

# Introduction to R for Biologists

Tidyverse ecosystem & making data tidy with tidyr

# Day 1 Outline

1. How to get set up using R
2. How and why to use RStudio & R Markdown (.Rmd)
3. Basics of programming
  - Data types
  - Functions
  - Troubleshooting
4. Intro to the Tidyverse
  - Tidy vs untidy data
  - Tidyverse-specific functions

# Day 1 Outline

1. How to get set up using R
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4. Intro to the Tidyverse
  - Tidy vs untidy data
  - Tidyverse-specific functions

# 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
placebo	4	12

not tidy

# Example: Contingency table

	survived	died
drug	15	3
placebo	4	12

not tidy

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

# Example: Contingency table

	survived	died
drug	15	3
placebo	4	12

not tidy

tidy

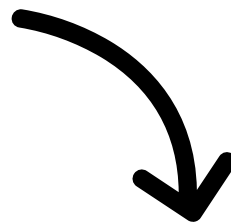
patient	treatment	outcome
1	drug	survived
2	drug	died
3	drug	survived
4	placebo	died
	⋮	



# tidyr library provides functions for transforming tables

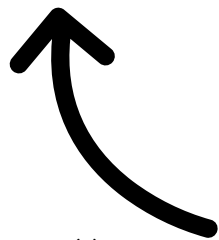
	survived	died
drug	15	3
placebo	4	12

`pivot_longer()`



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

`pivot_wider()`

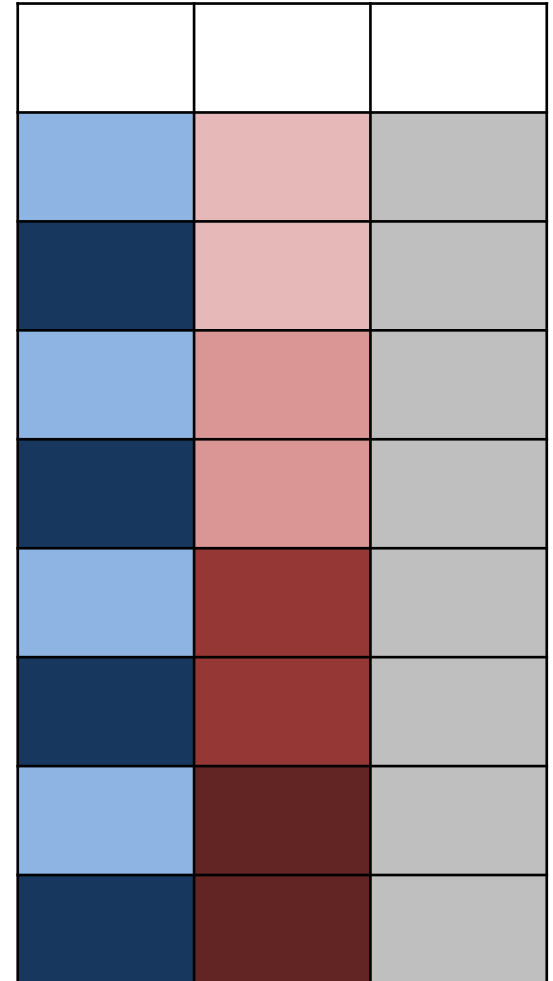
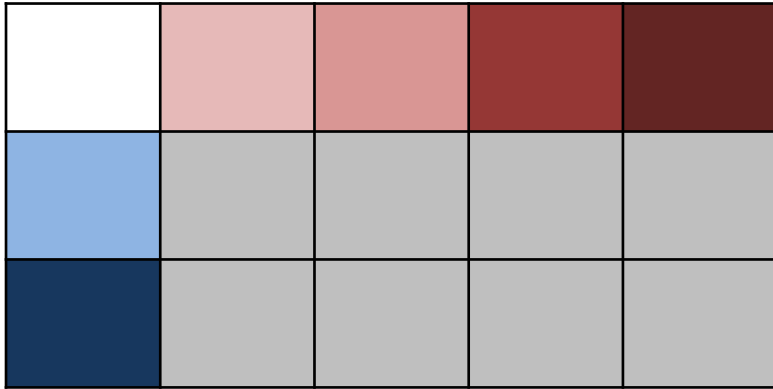


