Data Visualization - 5. Dplyr and Geoms

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Code Horizons

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Work with dplyr and ggplot

Load our libraries

```r
library(here)      # manage file paths
library(socviz)    # data and some useful functions
library(tidyverse) # your friend and mine
```

Tidyverse components

```
library(tidyverse)
Loading tidyverse: ggplot2
Loading tidyverse: tibble
Loading tidyverse: tidyr
Loading tidyverse: readr
Loading tidyverse: purrr
Loading tidyverse: dplyr
```

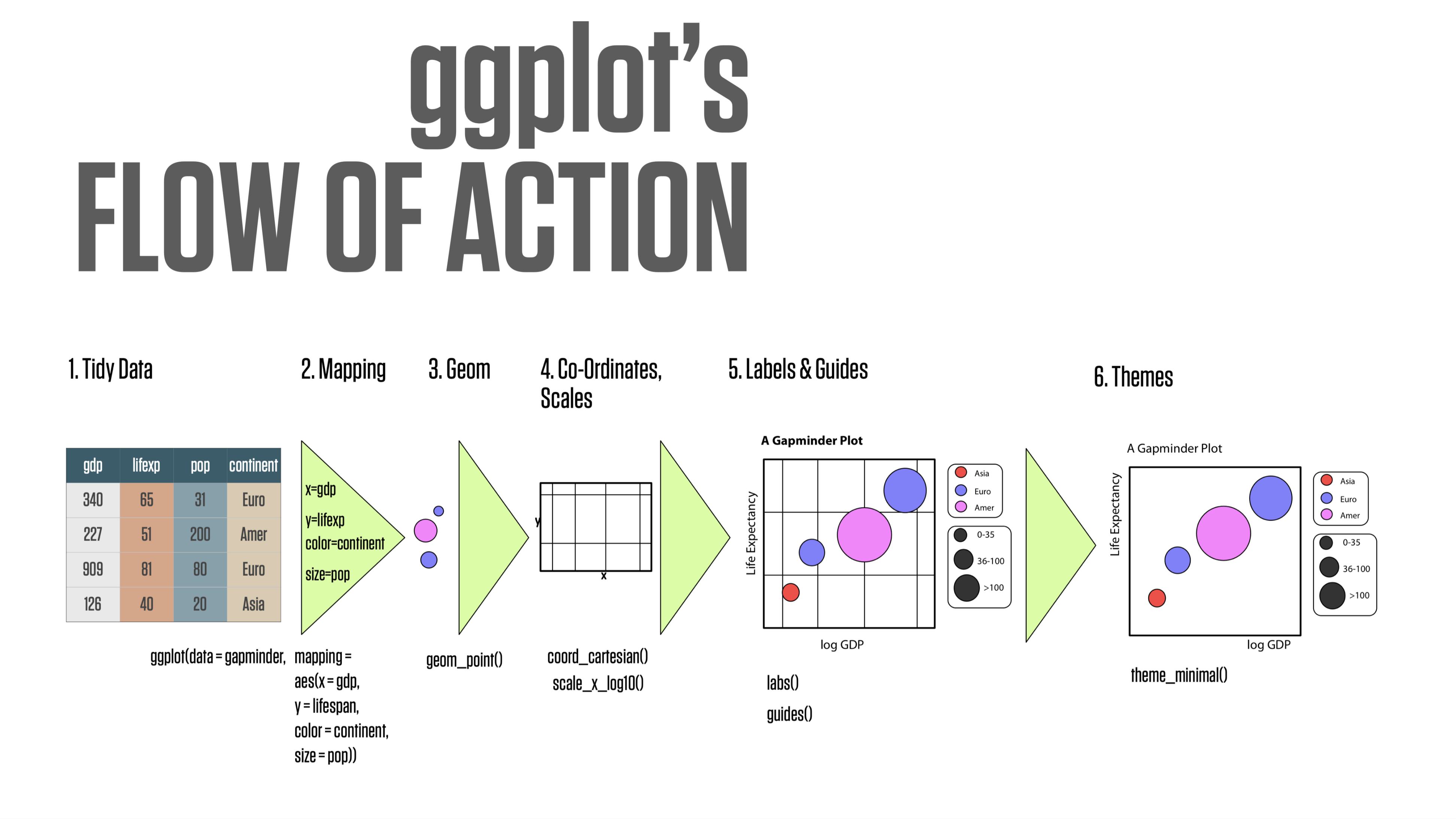
Load the package and ...

- **Draw graphs**
- **Nicer data tables**
- **Tidy your data**
- **Get data into R**
- **Fancy Iteration**
- **Action verbs for tables**

Other tidyverse components

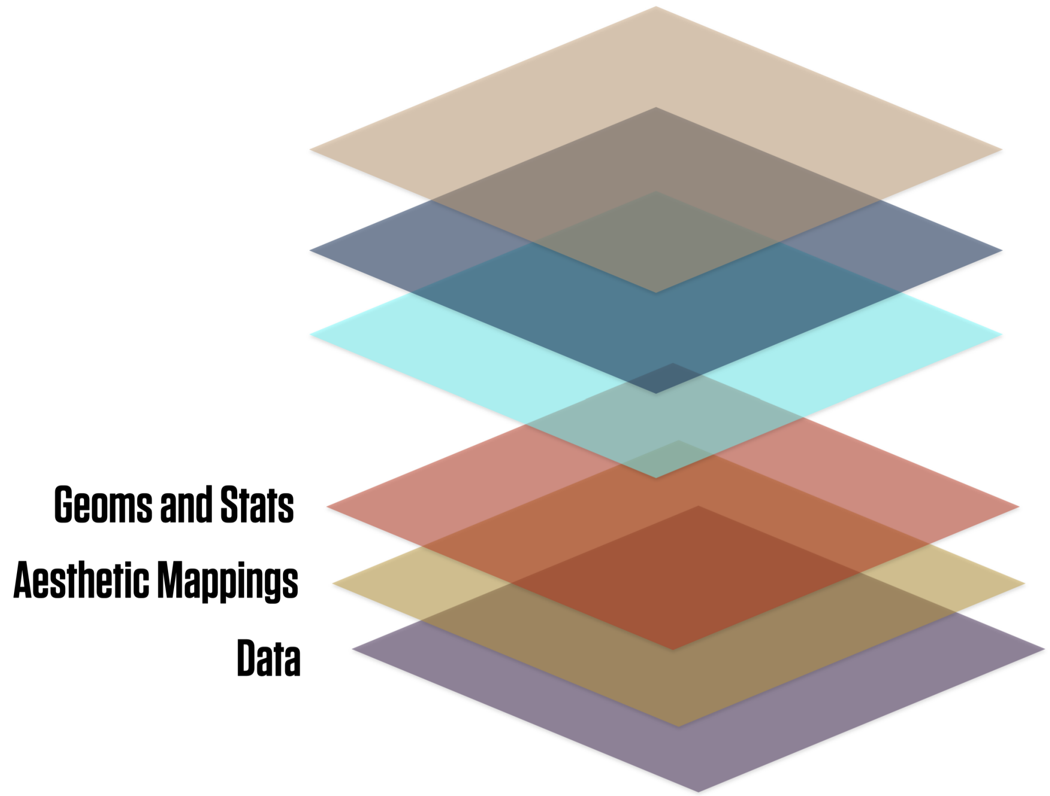
- `forcats` ◁ Deal with factors
- `haven` ◁ Import Stata, SPSS, etc
- `lubridate` ◁ Dates, Durations, Times
- `readxl` ◁ Import from spreadsheets
- `stringr` ◁ Strings and Regular Expressions
- `reprex` ◁ Make reproducible examples

Not all of these are attached when we do `library(tidyverse)`

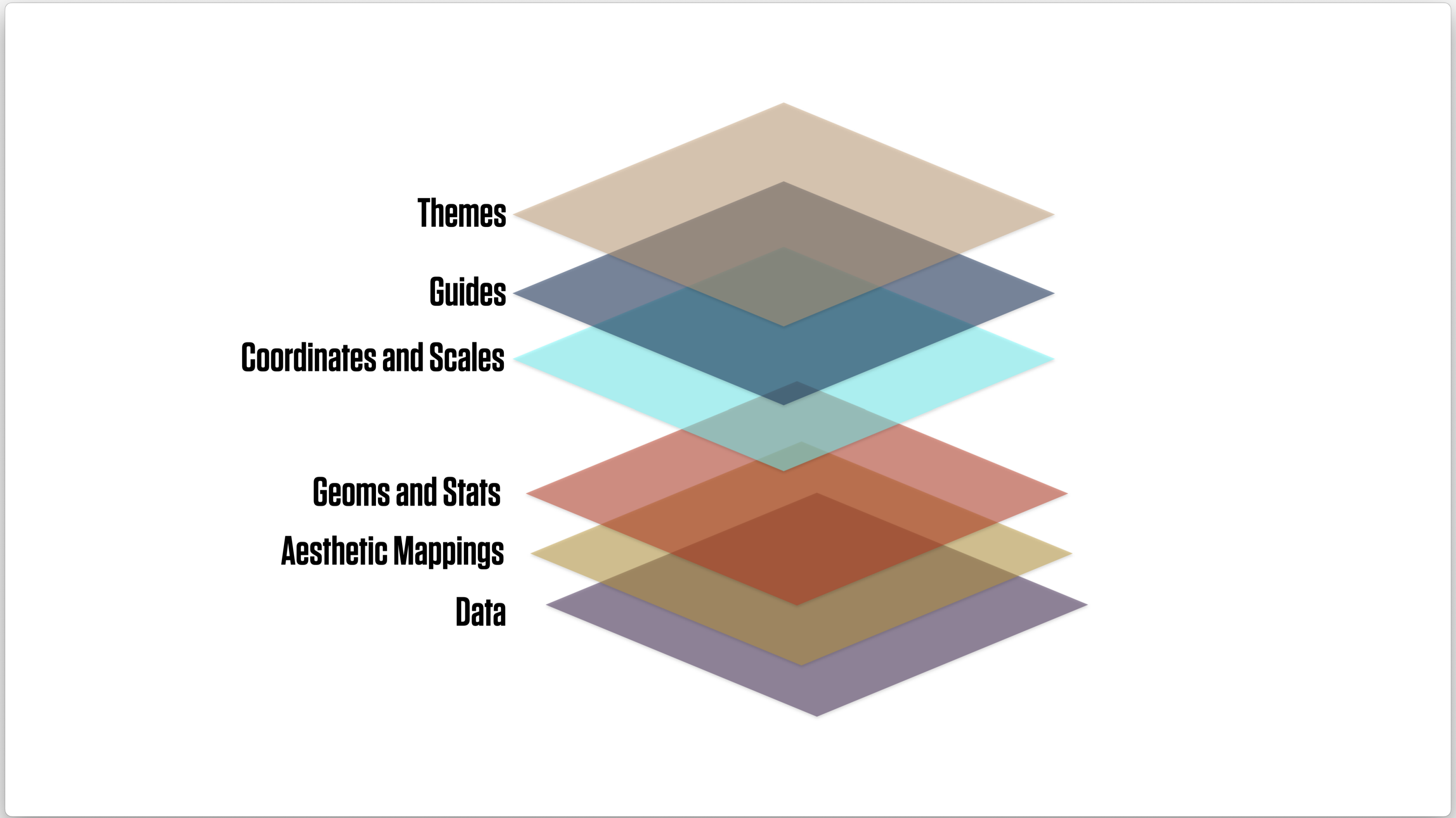


ggplot's flow of action

Thinking in terms of layers



Thinking in terms of layers



Thinking in terms of layers

Feeding data
to `ggplot`

Transform and summarize first. Then send your clean tables to ggplot.

Crosstabulation and beyond

U.S. General Social Survey data: `gss_sm`

```r
gss_sm
```

```
# A tibble: 2,867 × 32
    year    id ballot       age childs sibs   degree race  sex   region income16
   <dbl> <dbl> <labelled> <dbl>  <dbl> <labe> <fct>  <fct> <fct> <fct>  <fct>
 1  2016     1 1             47      3 2      Bache… White Male  New E… $170000…
 2  2016     2 2             61      0 3      High … White Male  New E… $50000 …
 3  2016     3 3             72      2 3      Bache… White Male  New E… $75000 …
 4  2016     4 1             43      4 3      High … White Fema… New E… $170000…
 5  2016     5 3             55      2 2      Gradu… White Fema… New E… $170000…
 6  2016     6 2             53      2 2      Junio… White Fema… New E… $60000 …
 7  2016     7 1             50      2 2      High … White Male  New E… $170000…
 8  2016     8 3             23      3 6      High … Other Fema… Middl… $30000 …
 9  2016     9 1             45      3 5      High … Black Male  Middl… $60000 …
10  2016    10 3             71      4 1      Junio… White Male  Middl… $60000 …
# ℹ 2,857 more rows
# ℹ 21 more variables: relig <fct>, marital <fct>, padeg <fct>, madeg <fct>,
#   partyid <fct>, polviews <fct>, happy <fct>, partners <fct>, grass <fct>,
#   zodiac <fct>, pres12 <labelled>, wtssall <dbl>, income_rc <fct>,
#   agegrp <fct>, ageq <fct>, siblings <fct>, kids <fct>, religion <fct>,
#   bigregion <fct>, partners_rc <fct>, obama <dbl>
```

We often want summary tables or graphs of data like this.

Two-way tables: Row percents

bigregion	Protestant	Catholic	Jewish	None	Other	Total
Northeast	32.4	33.3	5.5	23.0	5.7	100.0
Midwest	47.1	24.9	0.4	22.8	4.8	100.0
South	62.4	15.4	1.1	16.3	4.8	100.0
West	37.7	24.6	1.6	28.5	7.6	100.0

Two-way tables: Column percents

bigregion	Protestant	Catholic	Jewish	None	Other
Northeast	11.5	25.0	52.9	18.1	17.6
Midwest	23.7	26.5	5.9	25.4	20.8
South	47.4	24.7	21.6	27.5	31.4
West	17.4	23.9	19.6	29.1	30.2
Total	100.0	100.0	100.0	100.0	100.0

Two-way tables: Full marginals

bigregion	Protestant	Catholic	Jewish	None	Other
Northeast	5.5	5.7	0.9	3.9	1.0
Midwest	11.4	6.0	0.1	5.5	1.2
South	22.8	5.6	0.4	6.0	1.8
West	8.4	5.4	0.4	6.3	1.7

dplyr lets you work with tibbles

Remember, tibbles are tables of data where the columns can be of different types, such as numeric, logical, character, factor, etc.

We'll use dplyr to *transform* and *summarize* our data.

We'll use the pipe operator, ▷, to chain together sequences of actions on our tables.

dplyr’s core verbs

`dplyr` draws on the logic and language of **database queries**

Some **actions** to take on a single table

Group the data at the level we want, such as "*Religion by Region*" or "*Children by School*".

Subset either the rows or columns of or table—i.e. remove them before doing anything.

Mutate the data. That is, change something at the *current* level of grouping. Mutating adds new columns to the table, or changes the content of an existing column. It never changes the number of rows.

Summarize or aggregate the data. That is, make something new at a *higher* level of grouping. E.g., calculate means or counts by some grouping variable. This will generally result in a smaller, *summary* table. Usually this will have the same number of *rows* as there are *groups* being summarized.

For each action there's a function

Group using `group_by()`.

Subset has one action for rows and one for columns. We `filter()` rows and `select()` columns.

Mutate tables (i.e. add new columns, or re-make existing ones) using `mutate()`.

Summarize tables (i.e. perform aggregating calculations) using `summarize()`.

Group and Summarize

General Social Survey data: gss_sm

```r
## library(socviz) # if not loaded
gss_sm
```

```
# A tibble: 2,867 × 32
    year    id ballot       age childs sibs   degree race  sex   region income16
   <dbl> <dbl> <labelled> <dbl>  <dbl> <labe> <fct>  <fct> <fct> <fct>  <fct>
 1  2016     1 1             47      3 2      Bache… White Male  New E… $170000…
 2  2016     2 2             61      0 3      High … White Male  New E… $50000 …
 3  2016     3 3             72      2 3      Bache… White Male  New E… $75000 …
 4  2016     4 1             43      4 3      High … White Fema… New E… $170000…
 5  2016     5 3             55      2 2      Gradu… White Fema… New E… $170000…
 6  2016     6 2             53      2 2      Junio… White Fema… New E… $60000 …
 7  2016     7 1             50      2 2      High … White Male  New E… $170000…
 8  2016     8 3             23      3 6      High … Other Fema… Middl… $30000 …
 9  2016     9 1             45      3 5      High … Black Male  Middl… $60000 …
10  2016    10 3             71      4 1      Junio… White Male  Middl… $60000 …
# ℹ 2,857 more rows
# ℹ 21 more variables: relig <fct>, marital <fct>, padeg <fct>, madeg <fct>,
#   partyid <fct>, polviews <fct>, happy <fct>, partners <fct>, grass <fct>,
#   zodiac <fct>, pres12 <labelled>, wtssall <dbl>, income_rc <fct>,
#   agegrp <fct>, ageq <fct>, siblings <fct>, kids <fct>, religion <fct>,
#   bigregion <fct>, partners_rc <fct>, obama <dbl>
```

Notice how the tibble already tells us a lot.

Summarizing a Table

Here's what we're going to do:

1. Individual-Level GSS Data on Region and Religion

id	bigregion	religion
1014	Midwest	Protestant
1544	South	Protestant
665	Northeast	None
1618	South	None
2115	West	Catholic
417	South	Protestant
2045	West	Protestant
1863	Northeast	Other
1884	Midwest	Christian
1628	South	Protestant

2. Summary Count of Religious Preferences by Census Region

bigregion	religion	N
Northeast	Protestant	123
Northeast	Catholic	149
Northeast	Jewish	15
Northeast	None	97
Northeast	Christian	14
Northeast	Other	31

3. Percent Religious Preferences by Census Region

bigregion	religion	N	pct
Northeast	Protestant	123	28.3
Northeast	Catholic	149	34.3
Northeast	Jewish	15	3.4
Northeast	None	97	22.3
Northeast	Christian	14	3.2
Northeast	Other	31	7.1

Summarizing a Table

```
gss_sm |>
  select(id, bigregion, religion)
```

```
# A tibble: 2,867 × 3
      id bigregion religion
   <dbl> <fct>     <fct>
 1     1 Northeast None
 2     2 Northeast None
 3     3 Northeast Catholic
 4     4 Northeast Catholic
 5     5 Northeast None
 6     6 Northeast None
 7     7 Northeast None
 8     8 Northeast Catholic
 9     9 Northeast Protestant
10    10 Northeast None
# ℹ 2,857 more rows
```

We're just taking a look at the relevant columns here.

Group by *one* column or variable

```r
gss_sm |>
  group_by(bigregion)
```

```
# A tibble: 2,867 × 32
# Groups:   bigregion [4]
    year    id ballot       age childs sibs   degree race  sex   region income16
   <dbl> <dbl> <labelled> <dbl>  <dbl> <labe> <fct>  <fct> <fct> <fct>  <fct>
 1  2016     1 1             47      3 2      Bache… White Male  New E… $170000…
 2  2016     2 2             61      0 3      High … White Male  New E… $50000 …
 3  2016     3 3             72      2 3      Bache… White Male  New E… $75000 …
 4  2016     4 1             43      4 3      High … White Fema… New E… $170000…
 5  2016     5 3             55      2 2      Gradu… White Fema… New E… $170000…
 6  2016     6 2             53      2 2      Junio… White Fema… New E… $60000 …
 7  2016     7 1             50      2 2      High … White Male  New E… $170000…
 8  2016     8 3             23      3 6      High … Other Fema… Middl… $30000 …
 9  2016     9 1             45      3 5      High … Black Male  Middl… $60000 …
10  2016    10 3             71      4 1      Junio… White Male  Middl… $60000 …
# ℹ 2,857 more rows
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#   partyid <fct>, polviews <fct>, happy <fct>, partners <fct>, grass <fct>,
#   zodiac <fct>, pres12 <labelled>, wtssall <dbl>, income_rc <fct>,
#   agegrp <fct>, ageq <fct>, siblings <fct>, kids <fct>, religion <fct>,
```

Grouping just changes the logical structure of the tibble.

