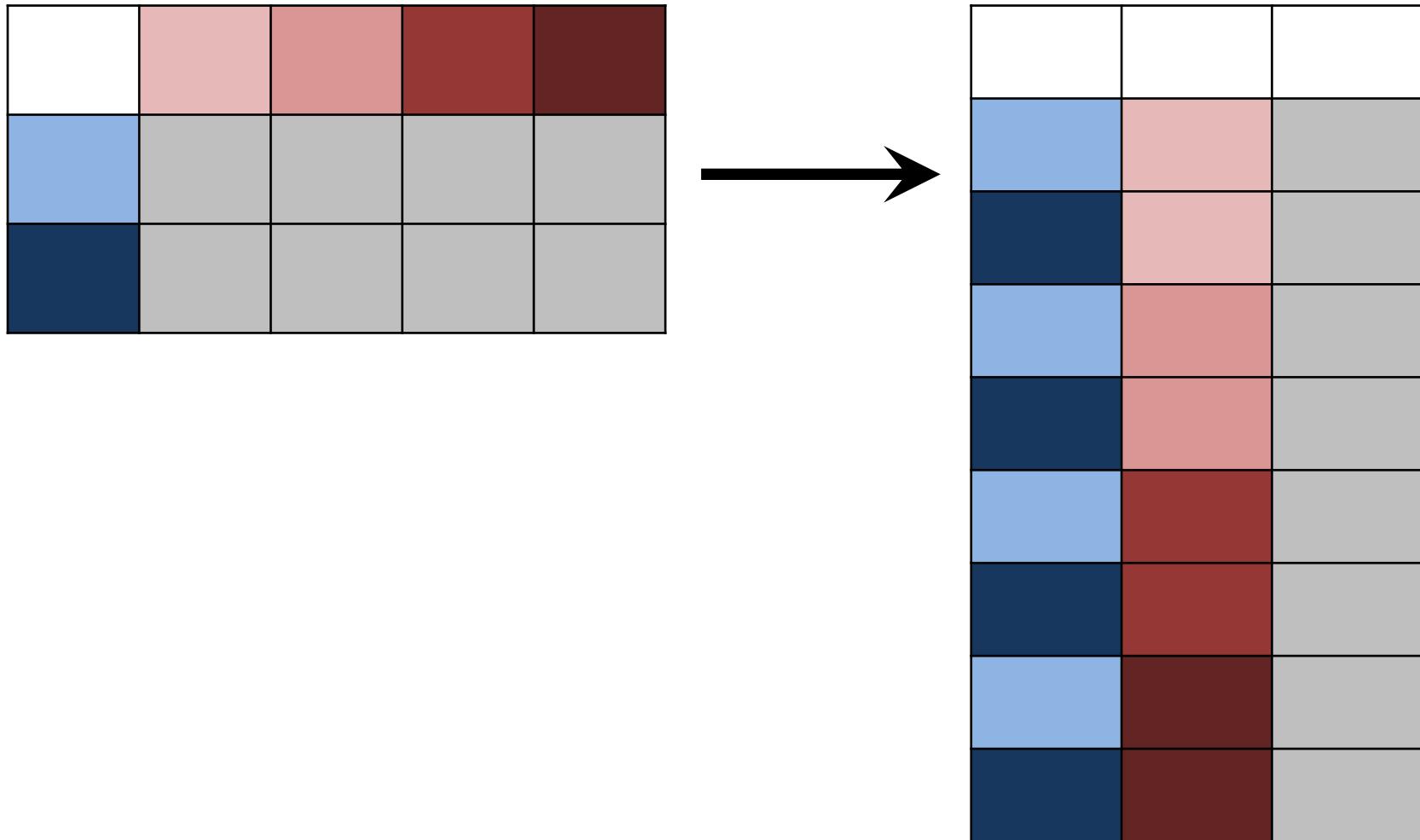
patient	treatment	outcome
1	drug	survived
2	drug	died
3	drug	survived
4	placebo	died
•	•	•

Making data sets longer or wider

We'll be discussing two functions:

- `pivot_longer()` — make a wide table long
- `pivot_wider()` — make a long table wide

pivot_longer()



pivot_longer()




```
data %>%  
  pivot_longer(cols, names_to = "A", values_to = "B")
```

pivot_longer()



columns




```
data %>%  
  pivot_longer(cols, names_to = "A", values_to = "B")
```

pivot_longer()



A


```
data %>%  
  pivot_longer(cols, names_to = "A", values_to = "B")
```

pivot_longer()




```
data %>%  
  pivot_longer(cols, names_to = "A", values_to = "B")
```