# Making data sets longer or wider

We'll be discussing two functions:

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

pivot\_longer()



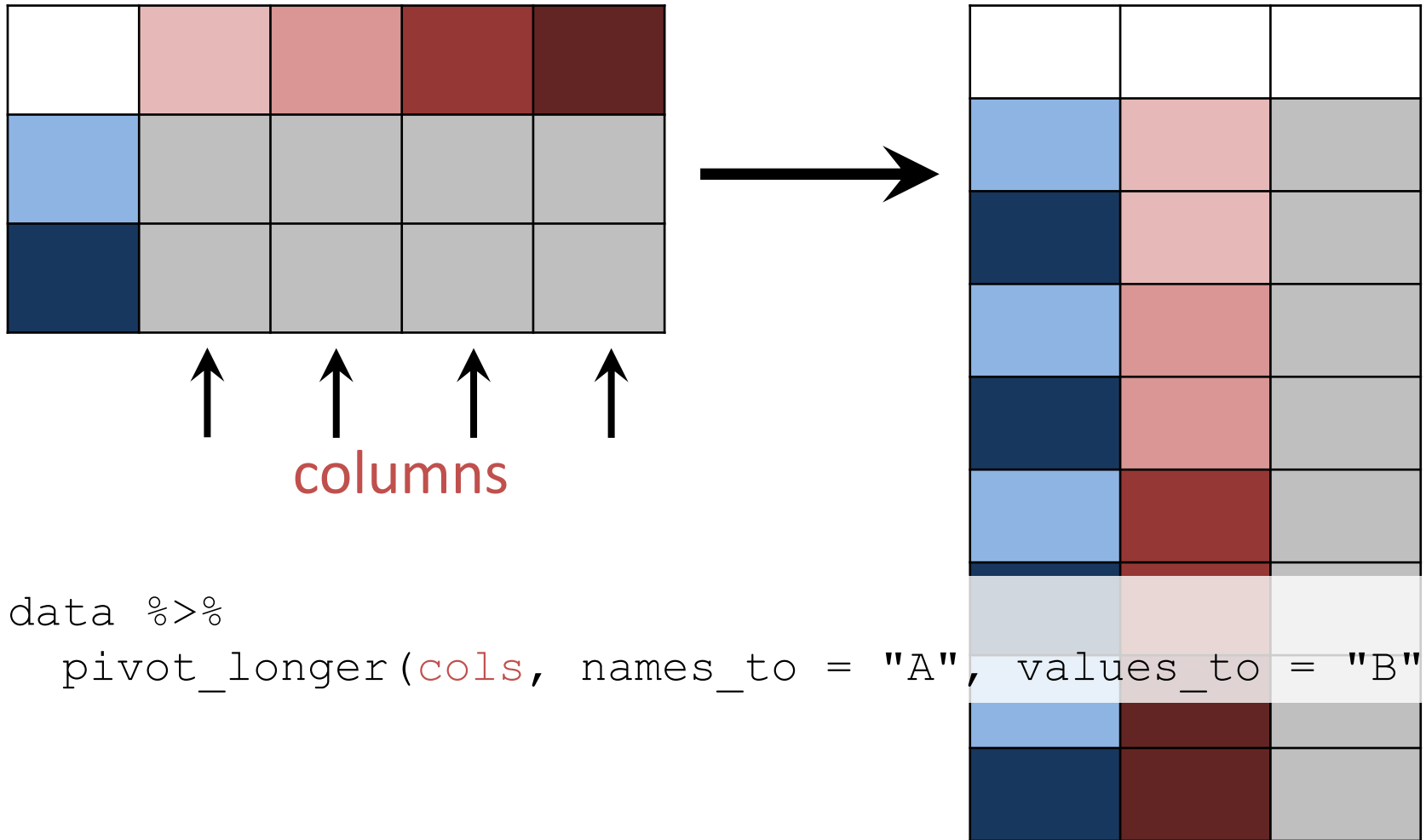
# `pivot_longer()`





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

# pivot\_longer()



# pivot\_longer()





A

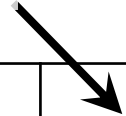
```
data %>%  
  pivot_longer(cols, names_to = "A", values_to = "B")
```