Group and summarize by *one* column

```
gss_sm
```

```
# A tibble: 2,867 × 32
    year    id ballot       age childs sibs   degree race  sex   region income16
   <dbl> <dbl> <labelled> <dbl>  <dbl> <labe> <fct>  <fct> <fct> <fct>  <fct>
 1  2016     1 1             47      3 2      Bache… White Male  New E… $170000…
 2  2016     2 2             61      0 3      High … White Male  New E… $50000 …
 3  2016     3 3             72      2 3      Bache… White Male  New E… $75000 …
 4  2016     4 1             43      4 3      High … White Fema… New E… $170000…
 5  2016     5 3             55      2 2      Gradu… White Fema… New E… $170000…
 6  2016     6 2             53      2 2      Junio… White Fema… New E… $60000 …
 7  2016     7 1             50      2 2      High … White Male  New E… $170000…
 8  2016     8 3             23      3 6      High … Other Fema… Middl… $30000 …
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# ℹ 2,857 more rows
# ℹ 21 more variables: relig <fct>, marital <fct>, padeg <fct>, madeg <fct>,
#   partyid <fct>, polviews <fct>, happy <fct>, partners <fct>, grass <fct>,
#   zodiac <fct>, pres12 <labelled>, wtssall <dbl>, income_rc <fct>,
#   agegrp <fct>, ageq <fct>, siblings <fct>, kids <fct>, religion <fct>,
#   bigregion <fct>, partners_rc <fct>, obama <dbl>
```

Group and summarize by *one* column

```r
gss_sm |>
  group_by(bigregion)
```

```
# A tibble: 2,867 × 32
# Groups:   bigregion [4]
    year    id ballot       age childs sibs   degree race  sex   region income16
   <dbl> <dbl> <labelled> <dbl>  <dbl> <labe> <fct>  <fct> <fct> <fct>  <fct>
 1  2016     1 1             47      3 2      Bache… White Male  New E… $170000…
 2  2016     2 2             61      0 3      High … White Male  New E… $50000 …
 3  2016     3 3             72      2 3      Bache… White Male  New E… $75000 …
 4  2016     4 1             43      4 3      High … White Fema… New E… $170000…
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10  2016    10 3             71      4 1      Junio… White Male  Middl… $60000 …
# ℹ 2,857 more rows
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#   partyid <fct>, polviews <fct>, happy <fct>, partners <fct>, grass <fct>,
#   zodiac <fct>, pres12 <labelled>, wtssall <dbl>, income_rc <fct>,
#   agegrp <fct>, ageq <fct>, siblings <fct>, kids <fct>, religion <fct>,
#   bigregion <fct>, partners_rc <fct>, obama <dbl>
```

Group and summarize by *one* column

```r
gss_sm |>
  group_by(bigregion) |>
  summarize(total = n())
```

```
# A tibble: 4 × 2
  bigregion total
  <fct>     <int>
1 Northeast   488
2 Midwest     695
3 South      1052
4 West        632
```

The function `n()` counts up the rows within each group.

All the other columns are dropped in the summary operation

Your original `gss_sm` table is untouched

Group and summarize by *two* columns

```r
gss_sm
```

```
# A tibble: 2,867 × 32
    year    id ballot       age childs sibs   degree race  sex   region income16
   <dbl> <dbl> <labelled> <dbl>  <dbl> <labe> <fct>  <fct> <fct> <fct>  <fct>
 1  2016     1 1             47      3 2      Bache… White Male  New E… $170000…
 2  2016     2 2             61      0 3      High … White Male  New E… $50000 …
 3  2016     3 3             72      2 3      Bache… White Male  New E… $75000 …
 4  2016     4 1             43      4 3      High … White Fema… New E… $170000…
 5  2016     5 3             55      2 2      Gradu… White Fema… New E… $170000…
 6  2016     6 2             53      2 2      Junio… White Fema… New E… $60000 …
 7  2016     7 1             50      2 2      High … White Male  New E… $170000…
 8  2016     8 3             23      3 6      High … Other Fema… Middl… $30000 …
 9  2016     9 1             45      3 5      High … Black Male  Middl… $60000 …
10  2016    10 3             71      4 1      Junio… White Male  Middl… $60000 …
# ℹ 2,857 more rows
# ℹ 21 more variables: relig <fct>, marital <fct>, padeg <fct>, madeg <fct>,
#   partyid <fct>, polviews <fct>, happy <fct>, partners <fct>, grass <fct>,
#   zodiac <fct>, pres12 <labelled>, wtssall <dbl>, income_rc <fct>,
#   agegrp <fct>, ageq <fct>, siblings <fct>, kids <fct>, religion <fct>,
#   bigregion <fct>, partners_rc <fct>, obama <dbl>
```

Group and summarize by *two* columns

```
gss_sm |>
  group_by(bigregion, religion)
```

```
# A tibble: 2,867 × 32
# Groups:   bigregion, religion [24]
    year    id ballot       age childs sibs   degree race  sex   region income16
   <dbl> <dbl> <labelled> <dbl>  <dbl> <labe> <fct>  <fct> <fct> <fct>  <fct>
 1  2016     1 1             47      3 2      Bache… White Male  New E… $170000…
 2  2016     2 2             61      0 3      High … White Male  New E… $50000 …
 3  2016     3 3             72      2 3      Bache… White Male  New E… $75000 …
 4  2016     4 1             43      4 3      High … White Fema… New E… $170000…
 5  2016     5 3             55      2 2      Gradu… White Fema… New E… $170000…
 6  2016     6 2             53      2 2      Junio… White Fema… New E… $60000 …
 7  2016     7 1             50      2 2      High … White Male  New E… $170000…
 8  2016     8 3             23      3 6      High … Other Fema… Middl… $30000 …
 9  2016     9 1             45      3 5      High … Black Male  Middl… $60000 …
10  2016    10 3             71      4 1      Junio… White Male  Middl… $60000 …
# ℹ 2,857 more rows
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#   partyid <fct>, polviews <fct>, happy <fct>, partners <fct>, grass <fct>,
#   zodiac <fct>, pres12 <labelled>, wtssall <dbl>, income_rc <fct>,
#   agegrp <fct>, ageq <fct>, siblings <fct>, kids <fct>, religion <fct>,
#   bigregion <fct>, partners_rc <fct>, obama <dbl>
```

Group and summarize by *two* columns

```r
gss_sm |>
  group_by(bigregion, religion) |>
  summarize(total = n())
```

```
# A tibble: 24 × 3
# Groups:   bigregion [4]
   bigregion religion   total
   <fct>     <fct>      <int>
 1 Northeast Protestant   158
 2 Northeast Catholic     162
 3 Northeast Jewish        27
 4 Northeast None         112
 5 Northeast Other         28
 6 Northeast <NA>           1
 7 Midwest   Protestant   325
 8 Midwest   Catholic     172
 9 Midwest   Jewish         3
10 Midwest   None         157
# ℹ 14 more rows
```

The function `n()` counts up the rows within the *innermost* (i.e. the rightmost) group.

Calculate frequencies

```r
gss_sm
```

```
# A tibble: 2,867 × 32
    year    id ballot       age childs sibs   degree race  sex   region income16
   <dbl> <dbl> <labelled> <dbl>  <dbl> <labe> <fct>  <fct> <fct> <fct>  <fct>
 1  2016     1 1             47      3 2      Bache… White Male  New E… $170000…
 2  2016     2 2             61      0 3      High … White Male  New E… $50000 …
 3  2016     3 3             72      2 3      Bache… White Male  New E… $75000 …
 4  2016     4 1             43      4 3      High … White Fema… New E… $170000…
 5  2016     5 3             55      2 2      Gradu… White Fema… New E… $170000…
 6  2016     6 2             53      2 2      Junio… White Fema… New E… $60000 …
 7  2016     7 1             50      2 2      High … White Male  New E… $170000…
 8  2016     8 3             23      3 6      High … Other Fema… Middl… $30000 …
 9  2016     9 1             45      3 5      High … Black Male  Middl… $60000 …
10  2016    10 3             71      4 1      Junio… White Male  Middl… $60000 …
# ℹ 2,857 more rows
# ℹ 21 more variables: relig <fct>, marital <fct>, padeg <fct>, madeg <fct>,
#   partyid <fct>, polviews <fct>, happy <fct>, partners <fct>, grass <fct>,
#   zodiac <fct>, pres12 <labelled>, wtssall <dbl>, income_rc <fct>,
#   agegrp <fct>, ageq <fct>, siblings <fct>, kids <fct>, religion <fct>,
#   bigregion <fct>, partners_rc <fct>, obama <dbl>
```

Calculate frequencies

```
gss_sm ▷
  group_by(bigregion, religion)
```

```
# A tibble: 2,867 × 32
# Groups:   bigregion, religion [24]
    year    id ballot       age childs sibs   degree race  sex   region income16
   <dbl> <dbl> <labelled> <dbl>  <dbl> <labe> <fct>  <fct> <fct> <fct>  <fct>
 1  2016     1 1             47      3 2      Bache… White Male  New E… $170000…
 2  2016     2 2             61      0 3      High … White Male  New E… $50000 …
 3  2016     3 3             72      2 3      Bache… White Male  New E… $75000 …
 4  2016     4 1             43      4 3      High … White Fema… New E… $170000…
 5  2016     5 3             55      2 2      Gradu… White Fema… New E… $170000…
 6  2016     6 2             53      2 2      Junio… White Fema… New E… $60000 …
 7  2016     7 1             50      2 2      High … White Male  New E… $170000…
 8  2016     8 3             23      3 6      High … Other Fema… Middl… $30000 …
 9  2016     9 1             45      3 5      High … Black Male  Middl… $60000 …
10  2016    10 3             71      4 1      Junio… White Male  Middl… $60000 …
# ℹ 2,857 more rows
# ℹ 21 more variables: relig <fct>, marital <fct>, padeg <fct>, madeg <fct>,
#   partyid <fct>, polviews <fct>, happy <fct>, partners <fct>, grass <fct>,
#   zodiac <fct>, pres12 <labelled>, wtssall <dbl>, income_rc <fct>,
#   agegrp <fct>, ageq <fct>, siblings <fct>, kids <fct>, religion <fct>,
#   bigregion <fct>, partners_rc <fct>, obama <dbl>
```

Calculate frequencies

```r
gss_sm |>
  group_by(bigregion, religion) |>
  summarize(total = n())
```

```
# A tibble: 24 × 3
# Groups:   bigregion [4]
   bigregion religion   total
   <fct>     <fct>      <int>
 1 Northeast Protestant   158
 2 Northeast Catholic     162
 3 Northeast Jewish        27
 4 Northeast None         112
 5 Northeast Other         28
 6 Northeast <NA>           1
 7 Midwest   Protestant   325
 8 Midwest   Catholic     172
 9 Midwest   Jewish         3
10 Midwest   None         157
# ℹ 14 more rows
```

Calculate frequencies

```r
gss_sm |>
  group_by(bigregion, religion) |>
  summarize(total = n()) |>
  mutate(freq = total / sum(total),
         pct = round((freq*100), 1))
```

```
# A tibble: 24 × 5
# Groups:   bigregion [4]
   bigregion religion   total    freq   pct
   <fct>     <fct>      <int>   <dbl> <dbl>
 1 Northeast Protestant   158 0.324    32.4
 2 Northeast Catholic     162 0.332    33.2
 3 Northeast Jewish        27 0.0553    5.5
 4 Northeast None         112 0.230    23
 5 Northeast Other         28 0.0574    5.7
 6 Northeast <NA>           1 0.00205   0.2
 7 Midwest   Protestant   325 0.468    46.8
 8 Midwest   Catholic     172 0.247    24.7
 9 Midwest   Jewish         3 0.00432   0.4
10 Midwest   None         157 0.226    22.6
# ℹ 14 more rows
```

The function `n()` counts up the rows

Which rows? The ones fed down the pipeline

The *innermost* (i.e. the rightmost) group.

Pipelines carry assumptions forward

```r
gss_sm |>
  group_by(bigregion, religion) |> #<<
  summarize(total = n()) |>
  mutate(freq = total / sum(total),
           pct = round((freq*100), 1))
```

```
# A tibble: 24 × 5
# Groups:   bigregion [4]
   bigregion religion   total    freq   pct
   <fct>     <fct>      <int>   <dbl> <dbl>
 1 Northeast Protestant   158 0.324    32.4
 2 Northeast Catholic     162 0.332    33.2
 3 Northeast Jewish        27 0.0553    5.5
 4 Northeast None         112 0.230    23  
 5 Northeast Other         28 0.0574    5.7
 6 Northeast <NA>           1 0.00205   0.2
 7 Midwest   Protestant   325 0.468    46.8
 8 Midwest   Catholic     172 0.247    24.7
 9 Midwest   Jewish         3 0.00432   0.4
10 Midwest   None         157 0.226    22.6
# ℹ 14 more rows
```

Groups are carried forward till summarized or explicitly ungrouped

Summary calculations are done on the innermost group, which then “disappears”.

Pipelines carry assumptions forward

```r
gss_sm |>
  group_by(bigregion, religion) |>
  summarize(total = n()) |>
  mutate(freq = total / sum(total),
         pct = round((freq*100), 1)) #<<
```

```
# A tibble: 24 × 5
# Groups:   bigregion [4]
   bigregion religion   total    freq   pct
   <fct>     <fct>      <int>   <dbl> <dbl>
 1 Northeast Protestant   158 0.324    32.4
 2 Northeast Catholic     162 0.332    33.2
 3 Northeast Jewish        27 0.0553    5.5
 4 Northeast None         112 0.230    23  
 5 Northeast Other         28 0.0574    5.7
 6 Northeast <NA>           1 0.00205   0.2
 7 Midwest   Protestant   325 0.468    46.8
 8 Midwest   Catholic     172 0.247    24.7
 9 Midwest   Jewish         3 0.00432   0.4
10 Midwest   None         157 0.226    22.6
# ℹ 14 more rows
```

`mutate()` is quite clever. See how we can immediately use `freq`, even though we are creating it in the same `mutate()` expression.

Convenience functions

```r
gss_sm |>
  group_by(bigregion, religion) |> #<<
  summarize(total = n()) |> #<<
  mutate(freq = total / sum(total),
           pct = round((freq*100), 1))
```

```
# A tibble: 24 × 5
# Groups:   bigregion [4]
   bigregion religion   total    freq   pct
   <fct>     <fct>      <int>   <dbl> <dbl>
 1 Northeast Protestant   158 0.324    32.4
 2 Northeast Catholic     162 0.332    33.2
 3 Northeast Jewish        27 0.0553    5.5
 4 Northeast None         112 0.230    23
 5 Northeast Other         28 0.0574    5.7
 6 Northeast <NA>           1 0.00205   0.2
 7 Midwest   Protestant   325 0.468    46.8
 8 Midwest   Catholic     172 0.247    24.7
 9 Midwest   Jewish         3 0.00432   0.4
10 Midwest   None         157 0.226    22.6
# ℹ 14 more rows
```

We're going to be doing this `group_by()` … `n()` step a lot. Some shorthand for it would be useful.

Three options for counting up rows

Use `n()`

```
gss_sm ▷
  group_by(bigregion, religion) ▷ #<<
  summarize(n = n()) #<<
```

```
# A tibble: 24 × 3
# Groups:   bigregion [4]
   bigregion religion       n
   <fct>     <fct>      <int>
 1 Northeast Protestant   158
 2 Northeast Catholic     162
 3 Northeast Jewish        27
 4 Northeast None         112
 5 Northeast Other         28
 6 Northeast <NA>           1
 7 Midwest   Protestant   325
 8 Midwest   Catholic     172
 9 Midwest   Jewish         3
10 Midwest   None         157
# ℹ 14 more rows
```

Group it yourself; result is grouped.

Use `tally()`

```
gss_sm ▷
  group_by(bigregion, religion) ▷
  tally() #<<
```

```
# A tibble: 24 × 3
# Groups:   bigregion [4]
   bigregion religion       n
   <fct>     <fct>      <int>
 1 Northeast Protestant   158
 2 Northeast Catholic     162
 3 Northeast Jewish        27
 4 Northeast None         112
 5 Northeast Other         28
 6 Northeast <NA>           1
 7 Midwest   Protestant   325
 8 Midwest   Catholic     172
 9 Midwest   Jewish         3
10 Midwest   None         157
# ℹ 14 more rows
```

More compact; result is grouped.

Use `count()`

```
gss_sm ▷
  count(bigregion, religion) #<<
```

```
# A tibble: 24 × 3
   bigregion religion       n
   <fct>     <fct>      <int>
 1 Northeast Protestant   158
 2 Northeast Catholic     162
 3 Northeast Jewish        27
 4 Northeast None         112
 5 Northeast Other         28
 6 Northeast <NA>           1
 7 Midwest   Protestant   325
 8 Midwest   Catholic     172
 9 Midwest   Jewish         3
10 Midwest   None         157
# ℹ 14 more rows
```

One step; result is not grouped.

Pass results on to ... a table

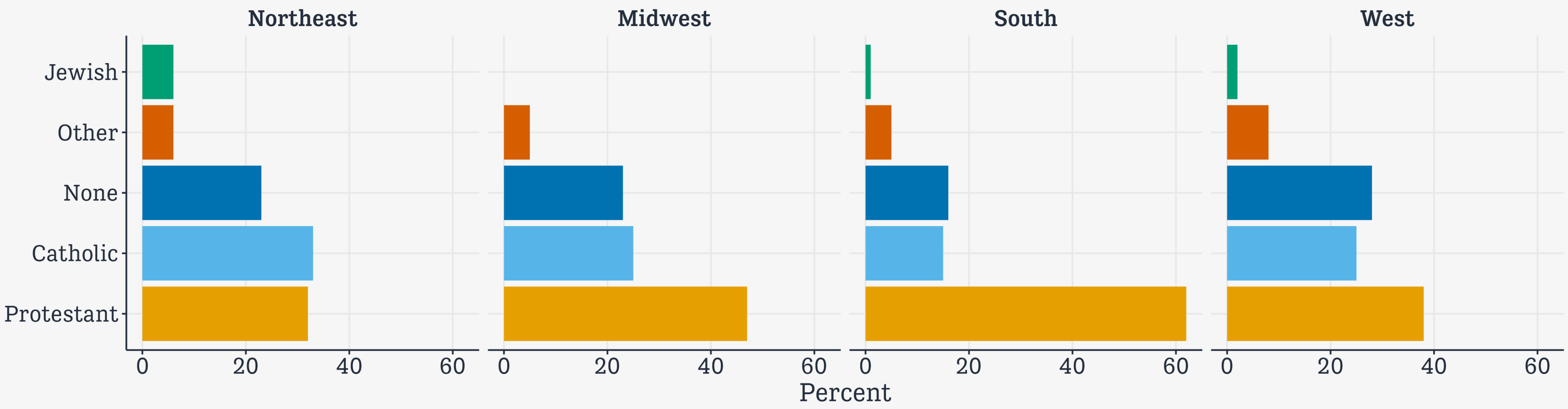
```r
gss_sm |>
  count(bigregion, religion) |>
  pivot_wider(names_from = bigregion, values_from = n) |>  #<<
  knitr::kable()
```

religion	Northeast	Midwest	South	West
Protestant	158	325	650	238
Catholic	162	172	160	155
Jewish	27	3	11	10
None	112	157	170	180
Other	28	33	50	48
NA	1	5	11	1

More on `pivot_wider()` and `kable()` soon ...

Pass results on to ... a graph

```r
gss_sm |>
  group_by(bigregion, religion) |>
  tally() |>
  mutate(pct = round((n/sum(n))*100), 1) |>
  drop_na() |>
  ggplot(mapping = aes(x = pct, y = reorder(religion, -pct), fill = religion)) + #<<
  geom_col() + #<<
    labs(x = "Percent", y = NULL) +
    guides(fill = "none") +
    facet_wrap(~ bigregion, nrow = 1)
```



Check by summarizing

```r
rel_by_region <- gss_sm |>
  count(bigregion, religion) |>
  mutate(pct = round((n/sum(n))*100, 1))

rel_by_region
```

```
# A tibble: 24 × 4
   bigregion religion       n   pct
   <fct>     <fct>      <int> <dbl>
 1 Northeast Protestant   158   5.5
 2 Northeast Catholic     162   5.7
 3 Northeast Jewish        27   0.9
 4 Northeast None         112   3.9
 5 Northeast Other         28   1  
 6 Northeast <NA>           1   0  
 7 Midwest   Protestant   325  11.3
 8 Midwest   Catholic     172   6  
 9 Midwest   Jewish         3   0.1
10 Midwest   None         157   5.5
# ℹ 14 more rows
```

Hm, did I sum over right group?