B

Example: Let's recreate the sitka data from a wide table

```
> head(sitka_wide)
  tree treat t152 t174 t201 t227 t258
1    1 ozone 4.51 4.98 5.41 5.90 6.15
2    2 ozone 4.24 4.20 4.68 4.92 4.96
3    3 ozone 3.98 4.36 4.79 4.99 5.03
4    4 ozone 4.36 4.77 5.10 5.30 5.36
5    5 ozone 4.34 4.95 5.42 5.97 6.28
6    6 ozone 4.59 5.08 5.36 5.76 6.00
```

Example: Let's recreate the sitka data from a wide table

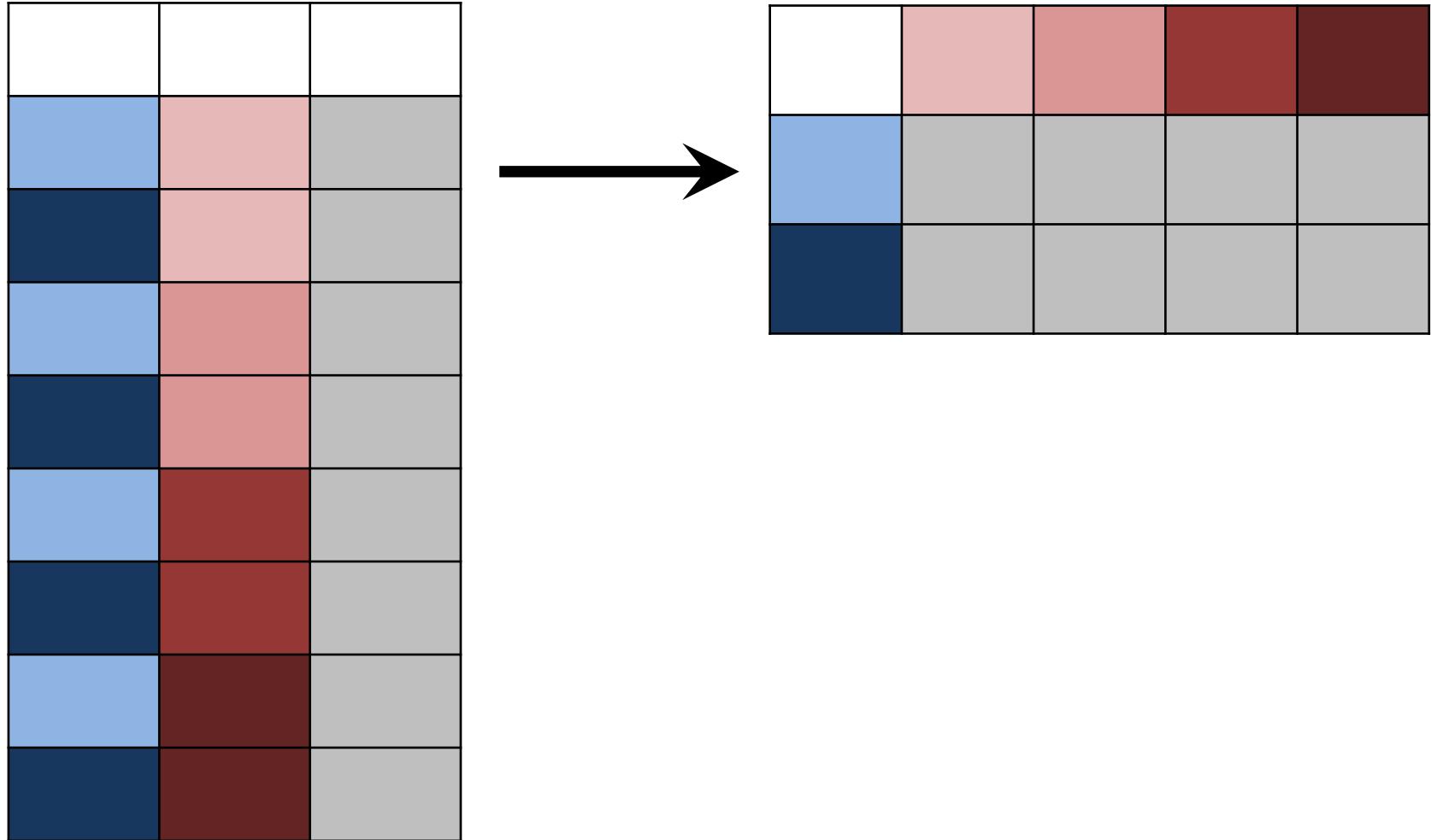
```
> head(sitka_wide)
   tree treat t152 t174 t201 t227 t258
1     1 ozone 4.51 4.98 5.41 5.90 6.15
2     2 ozone 4.24 4.20 4.68 4.92 4.96
3     3 ozone 3.98 4.36 4.79 4.99 5.03
4     4 ozone 4.36 4.77 5.10 5.30 5.36
5     5 ozone 4.34 4.95 5.42 5.97 6.28
6     6 ozone 4.59 5.08 5.36 5.76 6.00

sitka_wide %>%
  pivot_longer(
    t152:t258, names_to = "time", values_to = "size"
  )
```