# `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

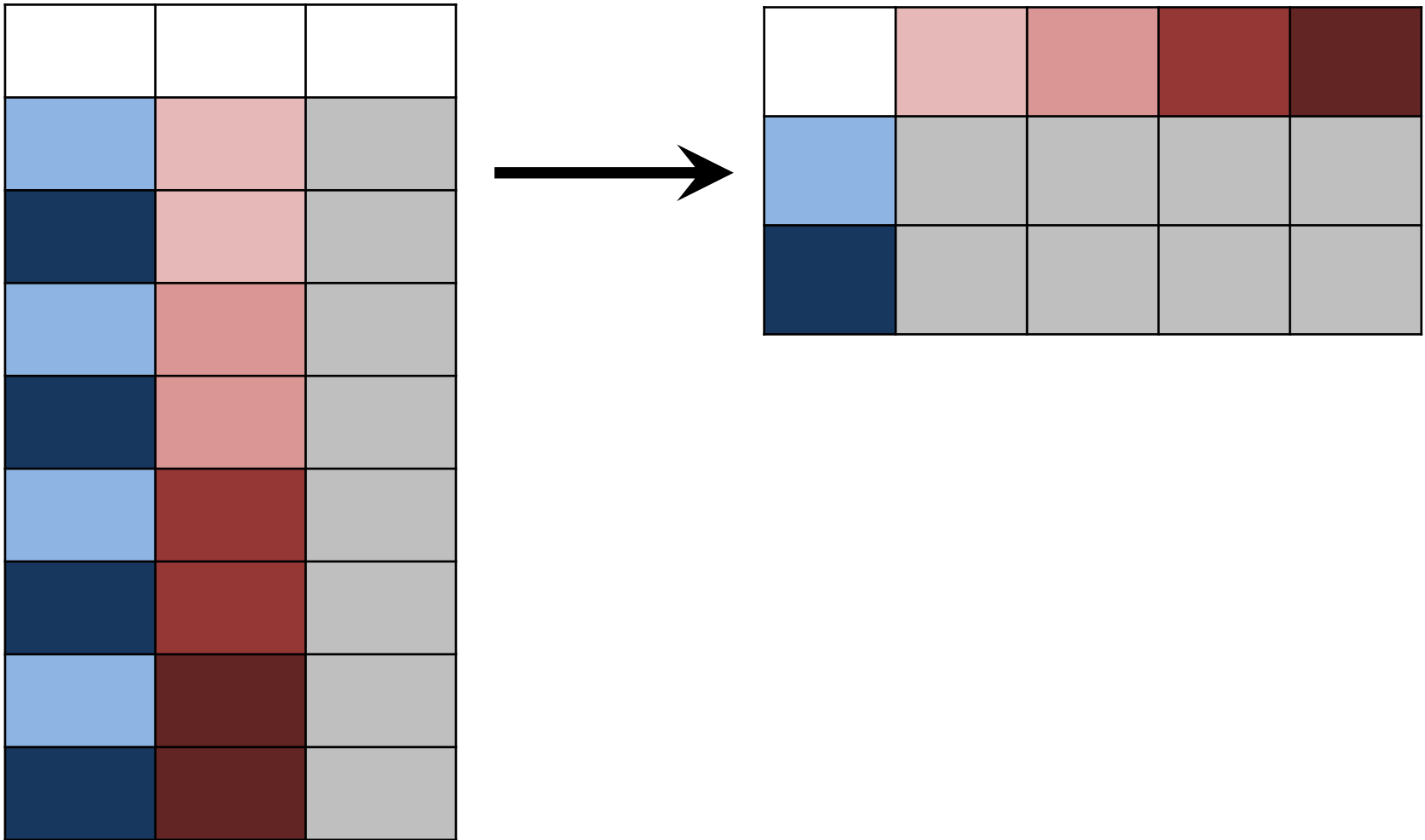
```
> head(sitka_wide)
  tree treat t152 t174 t201 t227 t258
1     1 ozone 4.51 4.98 5.41 5.90 6.15
2     2 ozone 4.24 4.20 4.68 4.92 4.96
3     3 ozone 3.98 4.36 4.79 4.99 5.03
4     4 ozone 4.36 4.77 5.10 5.30 5.36
5     5 ozone 4.34 4.95 5.42 5.97 6.28
6     6 ozone 4.59 5.08 5.36 5.76 6.00
```