Check by summarizing

```r
rel_by_region <- gss_sm |>
  count(bigregion, religion) |>
  mutate(pct = round((n/sum(n))*100, 1))

rel_by_region
```

```
# A tibble: 24 × 4
   bigregion religion        n   pct
   <fct>     <fct>       <int> <dbl>
 1 Northeast Protestant    158   5.5
 2 Northeast Catholic      162   5.7
 3 Northeast Jewish         27   0.9
 4 Northeast None          112   3.9
 5 Northeast Other          28   1
 6 Northeast <NA>            1   0
 7 Midwest   Protestant    325  11.3
 8 Midwest   Catholic      172   6
 9 Midwest   Jewish          3   0.1
10 Midwest   None          157   5.5
# ℹ 14 more rows
```

Hm, did I sum over right group?

```r
## Each region should sum to ~100
rel_by_region |>
  group_by(bigregion) |>
  summarize(total = sum(pct))
```

```
# A tibble: 4 × 2
  bigregion total
  <fct>     <dbl>
1 Northeast  17
2 Midwest    24.3
3 South      36.7
4 West       22
```

No! What has gone wrong here?

Check by summarizing

```r
rel_by_region <- gss_sm |>
  count(bigregion, religion) |> #<<
  mutate(pct = round((n/sum(n))*100, 1))
```

`count()` returns ungrouped results, so there are no groups carry forward to the `mutate()` step.

```r
rel_by_region |>
  summarize(total = sum(pct))
```

```
# A tibble: 1 × 1
  total
  <dbl>
1   100
```

With `count()`, the `pct` values here are the marginals for the whole table.

Check by summarizing

```r
rel_by_region <- gss_sm |>
  count(bigregion, religion) |> #<<
  mutate(pct = round((n/sum(n))*100, 1))
```

`count()` returns ungrouped results, so there are no groups carry forward to the **`mutate()`** step.

```r
rel_by_region |>
  summarize(total = sum(pct))
```

```
# A tibble: 1 × 1
  total
  <dbl>
1   100
```

With **`count()`**, the `pct` values here are the marginals for the whole table.

```r
rel_by_region <- gss_sm |>
  group_by(bigregion, religion) |> #<<
  tally() |> #<<
  mutate(pct = round((n/sum(n))*100, 1))
```

```r
# Check
rel_by_region |>
  group_by(bigregion) |>
  summarize(total = sum(pct))
```

```
# A tibble: 4 × 2
  bigregion total
  <fct>     <dbl>
1 Northeast 100
2 Midwest    99.9
3 South     100
4 West      100.
```

We get some rounding error because we used `round()` after summing originally.

Two lessons

Check your tables!

Pipelines feed their content forward, so you need to make sure your results are not incorrect.

Often, complex tables and graphs can be disturbingly plausible even when wrong.

So, figure out what the result should be and test it!

Starting with simple or toy cases can help with this process.

Two lessons

Inspect your pipes!

Understand pipelines by running them forward or peeling them back a step at a time.

This is a *very* effective way to understand your own and other people's code.

Use `dplyr` to make summary tables.

Then send your clean tables to `ggplot`.

Facets are often better than Guides

Let’s put that table in an object

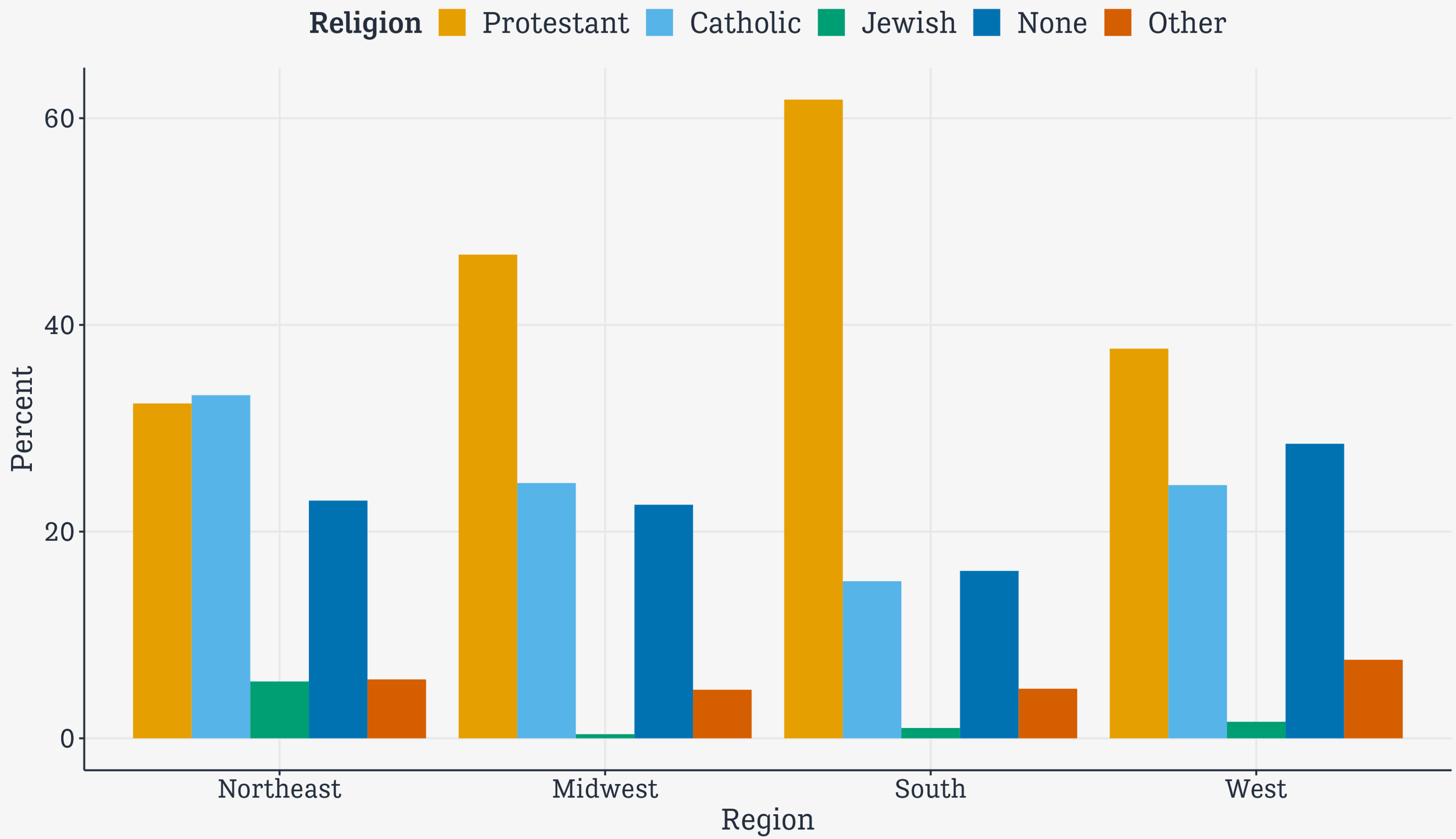
```r
rel_by_region <- gss_sm |>
  group_by(bigregion, religion) |>
  tally() |>
  mutate(pct = round((n/sum(n))*100, 1)) |>
  drop_na()


head(rel_by_region)
```

```
# A tibble: 6 × 4
# Groups:   bigregion [2]
  bigregion religion       n   pct
  <fct>     <fct>      <int> <dbl>
1 Northeast Protestant   158  32.4
2 Northeast Catholic     162  33.2
3 Northeast Jewish        27   5.5
4 Northeast None         112  23  
5 Northeast Other         28   5.7
6 Midwest   Protestant   325  46.8
```

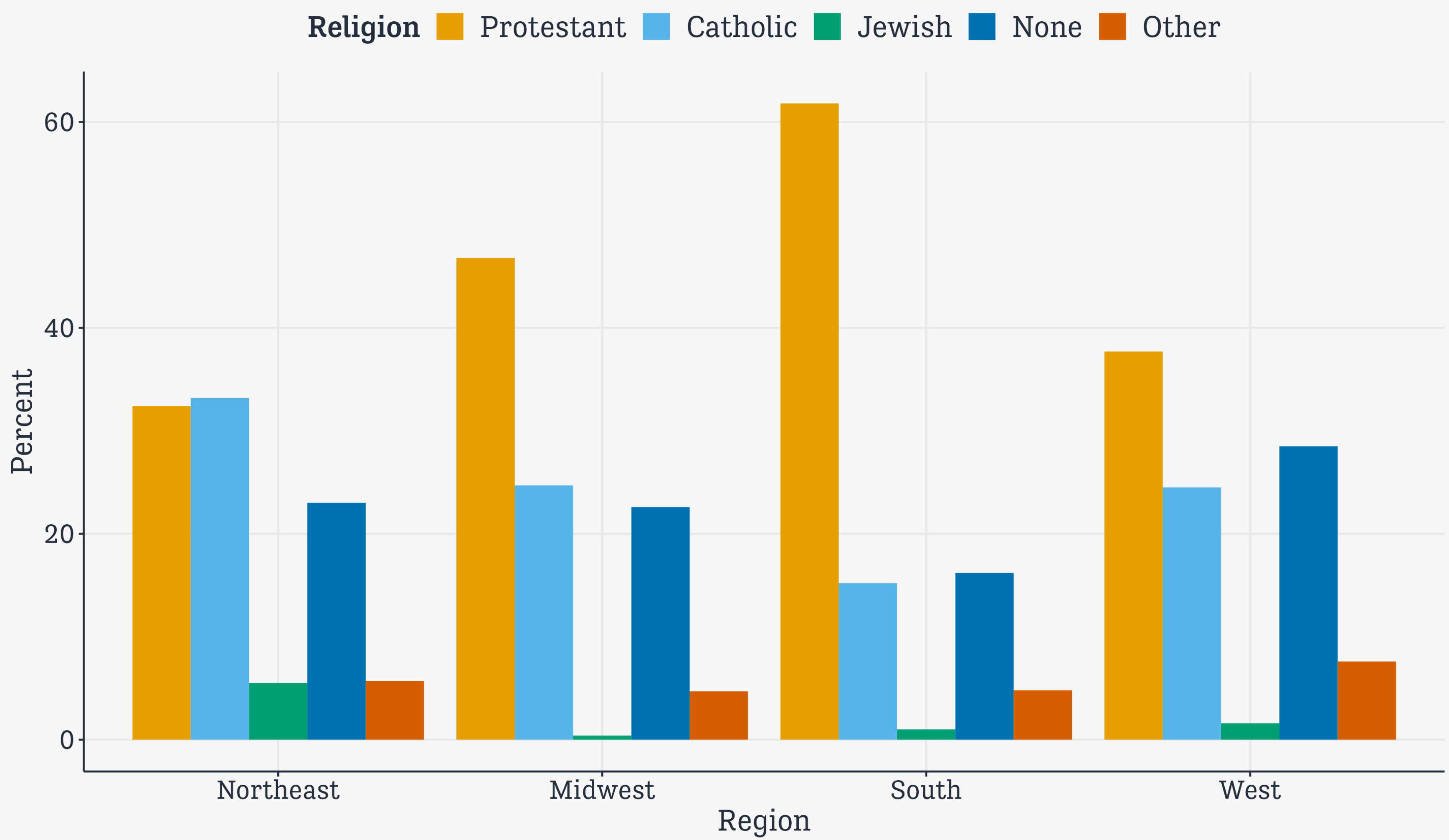
We might write ...

```r
p <- ggplot(data = rel_by_region,
                mapping = aes(x = bigregion,
                              y = pct,
                              fill = religion))
p_out <- p + geom_col(position = "dodge") +
    labs(x = "Region",
         y = "Percent",
         fill = "Religion")
```

We might write …



Is this an effective graph? Not really!



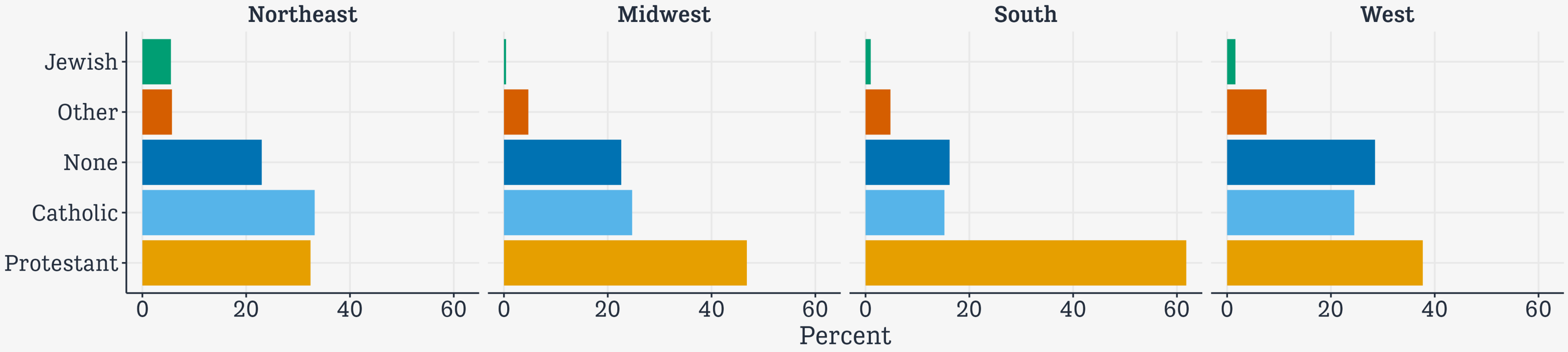
Try faceting instead

```r
p <- ggplot(data = rel_by_region,
                mapping = aes(x = pct, #<<
                              y = reorder(religion, -pct), #<<
                              fill = religion))
p_out_facet <- p + geom_col() +
  guides(fill = "none") +
  facet_wrap(~ bigregion, nrow = 1) +
  labs(x = "Percent",
       y = NULL)
```

Putting categories on the y-axis is a very useful trick.

Faceting reduces the number of guides the viewer needs to consult.

Try faceting instead



Try faceting instead

Try putting categories on the y-axis. (And reorder them by x.)

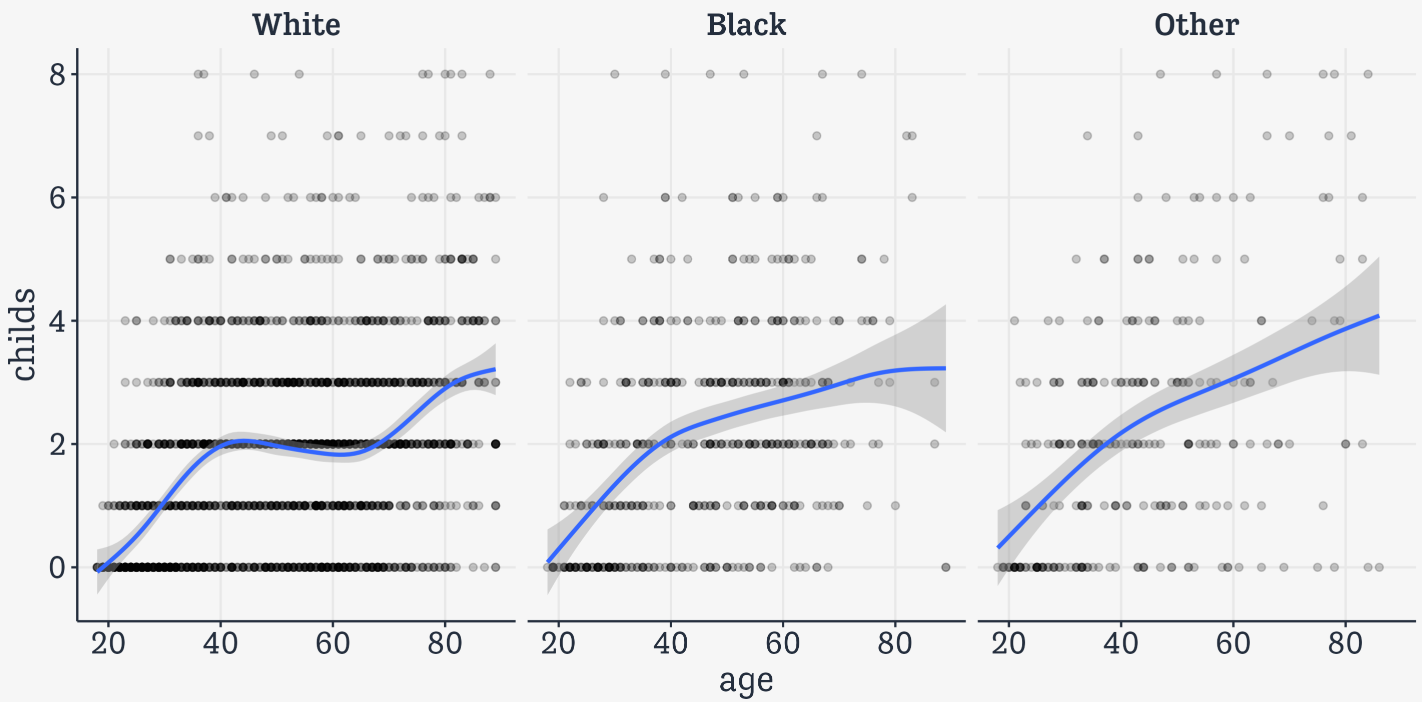
Try faceting variables instead of mapping them to color or shape.

Try to minimize the need for guides and legends.