Example: Let's recreate the sitka data from a wide table

```
> sitka_wide %>%
  pivot_longer(
    t152:t258, names_to = "time", values_to = "size"
  )
# A tibble: 395 x 4
  tree treat time   size
  <int> <fct> <chr> <dbl>
1     1 ozone t152   4.51
2     1 ozone t174   4.98
3     1 ozone t201   5.41
4     1 ozone t227   5.9 
5     1 ozone t258   6.15
6     2 ozone t152   4.24
7     2 ozone t174   4.2 
8     2 ozone t201   4.68
9     2 ozone t227   4.92
10    2 ozone t258   4.96
```

pivot_wider()



pivot_wider()

Blue	Pink	Grey
Dark Blue	Pink	Grey
Blue	Pink	Grey
Dark Blue	Pink	Grey
data %>% pivot_wider(names_from = "A", values_from = "B")		
Blue	Brown	Grey
Blue	Brown	Grey
Dark Blue	Brown	Grey



	Pink	Pink	Brown	Brown
Blue				
Dark Blue				
Blue				
Dark Blue				

```
data %>%  
pivot_wider(names_from = "A", values_from = "B")
```

pivot_wider()

A

	A	
	B	
1	blue	pink
2	dark blue	pink
3	blue	pink
4	dark blue	pink
5	blue	dark red
6	dark blue	dark red

data %>%
pivot_wider(names_from = "A", values_from = "B")

	A_1	A_2	A_3	A_4
	B			
1	blue			
2	dark blue			
3	blue			
4	dark blue			
5	blue		dark red	
6	dark blue		dark red	

```
data %>%  
pivot_wider(names_from = "A", values_from = "B")
```

pivot_wider()

B

blue	pink	grey		
dark blue	pink	grey		
blue	pink	grey		
dark blue	pink	grey		
blue	reddish-pink	grey	grey	grey
dark blue	reddish-pink	grey	grey	grey
blue	reddish-pink	grey	grey	grey
dark blue	reddish-brown	grey	grey	grey



	pink	pink	red	dark red
blue	grey	grey	grey	grey
dark blue	grey	grey	grey	grey
blue	grey	grey	grey	grey

```
data %>%
  pivot_wider(names_from = "A", values_from = "B")
```

Example: Let's turn the sitka data into a wide table

```
> head(sitka)
  size Time tree treat
1 4.51 152    1 ozone
2 4.98 174    1 ozone
3 5.41 201    1 ozone
4 5.90 227    1 ozone
5 6.15 258    1 ozone
6 4.24 152    2 ozone

sitka %>%
  pivot_wider(names_from="Time", values_from="size")
```

Example: Let's turn the Sitka data into a wide table

```
> sitka %>%
  pivot_wider(names_from="Time", values_from="size")

# A tibble: 79 x 7
  tree treat `152` `174` `201` `227` `258`
  <int> <fct> <dbl> <dbl> <dbl> <dbl> <dbl>
1     1 ozone   4.51   4.98   5.41   5.9    6.15
2     2 ozone   4.24   4.2    4.68   4.92   4.96
3     3 ozone   3.98   4.36   4.79   4.99   5.03
4     4 ozone   4.36   4.77   5.1    5.3    5.36
5     5 ozone   4.34   4.95   5.42   5.97   6.28
6     6 ozone   4.59   5.08   5.36   5.76   6
7     7 ozone   4.41   4.56   4.95   5.23   5.33
8     8 ozone   4.24   4.64   4.95   5.38   5.48
9     9 ozone   4.82   5.17   5.76   6.12   6.24
10    10 ozone  3.84   4.17   4.67   4.67   4.8
# ... with 69 more rows
```

Working with tidy data in R: tidyverse

Fundamental actions on data tables:

- make new columns — `mutate()`
- combine tables, adding columns — `left_join()`
- combine tables, adding rows — `bind_rows()`
- choose rows — `filter()`
- choose columns — `select()`
- arrange rows — `arrange()`
- calculate summary statistics — `summarize()`
- work on groups of data — `group_by()`