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

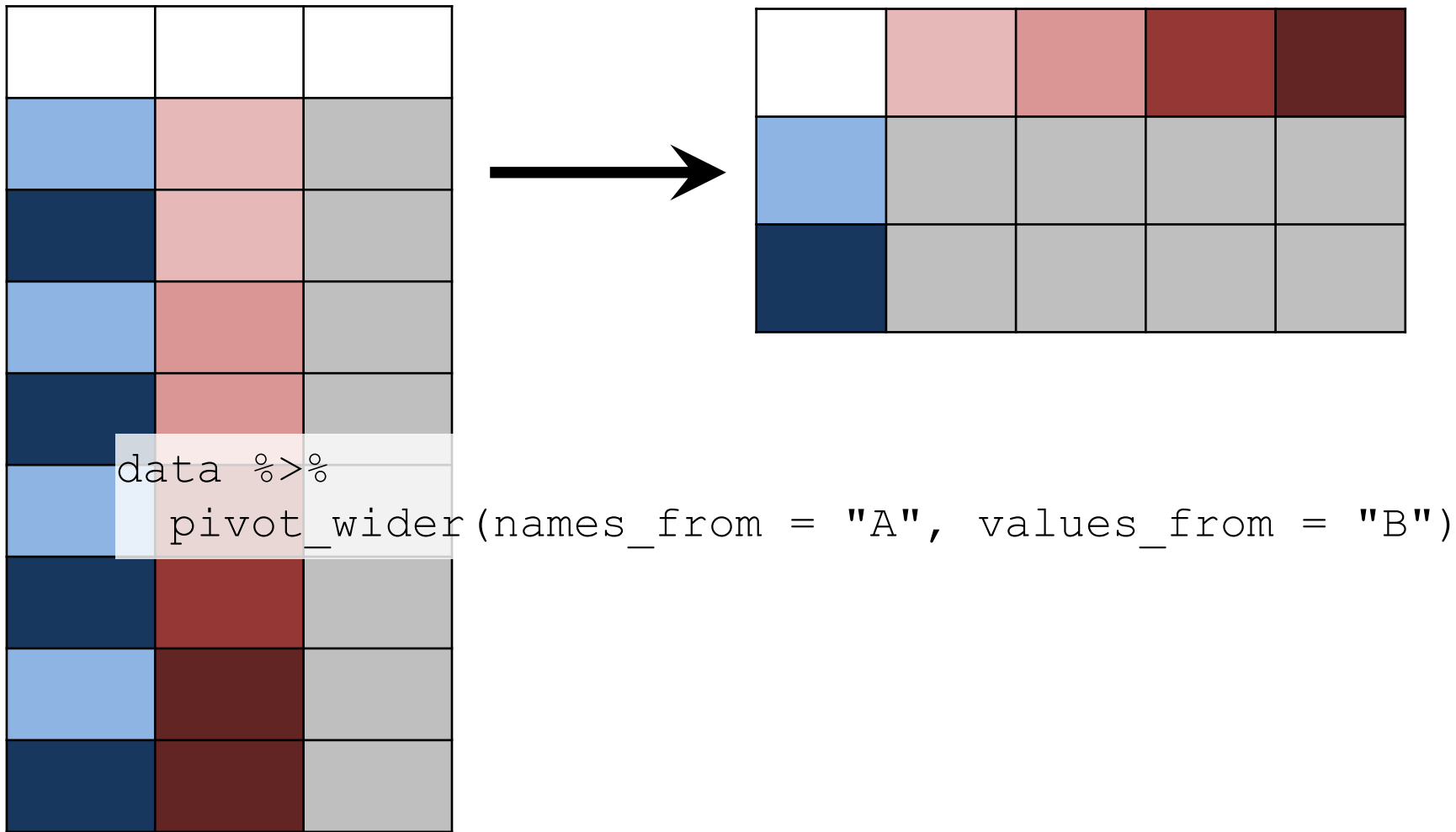
# Example: Let's recreate the sitka data from a wide table

```
> 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()`

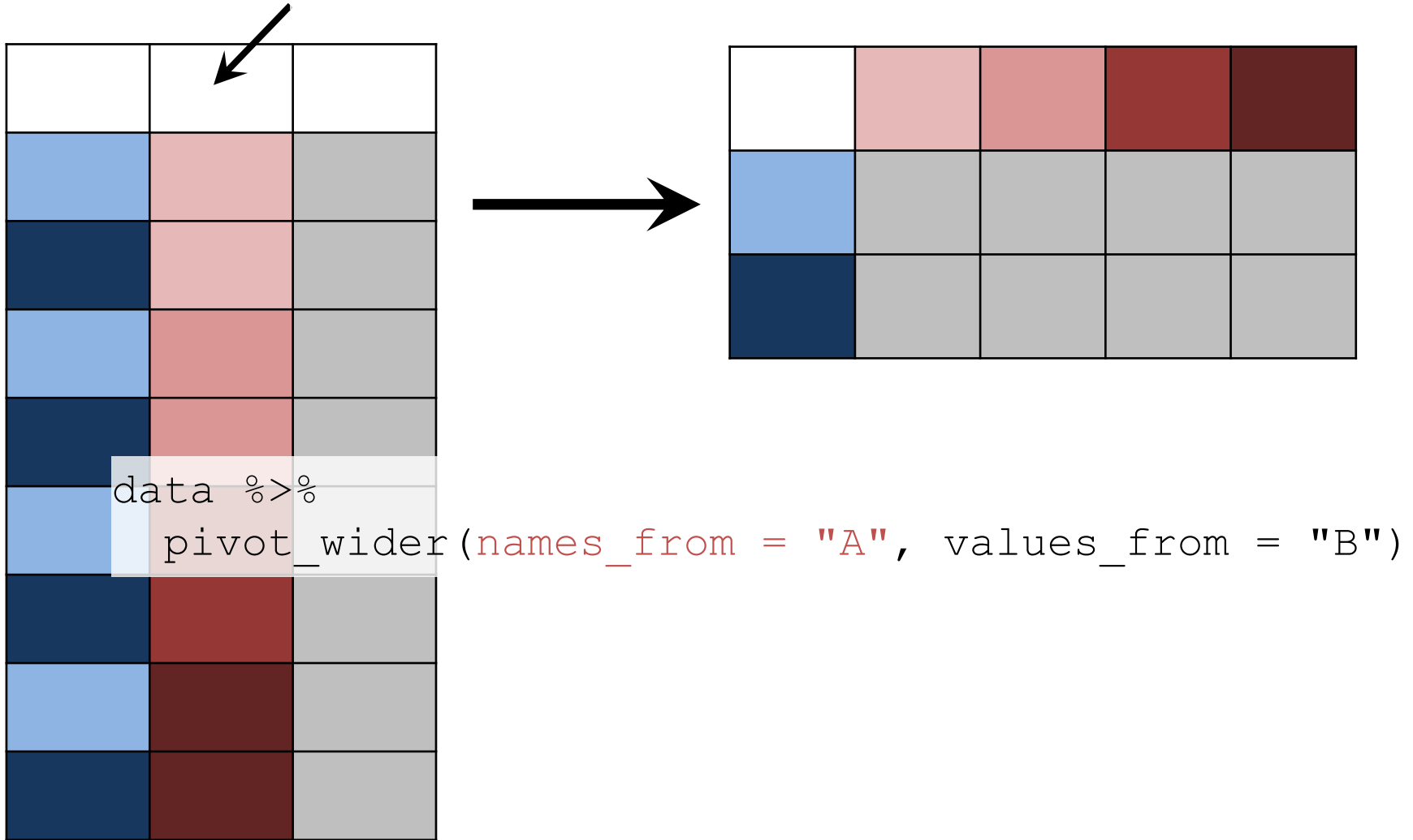


# `pivot_wider()`




# `pivot_wider()`

A




# `pivot_wider()`

B



light blue	light red	gray
dark blue	light red	gray
light blue	light red	gray
dark blue	light red	gray
light blue	dark red	gray
dark blue	dark red	gray
light blue	dark red	gray
dark blue	dark red	gray



	light red	light red	dark red	dark red
light blue	gray	gray	gray	gray
dark blue	gray	gray	gray	gray

```
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()`