Two kinds of facet

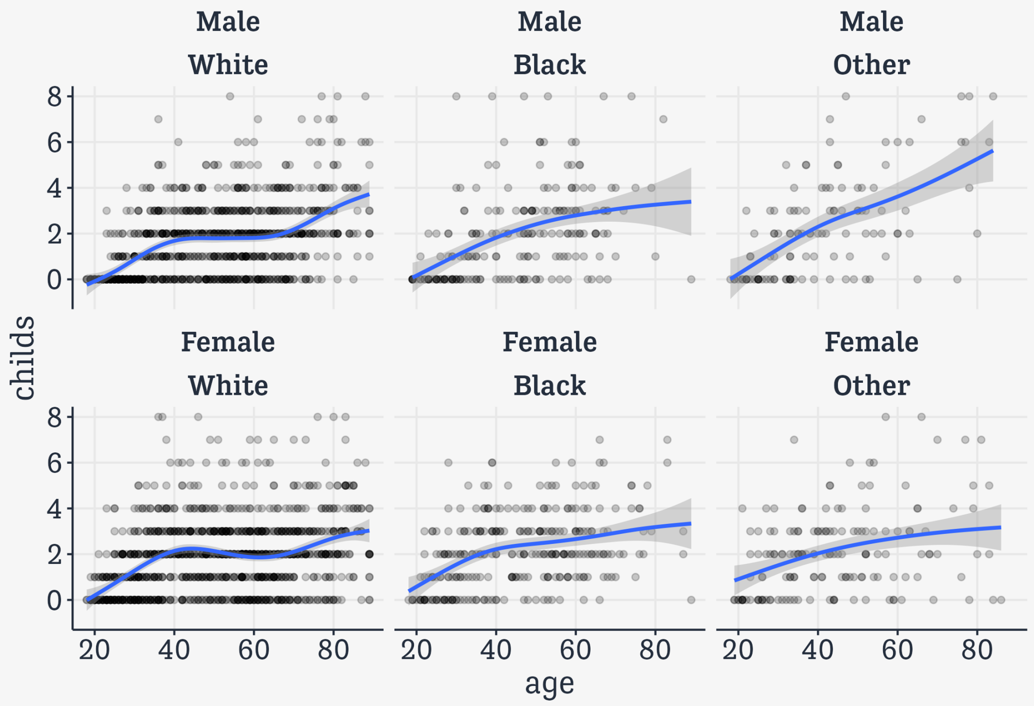
Facet Children vs Age, by Race

```r
p <-  ggplot(data = gss_sm,
             mapping = aes(x = age, y = childs))

p + geom_point(alpha = 0.2) +
  geom_smooth() +
  facet_wrap(~ race)
```



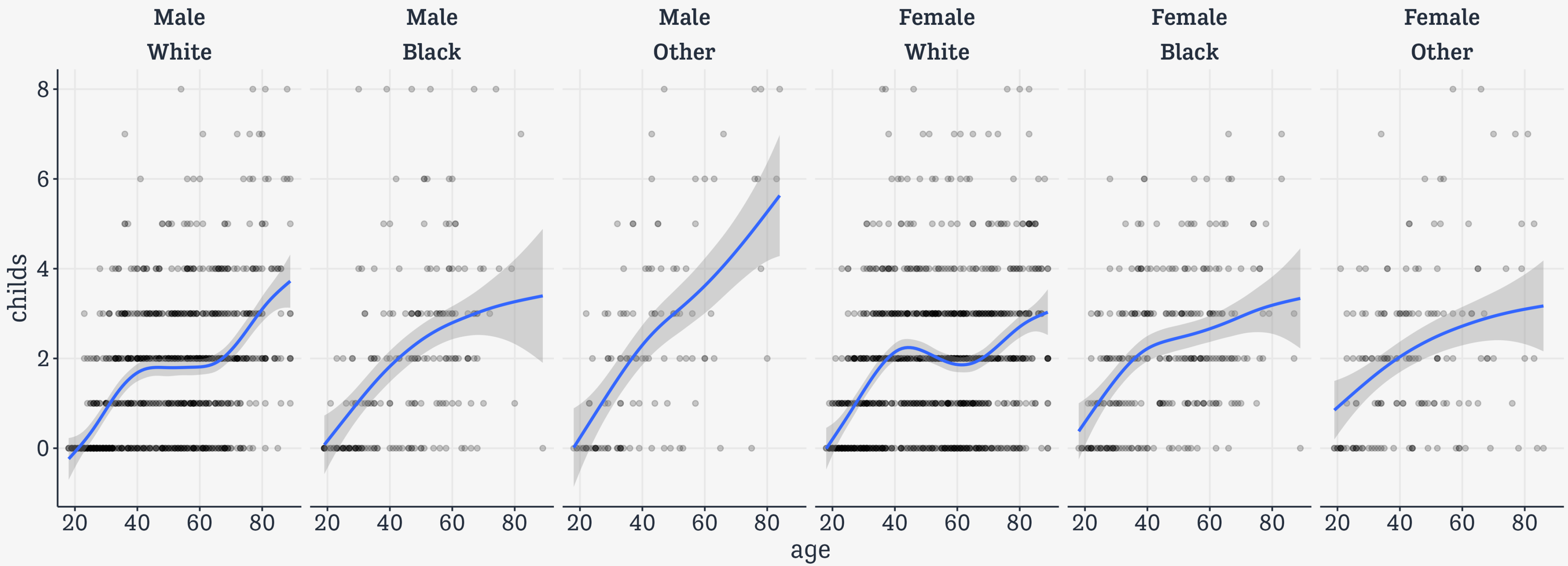
Facet by more than one variable

```r
p <- ggplot(data = gss_sm,
            mapping = aes(x = age, y = childs))

p + geom_point(alpha = 0.2) +
  geom_smooth() +
  facet_wrap(~ sex + race) #<<
```



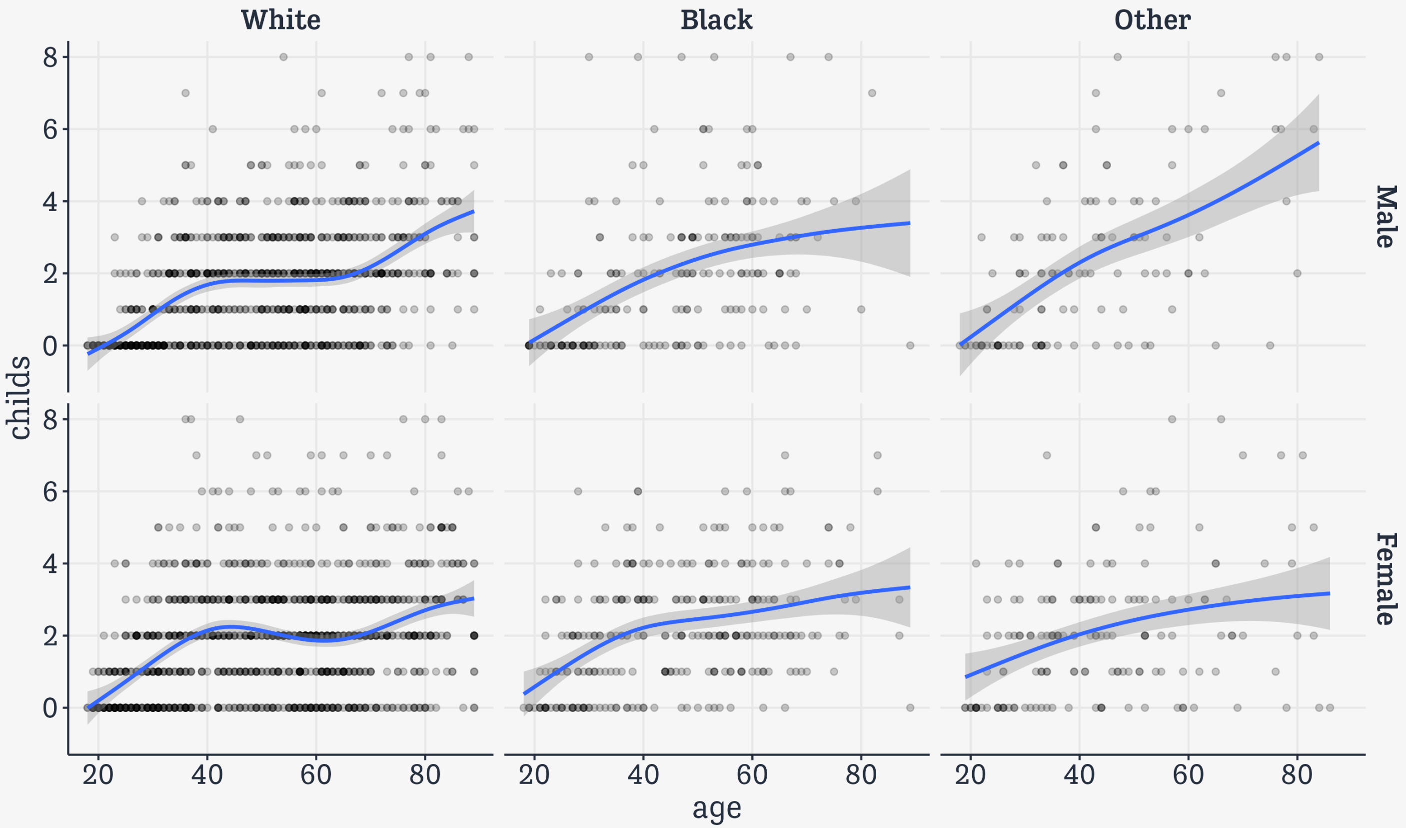
Arrange facet_wrap() quite freely

```r
p <-  ggplot(data = gss_sm,
             mapping = aes(x = age, y = childs))

p + geom_point(alpha = 0.2) +
  geom_smooth() +
  facet_wrap(~ sex + race, nrow = 1) #<<
```



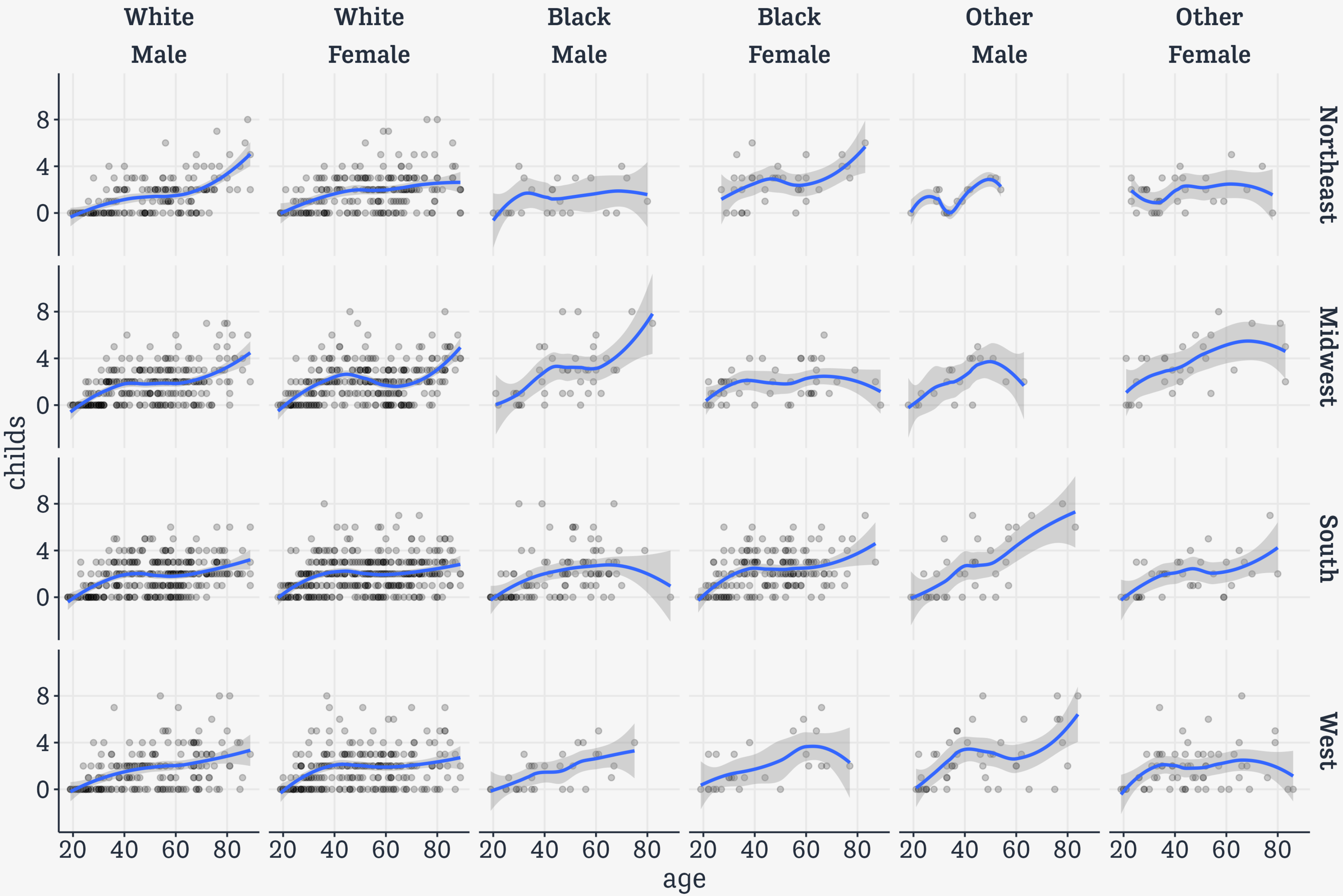
facet_grid() is more like a true crosstab

```r
p + geom_point(alpha = 0.2) +
  geom_smooth() +
  facet_grid(sex ~ race) #<<
```



Extend both to multi-way views

```r
p_out <- p + geom_point(alpha = 0.2) +
  geom_smooth() +
  facet_grid(bigregion ~ race + sex) #<<
```


White Male
White Female
Black Male
Black Female
Other Male
Other Female
Northeast
Midwest
South
West
childs
age
0
4
8
20
40
60
80

What we’ve built-up

Core Grammar

```
p <- ggplot(data = <DATA>,
            mapping=aes(<MAPPINGS>)) +

            <GEOM_FUNCTION>(
            mapping = aes(<MAPPINGS>),
            stat = <STAT>,
            position = <POSITION>) +

           <SCALE_FUNCTION> +

           <COORDINATE_FUNCTION> +

           <FACET_FUNCTION> +

           <THEME_FUNCTION>
```

Core grammar

Grouped data; faceting

Along with a few peeks at scale transformations, guide adjustments, and theme adjustment

```
p <- ggplot(data = gapminder,
            mapping = aes(x = year,
                          y = gdpPercap))

p + geom_line(aes(group = country)) +
    scale_y_log10() +
    coord_cartesian() +
    facet_wrap(~ continent) +
    theme_minimal()
```

All basic steps

dplyr and Pipelining

The elements of filtering and summarizing

```
gss_sm |>
  group_by(bigregion, religion) |>
  tally() |>
  mutate(freq = n / sum(n),
         pct = round((freq*100), 1))
```

```
# A tibble: 24 × 5
# Groups:   bigregion [4]
   bigregion religion        n    freq   pct
   <fct>     <fct>       <int>   <dbl> <dbl>
 1 Northeast Protestant    158 0.324    32.4
 2 Northeast Catholic      162 0.332    33.2
 3 Northeast Jewish         27 0.0553    5.5
 4 Northeast None          112 0.230    23
 5 Northeast Other          28 0.0574    5.7
 6 Northeast <NA>            1 0.00205   0.2
 7 Midwest   Protestant    325 0.468    46.8
 8 Midwest   Catholic      172 0.247    24.7
 9 Midwest   Jewish          3 0.00432   0.4
10 Midwest   None          157 0.226    22.6
# ℹ 14 more rows
```

Example and extension:
Organ Donation data

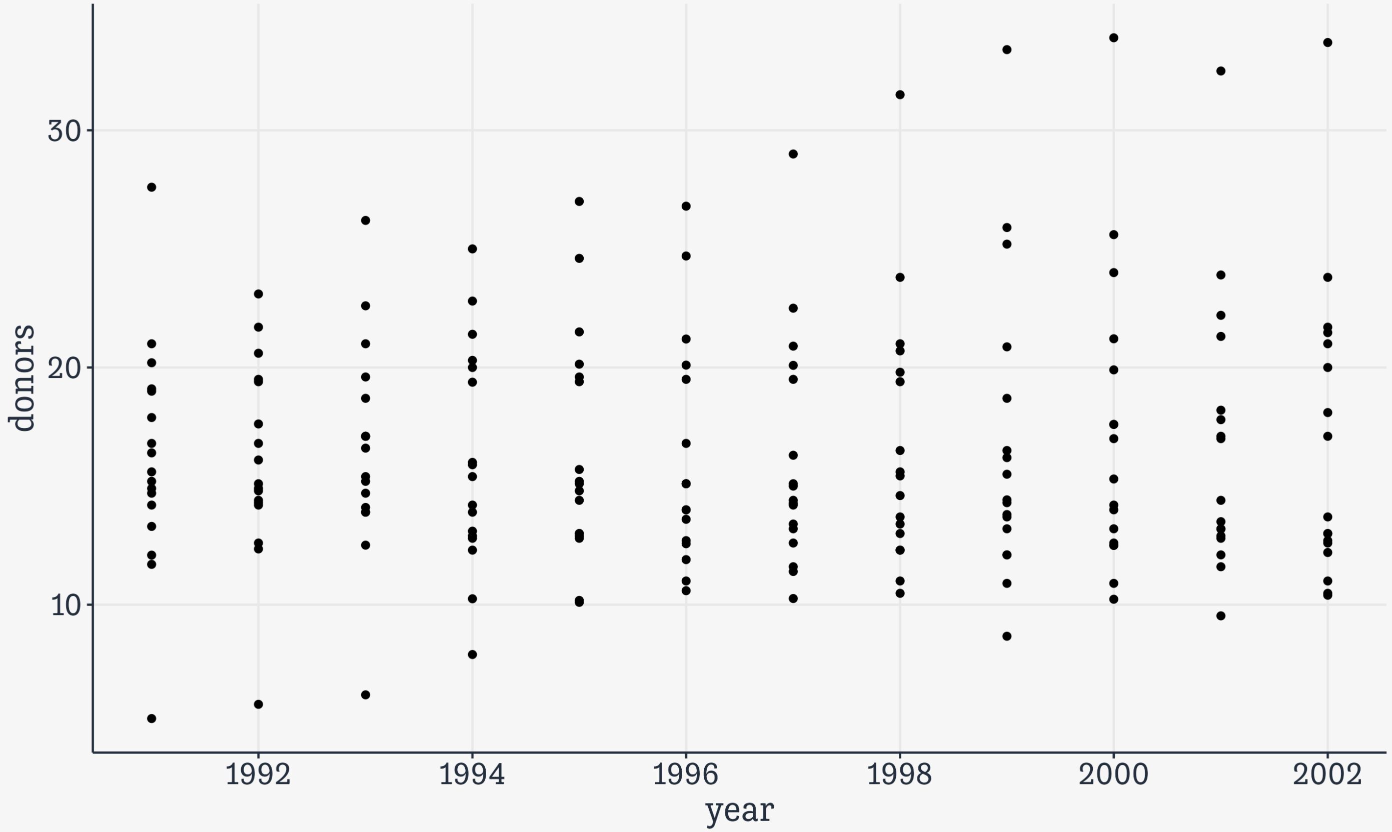
organdata is in the socviz package

```
organdata
```

```
# A tibble: 238 × 21
   country   year       donors   pop pop_dens   gdp gdp_lag health health_lag
   <chr>     <date>      <dbl> <int>    <dbl> <int>   <int>  <dbl>      <dbl>
 1 Australia NA          NA    17065    0.220 16774   16591   1300       1224
 2 Australia 1991-01-01  12.1  17284    0.223 17171   16774   1379       1300
 3 Australia 1992-01-01  12.4  17495    0.226 17914   17171   1455       1379
 4 Australia 1993-01-01  12.5  17667    0.228 18883   17914   1540       1455
 5 Australia 1994-01-01  10.2  17855    0.231 19849   18883   1626       1540
 6 Australia 1995-01-01  10.2  18072    0.233 21079   19849   1737       1626
 7 Australia 1996-01-01  10.6  18311    0.237 21923   21079   1846       1737
 8 Australia 1997-01-01  10.3  18518    0.239 22961   21923   1948       1846
 9 Australia 1998-01-01  10.5  18711    0.242 24148   22961   2077       1948
10 Australia 1999-01-01   8.67 18926    0.244 25445   24148   2231       2077
# ℹ 228 more rows
# ℹ 12 more variables: pubhealth <dbl>, roads <dbl>, cerebvas <int>,
#   assault <int>, external <int>, txp_pop <dbl>, world <chr>, opt <chr>,
#   consent_law <chr>, consent_practice <chr>, consistent <chr>, ccode <chr>
```

First look

```
p <- ggplot(data = organdata,
            mapping = aes(x = year, y = donors))
p + geom_point()
```

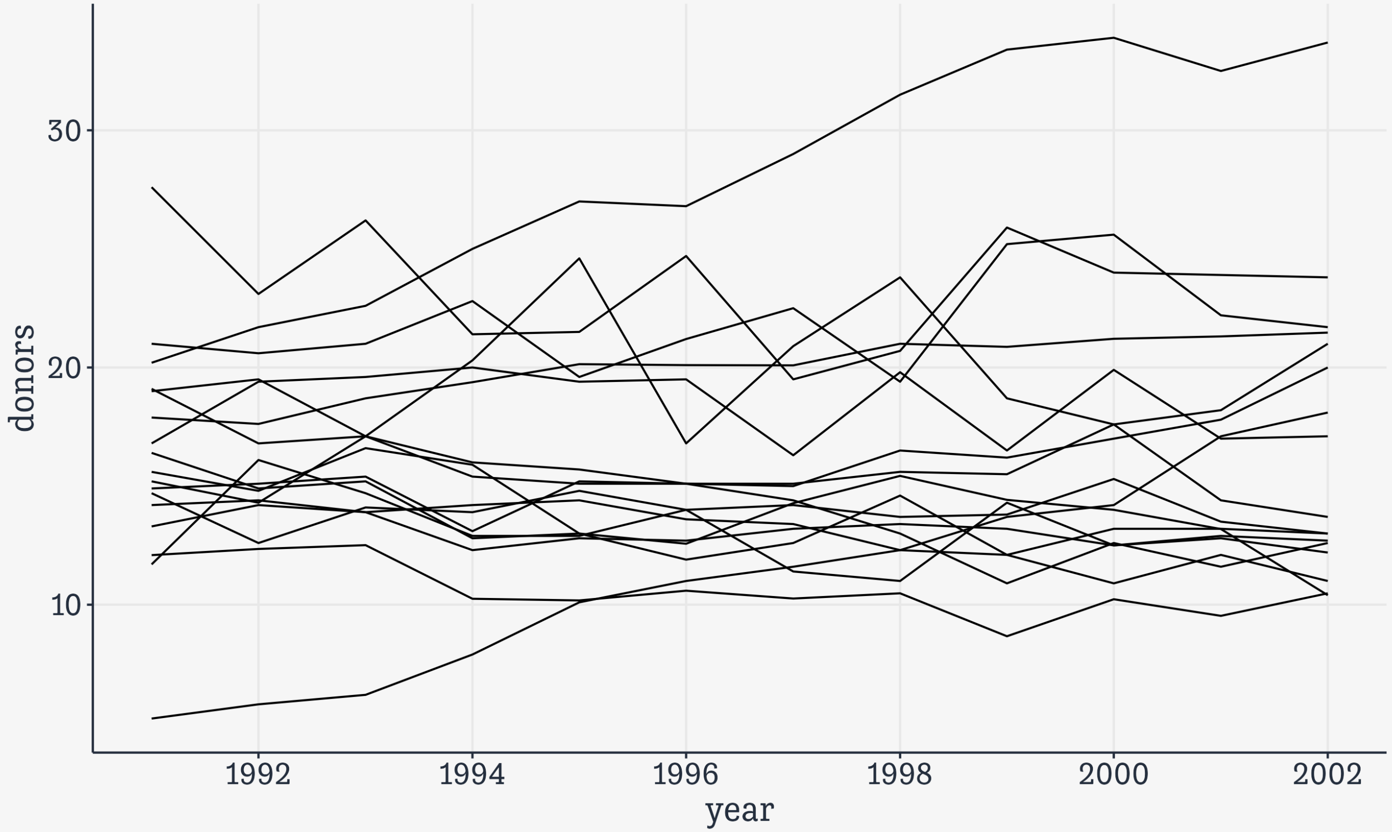


First look

```r
p <- ggplot(data = organdata,
            mapping = aes(x = year, y = donors))
p + geom_line()
```

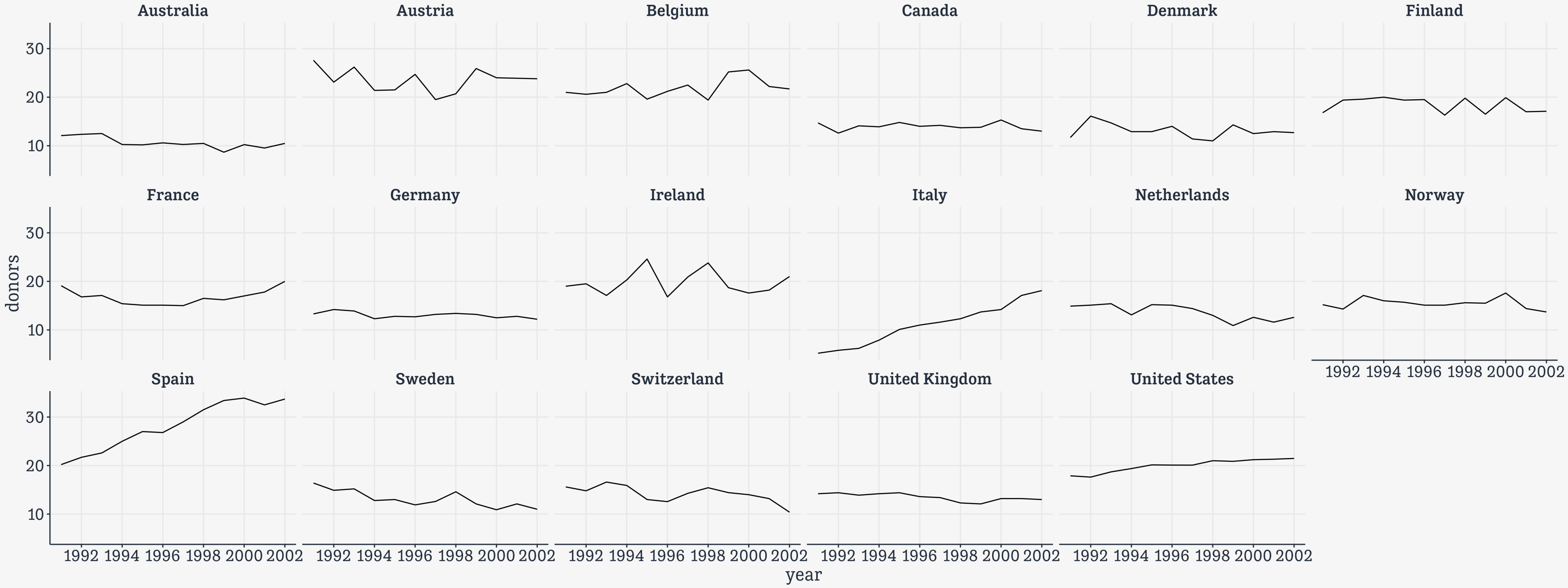
First look

```r
p <- ggplot(data = organdata,
            mapping = aes(x = year, y = donors))
p + geom_line(aes(group = country))
```



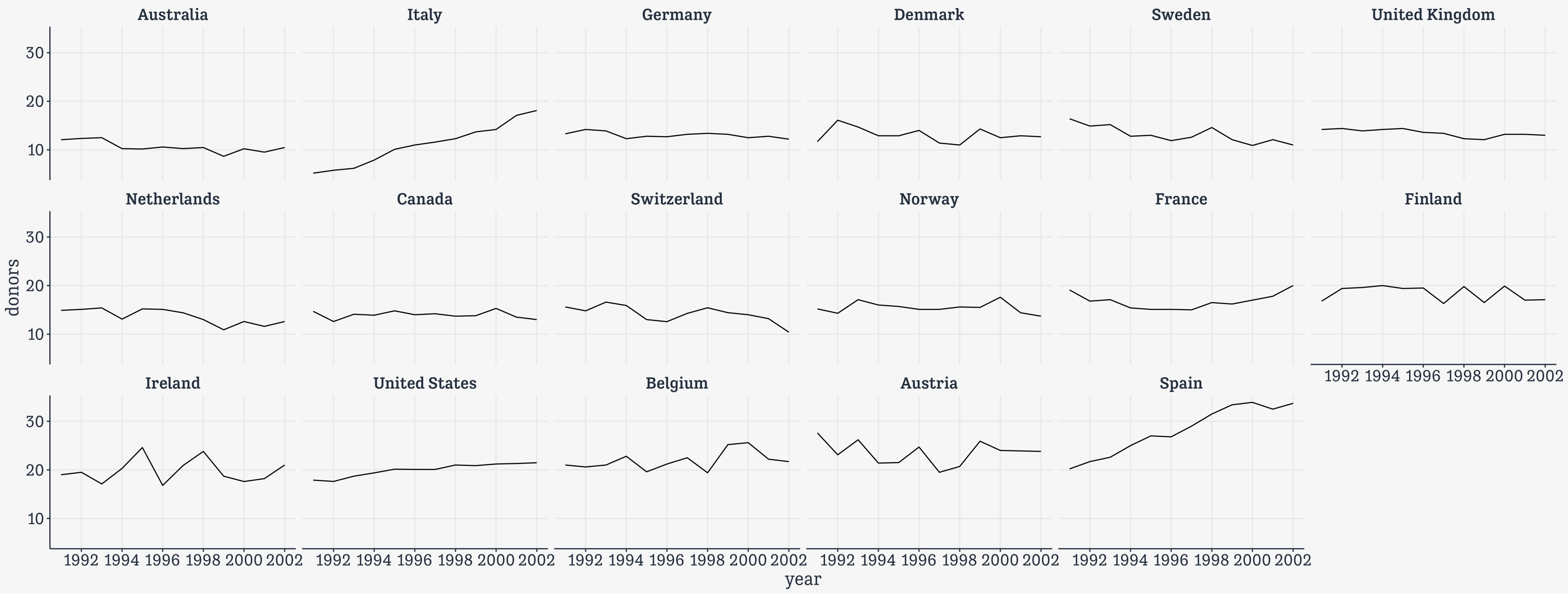
First look

```r
p <- ggplot(data = organdata,
            mapping = aes(x = year, y = donors))
p + geom_line() +
  facet_wrap(~ country, nrow = 3)
```



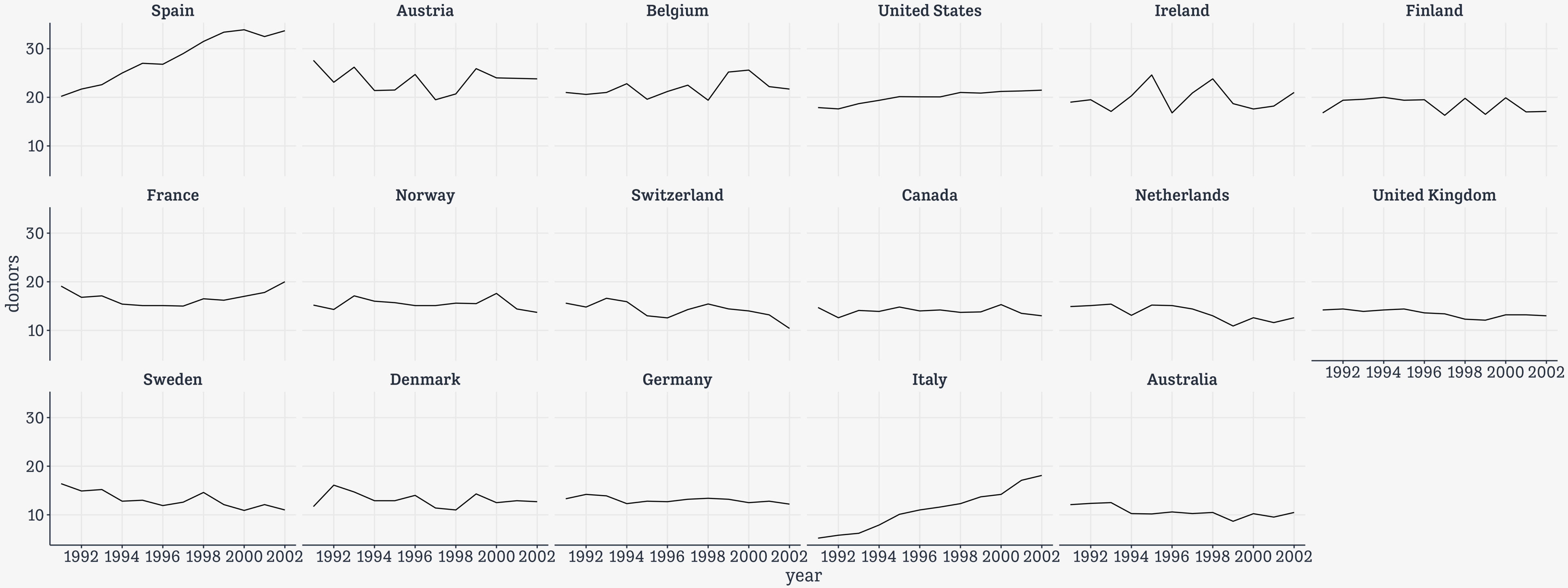
First look

```r
p ← ggplot(data = organdata,
           mapping = aes(x = year, y = donors))
p + geom_line() +
  facet_wrap(~ reorder(country, donors, na.rm = TRUE), nrow = 3)
```



First look

```r
p <- ggplot(data = organdata,
            mapping = aes(x = year, y = donors))
p + geom_line() +
  facet_wrap(~ reorder(country, -donors, na.rm = TRUE), nrow = 3)
```



Summarize better with `dplyr`

Summarize a bunch of variables

```
by_country <- organdata |>
  group_by(consent_law, country)  |>
    summarize(donors_mean= mean(donors, na.rm = TRUE),
              donors_sd = sd(donors, na.rm = TRUE),
              gdp_mean = mean(gdp, na.rm = TRUE),
              health_mean = mean(health, na.rm = TRUE),
              roads_mean = mean(roads, na.rm = TRUE),
              cerebvas_mean = mean(cerebvas, na.rm = TRUE))

head(by_country)
```

```
# A tibble: 6 × 8
# Groups:   consent_law [1]
  consent_law country     donors_mean donors_sd gdp_mean health_mean roads_mean
  <chr>       <chr>             <dbl>     <dbl>    <dbl>       <dbl>      <dbl>
1 Informed    Australia          10.6     1.14    22179.       1958.      105. 
2 Informed    Canada             14.0     0.751   23711.       2272.      109. 
3 Informed    Denmark            13.1     1.47    23722.       2054.      102. 
4 Informed    Germany            13.0     0.611   22163.       2349.      113. 
5 Informed    Ireland            19.8     2.48    20824.       1480.      118. 
6 Informed    Netherlands        13.7     1.55    23013.       1993.       76.1
# ℹ 1 more variable: cerebvas_mean <dbl>
```

This works, but there’s so much repetition! It’s an open invitation to make mistakes copying and pasting.

DRY:

Don't Repeat Yourself

Use across() and where() instead

```r
  by_country <- organdata |>
    group_by(consent_law, country) |>
      summarize(across(where(is.numeric),#<<
                        list(mean = ~ mean(.x, na.rm = TRUE),
                             sd = ~ sd(.x, na.rm = TRUE))))
head(by_country)
```

```
# A tibble: 6 × 28
# Groups:   consent_law [1]
  consent_law country     donors_mean donors_sd pop_mean pop_sd pop_dens_mean
  <chr>       <chr>             <dbl>     <dbl>    <dbl>  <dbl>         <dbl>
1 Informed    Australia          10.6     1.14    18318.  831.          0.237
2 Informed    Canada             14.0     0.751   29608. 1193.          0.297
3 Informed    Denmark            13.1     1.47     5257.   80.6        12.2
4 Informed    Germany            13.0     0.611   80255. 5158.         22.5
5 Informed    Ireland            19.8     2.48     3674.  132.          5.23
6 Informed    Netherlands        13.7     1.55    15548.  373.         37.4
# ℹ 21 more variables: pop_dens_sd <dbl>, gdp_mean <dbl>, gdp_sd <dbl>,
#   gdp_lag_mean <dbl>, gdp_lag_sd <dbl>, health_mean <dbl>, health_sd <dbl>,
#   health_lag_mean <dbl>, health_lag_sd <dbl>, pubhealth_mean <dbl>,
#   pubhealth_sd <dbl>, roads_mean <dbl>, roads_sd <dbl>, cerebvas_mean <dbl>,
#   cerebvas_sd <dbl>, assault_mean <dbl>, assault_sd <dbl>,
#   external_mean <dbl>, external_sd <dbl>, txp_pop_mean <dbl>,
#   txp_pop_sd <dbl>
```

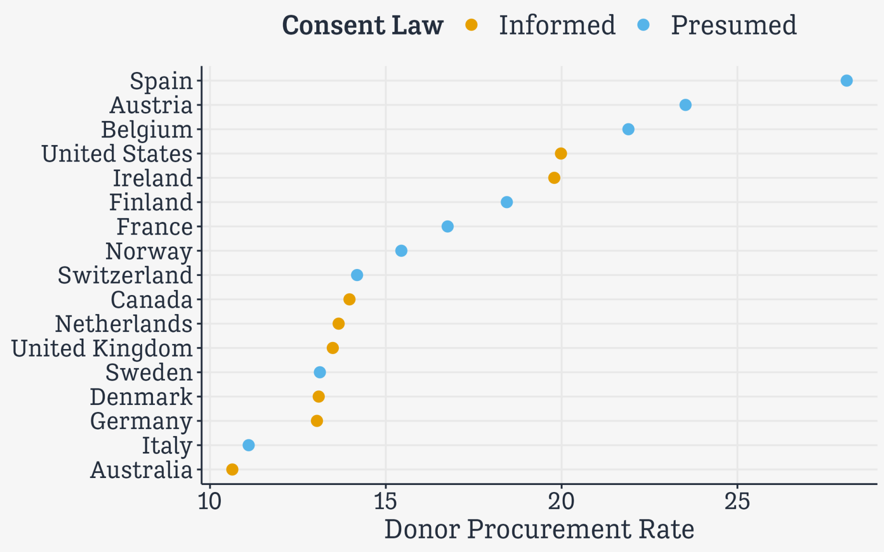
Use across() and where() instead

```r
by_country ← organdata ▷
  group_by(consent_law, country) ▷
    summarize(across(where(is.numeric),#<<
                       list(mean = ~ mean(.x, na.rm = TRUE),
                            sd = ~ sd(.x, na.rm = TRUE))),
              .groups = "drop") #<<
head(by_country)
```

```
# A tibble: 6 × 28
  consent_law country     donors_mean donors_sd pop_mean pop_sd pop_dens_mean
  <chr>       <chr>             <dbl>     <dbl>    <dbl>  <dbl>         <dbl>
1 Informed    Australia          10.6     1.14    18318.  831.          0.237
2 Informed    Canada             14.0     0.751   29608. 1193.          0.297
3 Informed    Denmark            13.1     1.47     5257.   80.6        12.2
4 Informed    Germany            13.0     0.611   80255. 5158.         22.5
5 Informed    Ireland            19.8     2.48     3674.  132.          5.23
6 Informed    Netherlands        13.7     1.55    15548.  373.         37.4
# ℹ 21 more variables: pop_dens_sd <dbl>, gdp_mean <dbl>, gdp_sd <dbl>,
#   gdp_lag_mean <dbl>, gdp_lag_sd <dbl>, health_mean <dbl>, health_sd <dbl>,
#   health_lag_mean <dbl>, health_lag_sd <dbl>, pubhealth_mean <dbl>,
#   pubhealth_sd <dbl>, roads_mean <dbl>, roads_sd <dbl>, cerebvas_mean <dbl>,
#   cerebvas_sd <dbl>, assault_mean <dbl>, assault_sd <dbl>,
#   external_mean <dbl>, external_sd <dbl>, txp_pop_mean <dbl>,
#   txp_pop_sd <dbl>
```

Plot our summary data

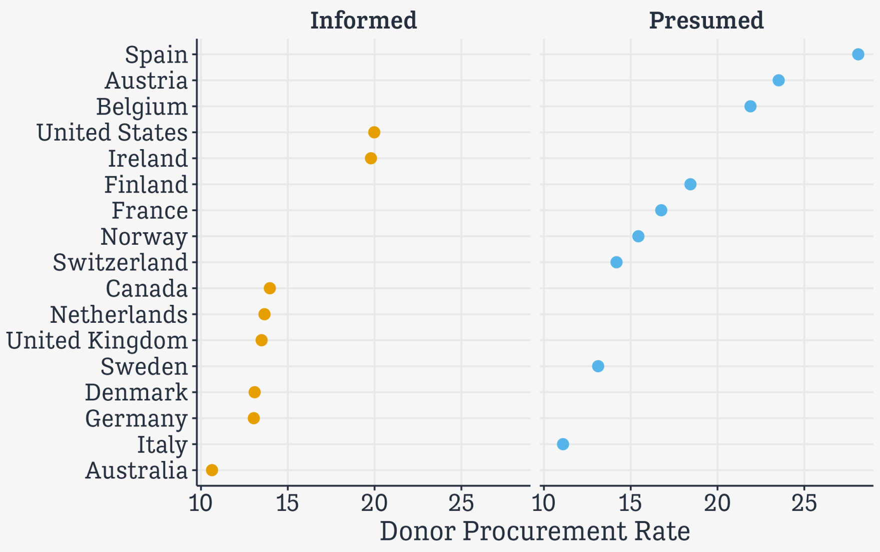
```r
by_country |>
  ggplot(mapping =
           aes(x = donors_mean,
               y = reorder(country, donors_mean
               color = consent_law)) +
  geom_point(size=3) +
  labs(x = "Donor Procurement Rate",
       y = NULL,
       color = "Consent Law")
```



What about faceting it instead?

The problem is that countries can only be in one Consent Law category.

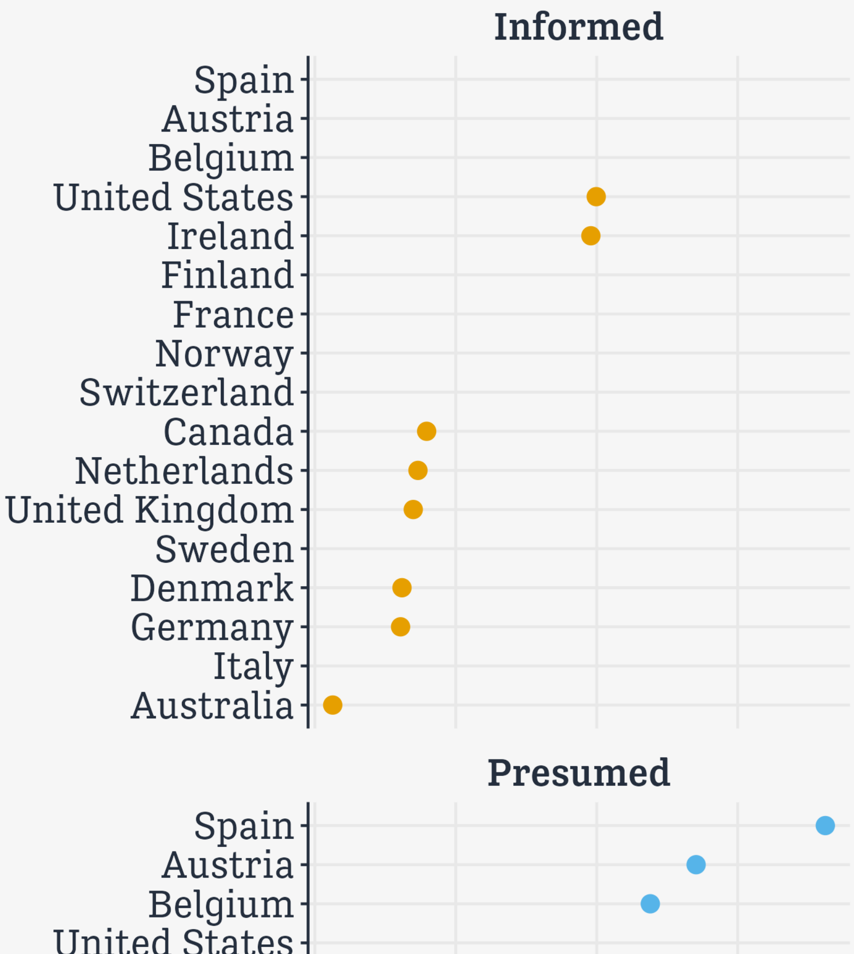
```
by_country ▷
  ggplot(mapping =
           aes(x = donors_mean,
               y = reorder(country, donors_mean
               color = consent_law)) +
  geom_point(size=3) +
  guides(color = "none") +
  facet_wrap(~ consent_law) + #<<
  labs(x = "Donor Procurement Rate",
       y = NULL,
       color = "Consent Law")
```



What about faceting it instead?

Restricting to one column doesn’t fix it.

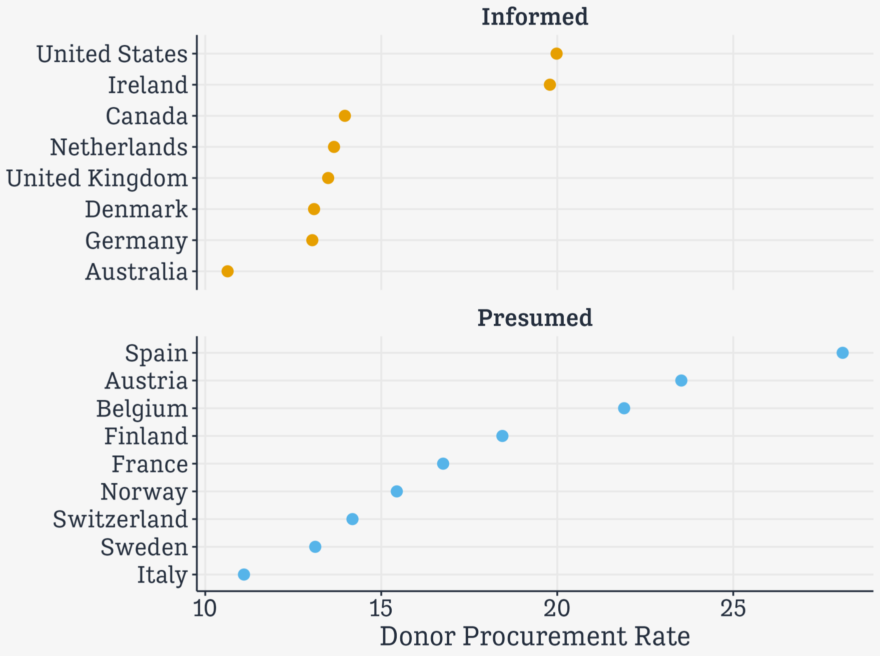
```r
by_country |>
  ggplot(mapping =
           aes(x = donors_mean,
               y = reorder(country, donors_mean
               color = consent_law)) +
  geom_point(size=3) +
  guides(color = "none") +
  facet_wrap(~ consent_law, ncol = 1) + #<<
  labs(x = "Donor Procurement Rate",
       y = NULL,
       color = "Consent Law")
```



Allow the y-scale to vary

Normally the point of a facet is to preserve comparability between panels by not allowing the scales to vary. But for categorical measures it can be useful to allow this.

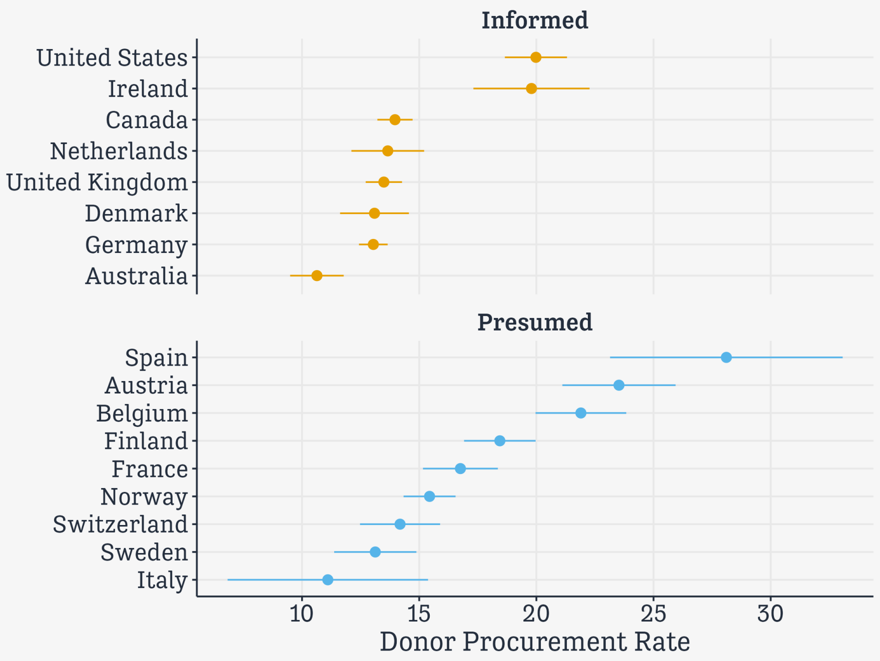
```
by_country |>
  ggplot(mapping =
           aes(x = donors_mean,
               y = reorder(country, donors_mean
               color = consent_law)) +
  geom_point(size=3) +
  guides(color = "none") +
  facet_wrap(~ consent_law,
             ncol = 1,
             scales = "free_y") +  #<<
  labs(x = "Donor Procurement Rate",
       y = NULL,
       color = "Consent Law")
```



Again, these methods are general

:::: {.columns} ::: {.column width="50%"}

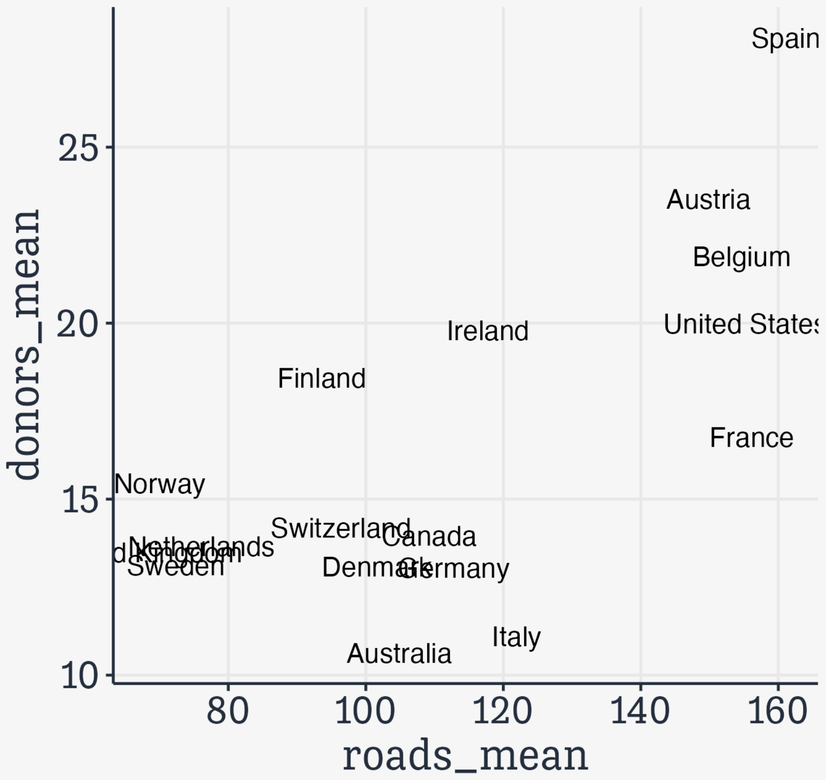
```
by_country |>
  ggplot(mapping =
           aes(x = donors_mean,
               y = reorder(country, donors_mean
               color = consent_law)) +
  geom_pointrange(mapping = #<<
                    aes(xmin = donors_mean - do
                        xmax = donors_mean + do
  guides(color = "none") +
  facet_wrap(~ consent_law,
             ncol = 1,
             scales = "free_y") +
  labs(x = "Donor Procurement Rate",
       y = NULL,
       color = "Consent Law")
```



Plot text directly

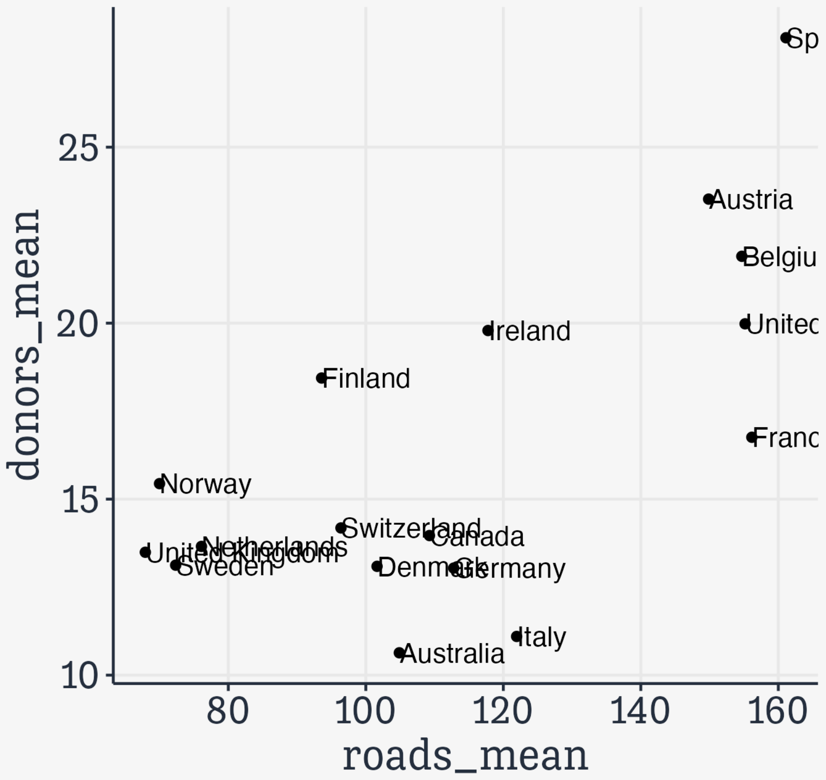
`geom_text()` for basic labels

```r
by_country |>
  ggplot(mapping = aes(x = roads_mean,
                       y = donors_mean)) +
  geom_text(mapping = aes(label = country))
```



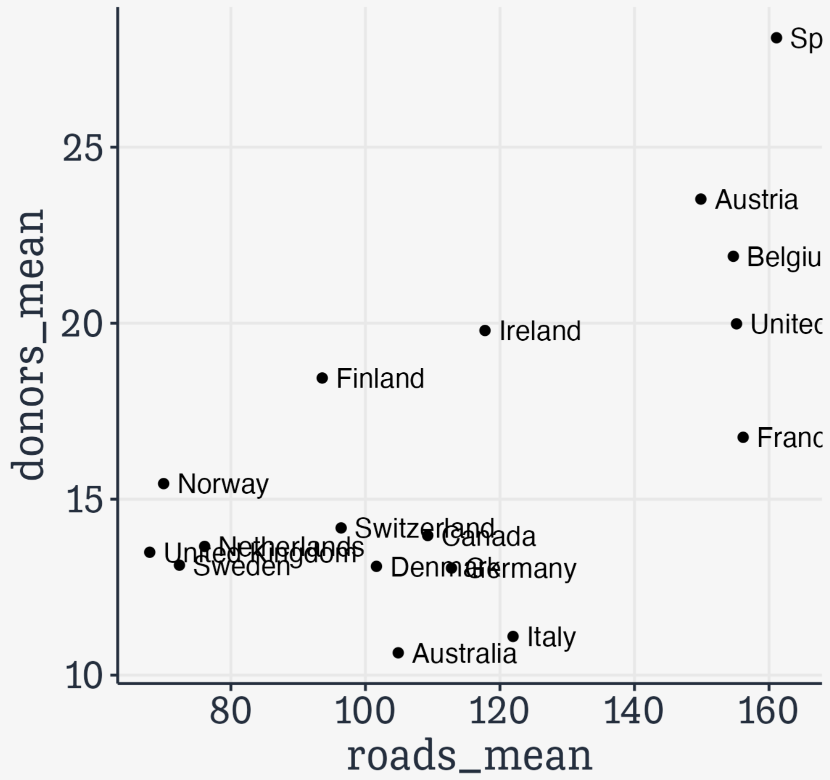
It’s not very flexible

```
by_country |>
  ggplot(mapping = aes(x = roads_mean,
                       y = donors_mean)) +
  geom_point() +
  geom_text(mapping = aes(label = country),
            hjust = 0)
```



There are tricks, but they're limited

```
by_country ▷
  ggplot(mapping = aes(x = roads_mean,
                       y = donors_mean)) +
  geom_point() +
  geom_text(mapping = aes(x = roads_mean + 2,
                          label = country),
            hjust = 0)
```



We'll use `ggrepel` instead

The `ggrepel` package provides `geom_text_repel()` and `geom_label_repel()`

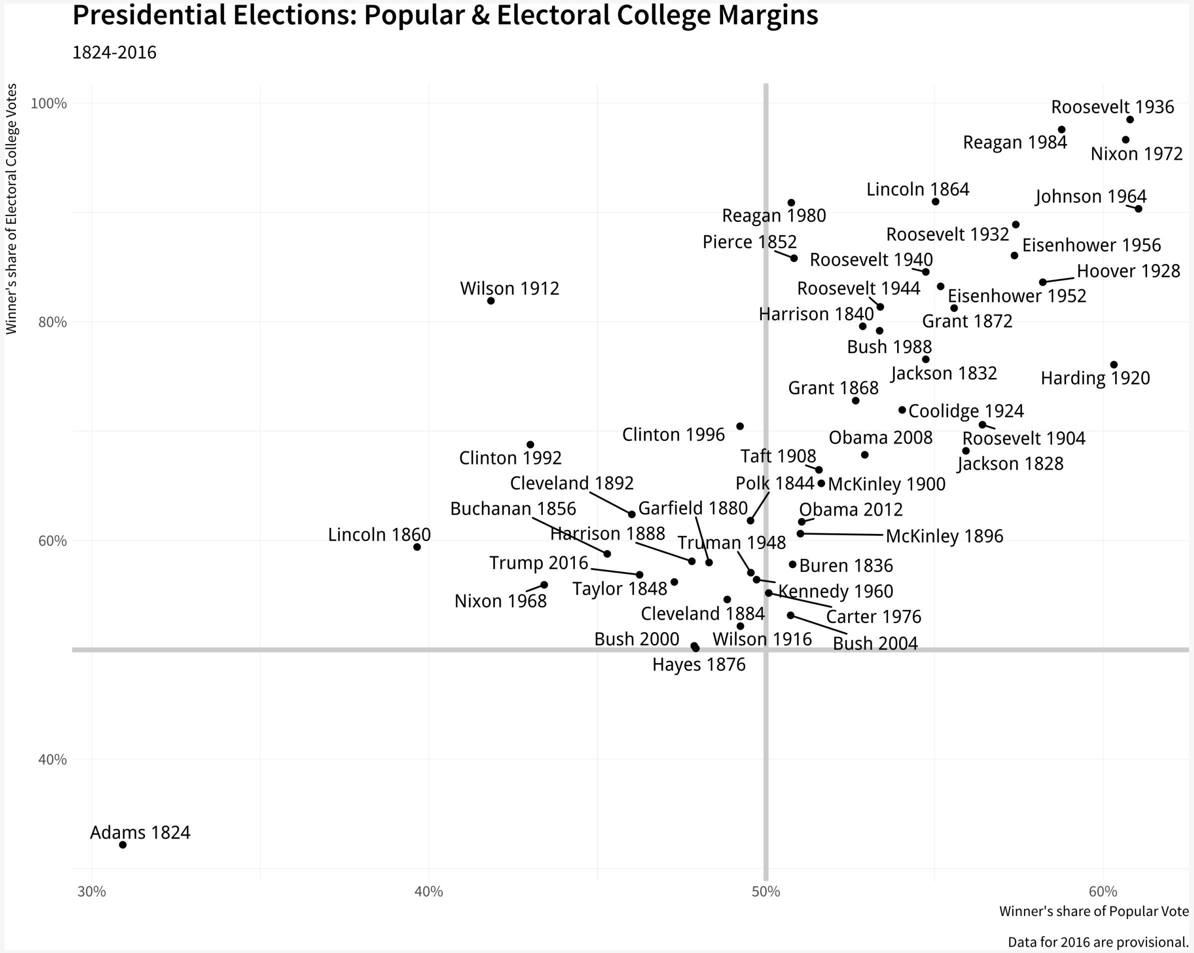
Example: U.S. Historic Presidential Elections

elections_historic is in socviz

```r
elections_historic
```

```
# A tibble: 49 × 19
   election  year winner      win_party ec_pct popular_pct popular_margin  votes
      <int> <int> <chr>       <chr>      <dbl>       <dbl>          <dbl>  <int>
 1       10  1824 John Quinc… D.-R.      0.322       0.309        -0.104  1.13e5
 2       11  1828 Andrew Jac… Dem.       0.682       0.559         0.122  6.43e5
 3       12  1832 Andrew Jac… Dem.       0.766       0.547         0.178  7.03e5
 4       13  1836 Martin Van… Dem.       0.578       0.508         0.142  7.63e5
 5       14  1840 William He… Whig       0.796       0.529         0.0605 1.28e6
 6       15  1844 James Polk  Dem.       0.618       0.495         0.0145 1.34e6
 7       16  1848 Zachary Ta… Whig       0.562       0.473         0.0479 1.36e6
 8       17  1852 Franklin P… Dem.       0.858       0.508         0.0695 1.61e6
 9       18  1856 James Buch… Dem.       0.588       0.453         0.122  1.84e6
10       19  1860 Abraham Li… Rep.       0.594       0.396         0.101  1.86e6
# ℹ 39 more rows
# ℹ 11 more variables: margin <int>, runner_up <chr>, ru_part <chr>,
#   turnout_pct <dbl>, winner_lname <chr>, winner_label <chr>, ru_lname <chr>,
#   ru_label <chr>, two_term <lgl>, ec_votes <dbl>, ec_denom <dbl>
```

We’ll draw a plot like this



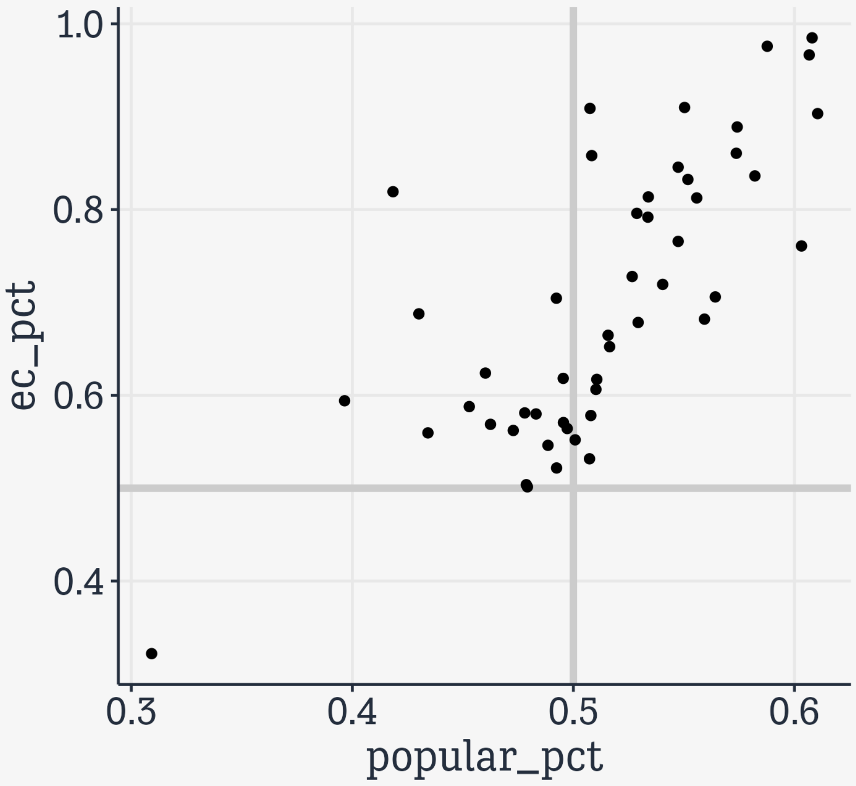
Presidential elections

Keep things neat

```r
## The packages we'll use in addition to ggplot
library(ggrepel) #<<
library(scales) #<<

p_title <- "Presidential Elections: Popular & Electoral College Margins"
p_subtitle <- "1824-2016"
p_caption <- "Data for 2016 are provisional."
x_label <- "Winner's share of Popular Vote"
y_label <- "Winner's share of Electoral College Votes"
```

Base Layer, Lines, Points

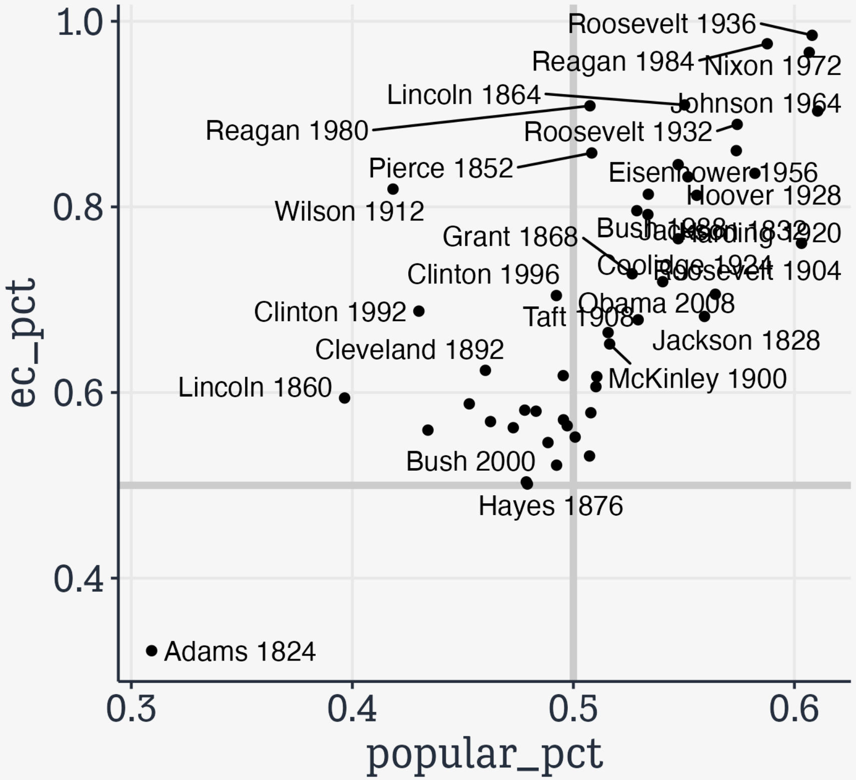
```r
p <- ggplot(data = elections_historic,
            mapping = aes(x = popular_pct,
                          y = ec_pct,
                          label = winner_label))

p + geom_hline(yintercept = 0.5,
               linewidth = 1.4,
               color = "gray80") +
    geom_vline(xintercept = 0.5,
               linewidth = 1.4,
               color = "gray80") +
    geom_point()
```



Add the labels

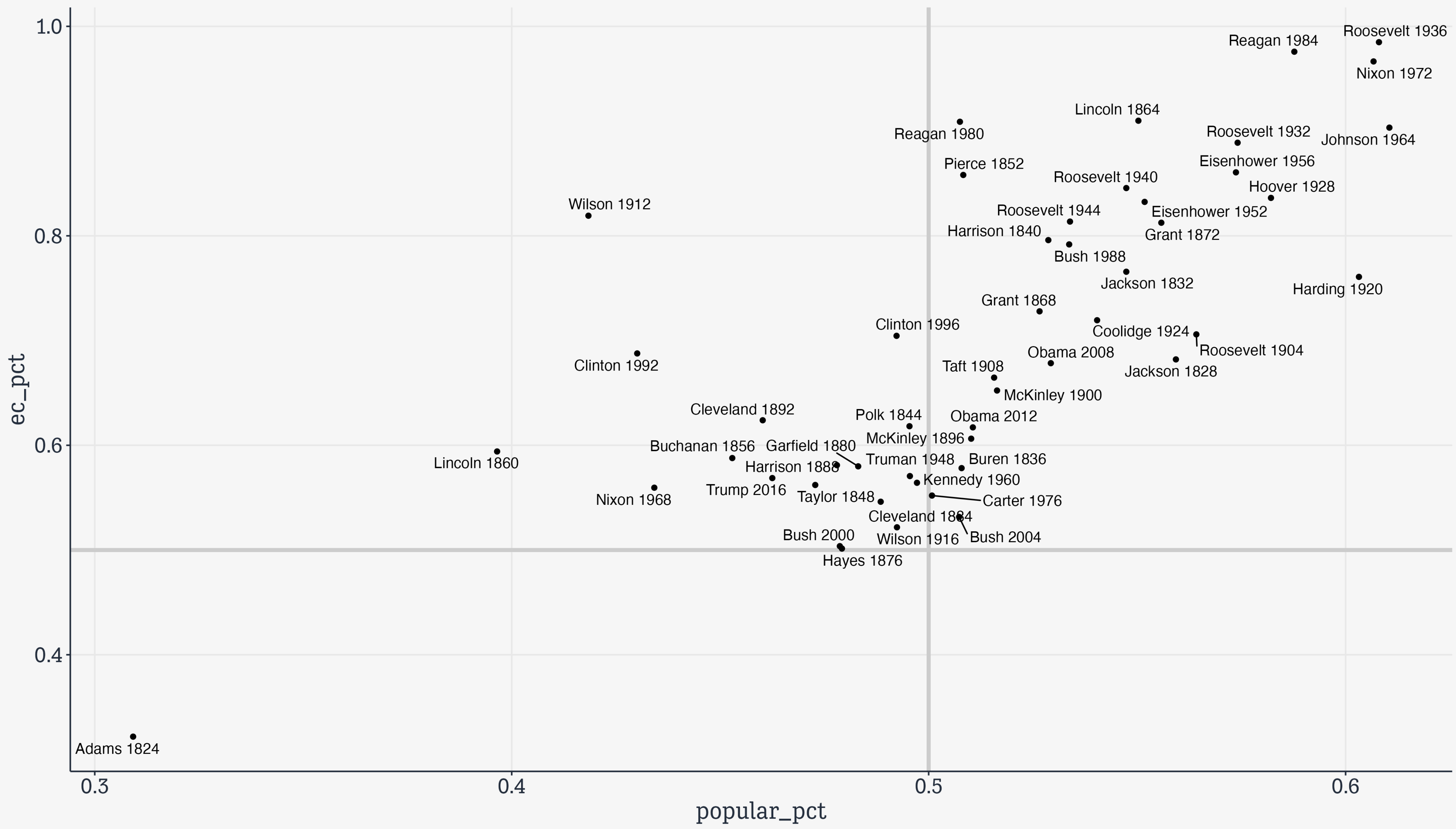
This looks terrible here because `geom_text_repel()` uses the dimensions of the available graphics device to iteratively figure out the labels. Let's allow it to draw on the whole slide.

```
p ← ggplot(data = elections_historic,
           mapping = aes(x = popular_pct,
                         y = ec_pct,
                         label = winner_label))

p + geom_hline(yintercept = 0.5,
               linewidth = 1.4, color = "gray80
  geom_vline(xintercept = 0.5,
             linewidth = 1.4, color = "gray80")
  geom_point() +
  geom_text_repel()
```



Labeling is with respect to the plot size

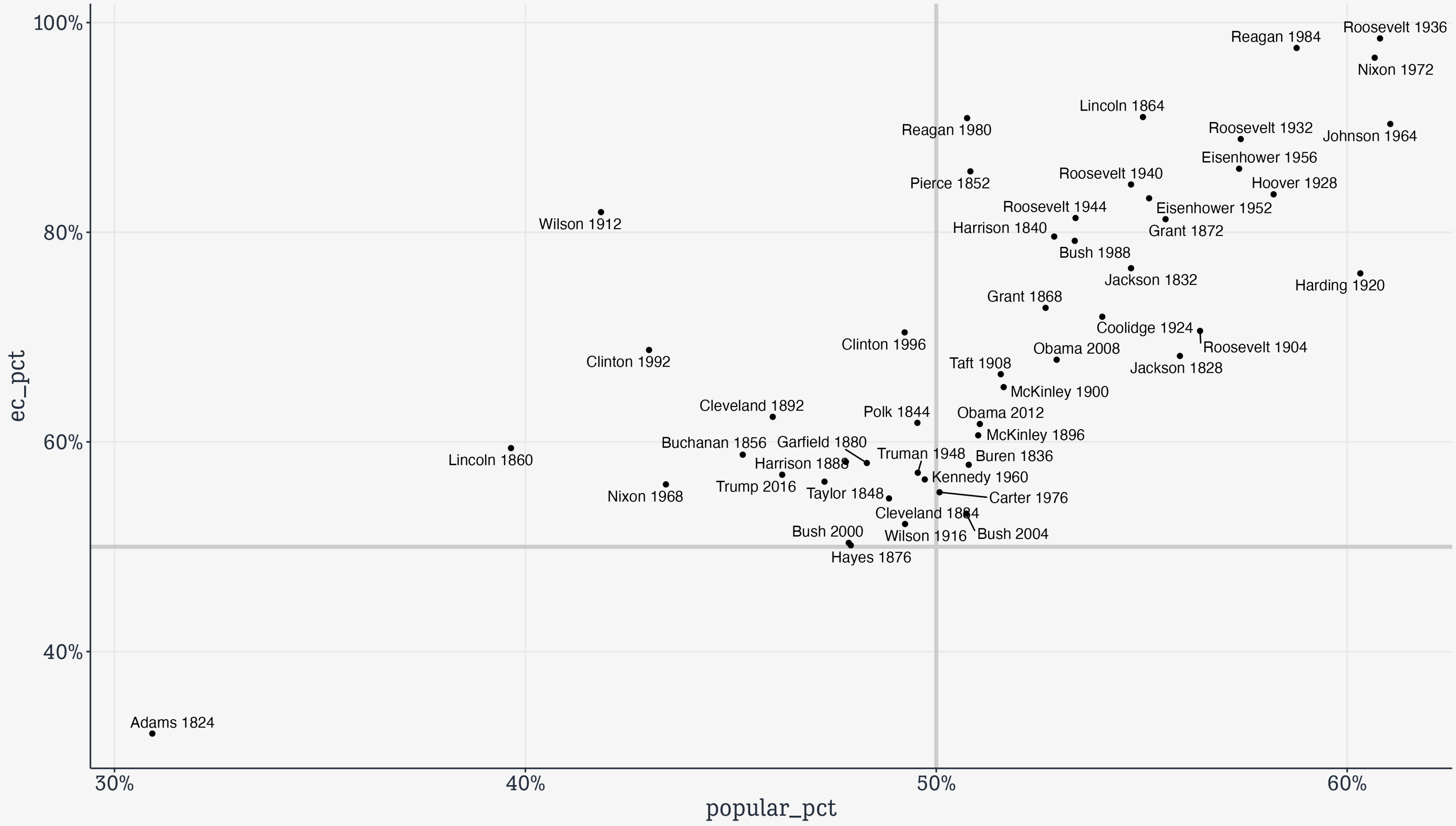
```
p ← ggplot(data = elections_historic,
           mapping  = aes(x = popular_pct,
                          y = ec_pct,
                          label = winner_label))

p_out ← p +
  geom_hline(yintercept = 0.5,
             linewidth = 1.4,
             color = "gray80") +
  geom_vline(xintercept = 0.5,
             linewidth = 1.4,
             color = "gray80") +
  geom_point() +
  geom_text_repel() #<<
```



ec_pct
popular_pct
1.0
0.8
0.6
0.4
0.3
0.4
0.5
0.6
Roosevelt 1936
Reagan 1984
Nixon 1972
Lincoln 1864
Reagan 1980
Roosevelt 1932
Johnson 1964
Pierce 1852
Eisenhower 1956
Roosevelt 1940
Hoover 1928
Wilson 1912
Roosevelt 1944
Eisenhower 1952
Harrison 1840
Grant 1872
Bush 1988
Jackson 1832
Harding 1920
Grant 1868
Clinton 1996
Coolidge 1924
Roosevelt 1904
Obama 2008
Clinton 1992
Taft 1908
Jackson 1828
McKinley 1900
Cleveland 1892
Polk 1844
Obama 2012
McKinley 1896
Buchanan 1856
Garfield 1880
Lincoln 1860
Harrison 1888
Truman 1948
Buren 1836
Kennedy 1960
Trump 2016
Nixon 1968
Taylor 1848
Carter 1976
Cleveland 1884
Bush 2000
Wilson 1916
Bush 2004
Hayes 1876
Adams 1824

Adjust the Scales

```
p ← ggplot(data = elections_historic,
            mapping  = aes(x = popular_pct,
                           y = ec_pct,
                           label = winner_label))
p_out ← p + geom_hline(yintercept = 0.5,
                        linewidth = 1.4,
                        color = "gray80") +
    geom_vline(xintercept = 0.5,
               linewidth = 1.4,
               color = "gray80") +
    geom_point() +
    geom_text_repel() +
    scale_x_continuous(labels = label_percent()) + #<<
    scale_y_continuous(labels = label_percent()) #<<
```



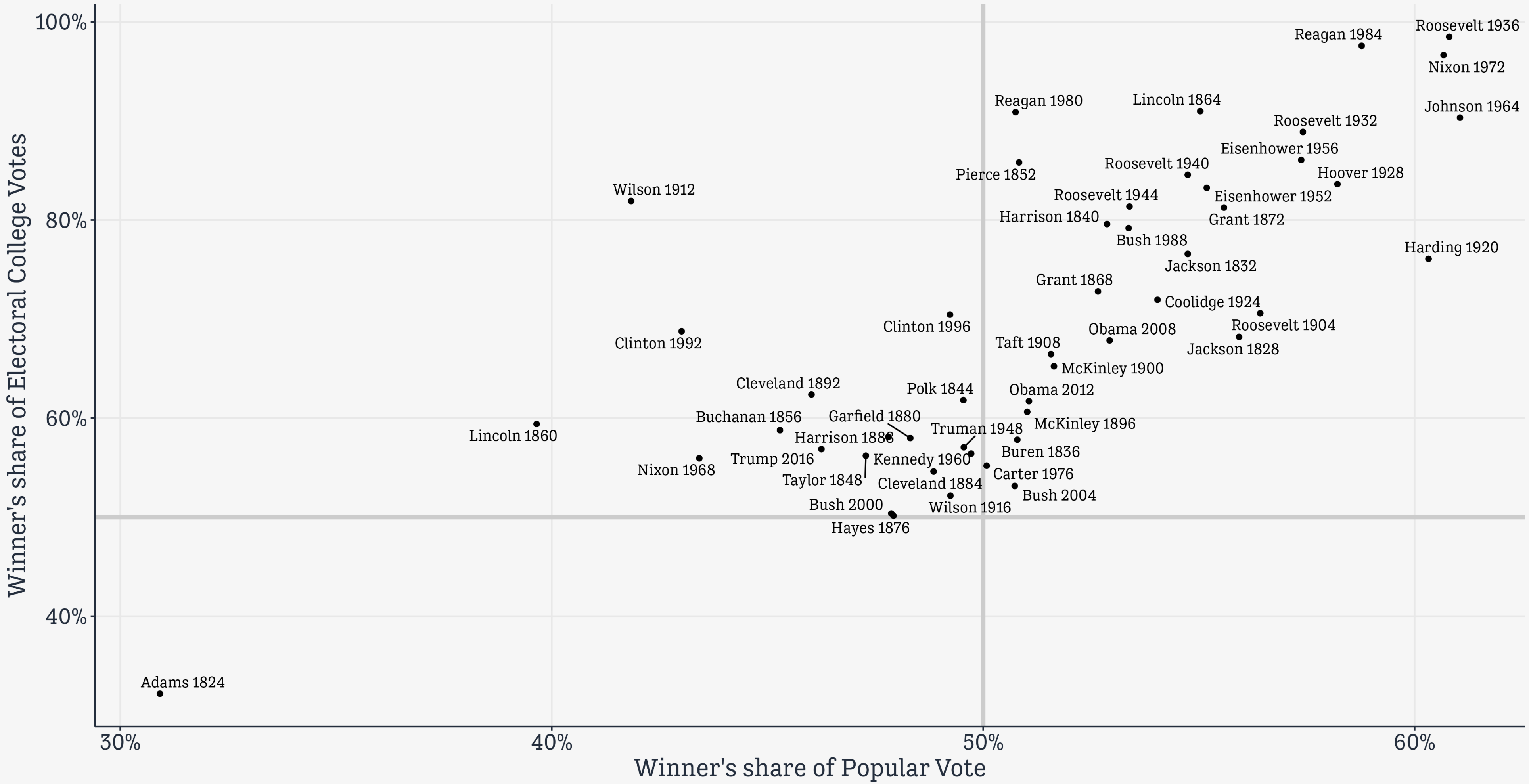
ec_pct
popular_pct
100%
80%
60%
40%
30%
40%
50%
60%
Roosevelt 1936
Reagan 1984
Nixon 1972
Lincoln 1864
Reagan 1980
Roosevelt 1932
Johnson 1964
Eisenhower 1956
Pierce 1852
Roosevelt 1940
Hoover 1928
Wilson 1912
Roosevelt 1944
Eisenhower 1952
Harrison 1840
Grant 1872
Bush 1988
Jackson 1832
Harding 1920
Grant 1868
Coolidge 1924
Clinton 1996
Roosevelt 1904
Obama 2008
Clinton 1992
Taft 1908
Jackson 1828
McKinley 1900
Cleveland 1892
Polk 1844
Obama 2012
McKinley 1896
Buchanan 1856
Garfield 1880
Lincoln 1860
Truman 1948
Buren 1836
Harrison 1888
Kennedy 1960
Trump 2016
Nixon 1968
Taylor 1848
Carter 1976
Cleveland 1884
Bush 2000
Bush 2004
Wilson 1916
Hayes 1876
Adams 1824

Add the labels

```r
p <- ggplot(data = elections_historic,
            mapping  = aes(x = popular_pct,
                           y = ec_pct,
                           label = winner_label))
p_out <- p + geom_hline(yintercept = 0.5,
                        linewidth = 1.4,
                        color = "gray80") +
  geom_vline(xintercept = 0.5,
             linewidth = 1.4,
             color = "gray80") +
  geom_point() +
  geom_text_repel(mapping = aes(family = "Tenso Slide")) +#<<
  scale_x_continuous(labels = label_percent()) +
  scale_y_continuous(labels = label_percent()) +
  labs(x = x_label, y = y_label,  #<<
       title = p_title,
       subtitle = p_subtitle,
       caption = p_caption)
```

Presidential Elections: Popular & Electoral College Margins

1824-2016

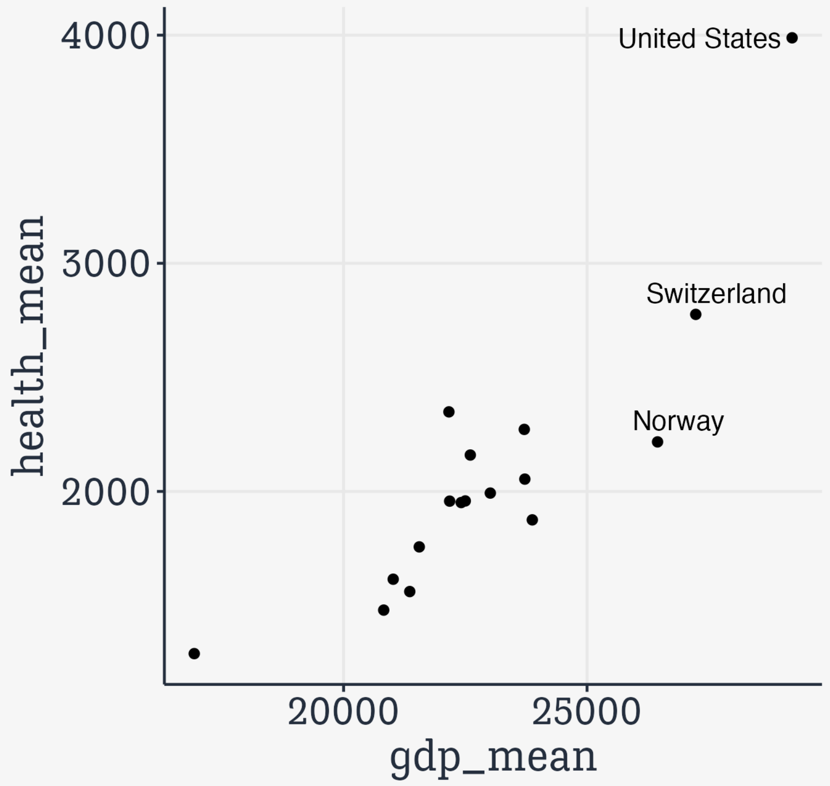


Data for 2016 are provisional.

Labeling points of interest

Option 1: On the fly in ggplot

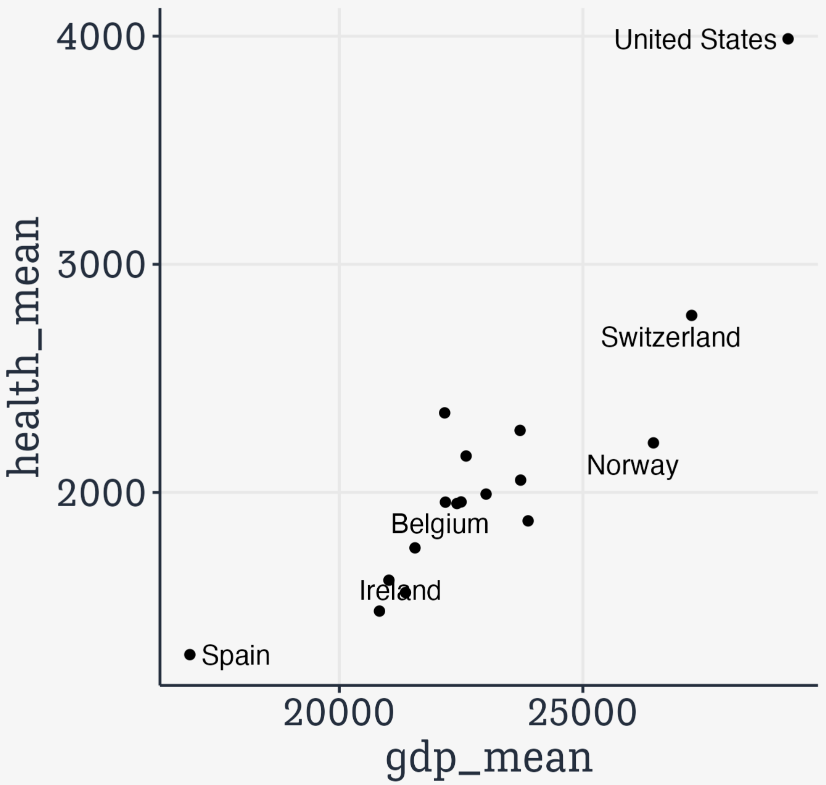
```
by_country ▷
  ggplot(mapping = aes(x = gdp_mean,
                       y = health_mean)) +
  geom_point() +
  geom_text_repel(data = subset(by_country, gdp_
                  mapping = aes(label = country
```



Option 1: On the fly inside ggplot

Stuffing everything into the subset() call might get messy

```
by_country ▷
  ggplot(mapping = aes(x = gdp_mean,
                       y = health_mean)) +
  geom_point() +
  geom_text_repel(data = subset(by_country,
                                gdp_mean > 2500
                                  health_mean <
                                  country %in%
                mapping = aes(label = country
```



Option 2: Use dplyr first

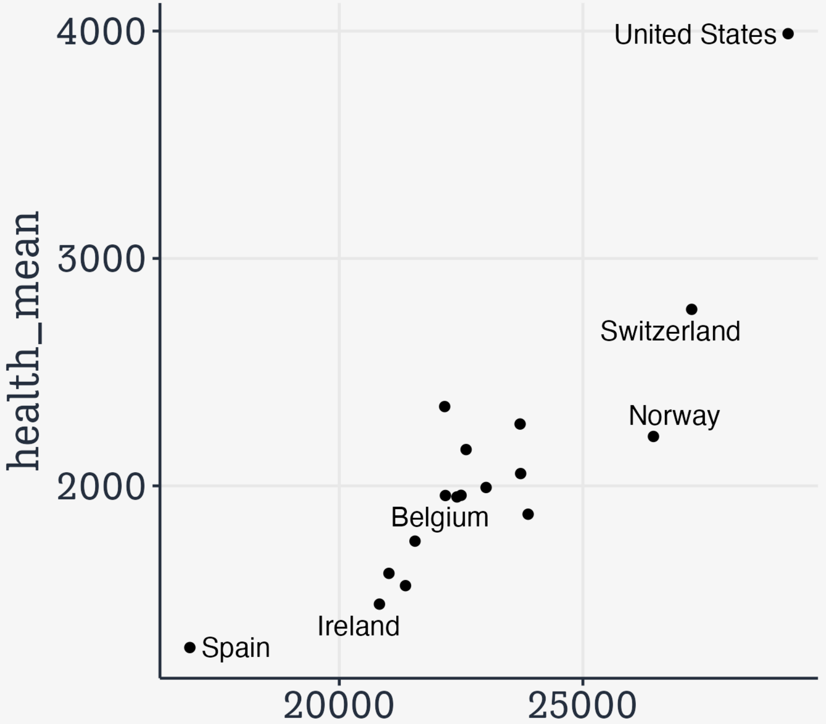
```
df_hl <- by_country |>
  filter(gdp_mean > 25000 |
           health_mean < 1500 |
           country %in% "Belgium")

df_hl
```

```
# A tibble: 6 × 28
  consent_law country       donors_mean donors_sd pop_mean  pop_sd pop_dens_mean
  <chr>       <chr>               <dbl>     <dbl>    <dbl>   <dbl>         <dbl>
1 Informed    Ireland              19.8      2.48    3674.   132.           5.23
2 Informed    United States        20.0      1.33  269330. 12545.           2.80
3 Presumed    Belgium              21.9      1.94   10153.   109.          30.7 
4 Presumed    Norway               15.4      1.11    4386.    97.3          1.35
5 Presumed    Spain                28.1      4.96   39666.   951.           7.84
6 Presumed    Switzerland          14.2      1.71    7037.   170.          17.0 
# ℹ 21 more variables: pop_dens_sd <dbl>, gdp_mean <dbl>, gdp_sd <dbl>,
#   gdp_lag_mean <dbl>, gdp_lag_sd <dbl>, health_mean <dbl>, health_sd <dbl>,
#   health_lag_mean <dbl>, health_lag_sd <dbl>, pubhealth_mean <dbl>,
#   pubhealth_sd <dbl>, roads_mean <dbl>, roads_sd <dbl>, cerebvas_mean <dbl>,
#   cerebvas_sd <dbl>, assault_mean <dbl>, assault_sd <dbl>,
#   external_mean <dbl>, external_sd <dbl>, txp_pop_mean <dbl>,
#   txp_pop_sd <dbl>
```

Option 2: Use `dplyr` first

This makes things neater. A `geom` can be fully “autonomous”. Each one can have its own `mapping` call *and* its own `data` source. This can be very useful when building up plots overlaying several sources or subsets of data.

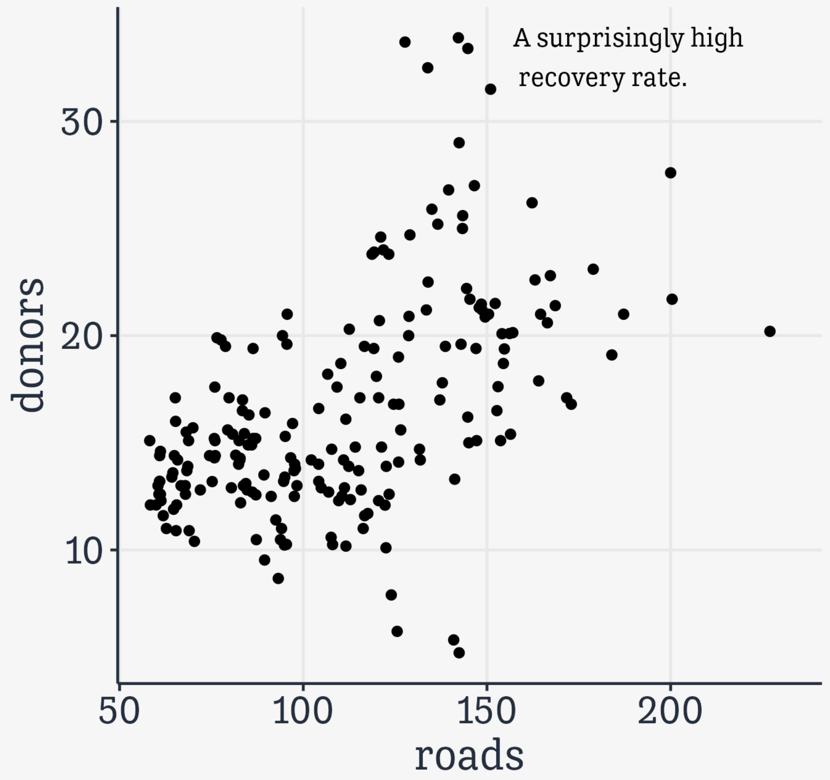
```
by_country ▷
  ggplot(mapping = aes(x = gdp_mean,
                       y = health_mean)) +
  geom_point() +
  geom_text_repel(data = df_hl,
                  mapping = aes(label = country
```



Write and draw
inside the plot area

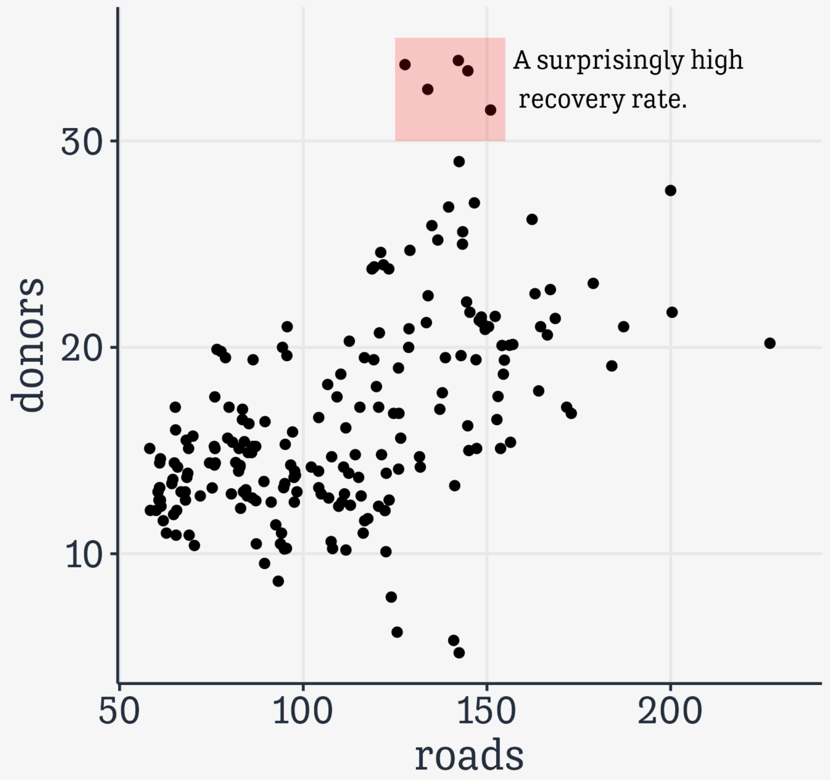
annotate() can imitate geoms

```
organdata ▷
  ggplot(mapping = aes(x = roads,
                       y = donors)) +
  geom_point() +
  annotate(geom = "text",
           family = "Tenso Slide",
           x = 157,
           y = 33,
           label = "A surprisingly high \n reco
           hjust = 0)
```



annotate() can imitate geoms

```r
organdata ▷
  ggplot(mapping = aes(x = roads,
                       y = donors)) +
  geom_point() +
  annotate(geom = "rect",
           xmin = 125, xmax = 155,
           ymin = 30, ymax = 35,
           fill = "red",
           alpha = 0.2) +
  annotate(geom = "text",
           x = 157, y = 33,
           family = "Tenso Slide",
           label = "A surprisingly high \n reco
           hjust = 0)
```



Scales, Guides, and Themes

Every mapped variable has a scale

Aesthetic mappings link quantities or categories in your data to things you can see on the graph. Thus, they have a scale associated with that representation.

Scale functions manage this relationship. Remember: not just `x` and `y` but also `color`, `fill`, `shape`, `size`, and `alpha` are scales.

If it can represent your data, it has a scale, and a *scale function* to manage it.

This means you control things like color schemes *for data mappings* through scale functions

Because those colors are representing features of your data.

Naming conventions for scale functions

In general, scale functions are named like this:

scale_<MAPPING>_<KIND>()

Naming conventions

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scale_<MAPPING>_<KIND>()

We already know there are a lot of **mappings**

x, y, color, size, shape, and so on.

Naming conventions

In general, scale functions are named like this:

scale_<MAPPING>_<KIND>()

We already know there are a lot of **mappings**

x, y, color, size, shape, and so on.

And there are many **kinds** of scale as well.

discrete, continuous, log10, date, binned, and many others.

So there's a whole zoo of scale functions.

The naming convention helps us keep track.

Naming conventions

scale_<MAPPING>_<KIND>()

scale_x_continuous()

scale_y_continous()

scale_x_discrete()

scale_y_discrete()

scale_x_log10()

scale_x_sqrt()

Naming conventions

scale_<MAPPING>_<KIND>()

scale_x_continuous()

scale_y_continous()

scale_x_discrete()

scale_y_discrete()

scale_x_log10()

scale_x_sqrt()

scale_color_discrete()

scale_color_gradient()

scale_color_gradient2()

scale_color_brewer()

scale_fill_discrete()

scale_fill_gradient()

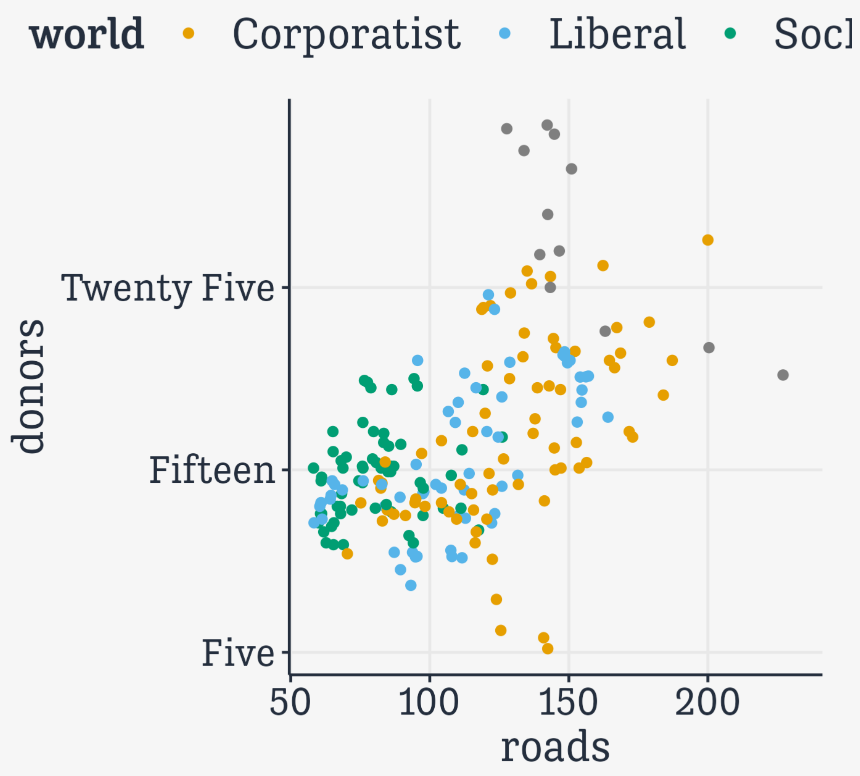
scale_fill_gradient2()

scale_fill_brewer()

Scale functions in practice

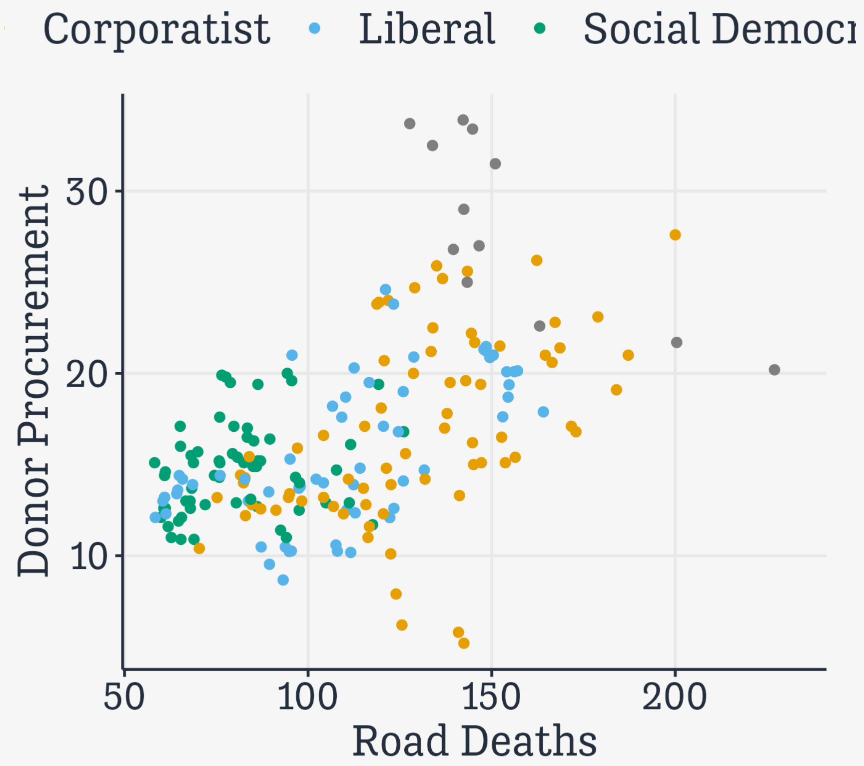
Scale functions take arguments appropriate to their mapping and kind

```
organdata |>
  ggplot(mapping = aes(x = roads,
                       y = donors,
                       color = world)) +
  geom_point() +
  scale_y_continuous(breaks = c(5, 15, 25),
                     labels = c("Five",
                                "Fifteen",
                                "Twenty Five"))
```



More usefully ...

```r
organdata |>
  ggplot(mapping = aes(x = roads,
                       y = donors,
                       color = world)) +
  geom_point() +
  scale_color_discrete(labels =
                         c("Corporatist",
                           "Liberal",
                           "Social Democratic",
                           "Unclassified")) +
  labs(x = "Road Deaths",
       y = "Donor Procurement",
       color = "Welfare State")
```



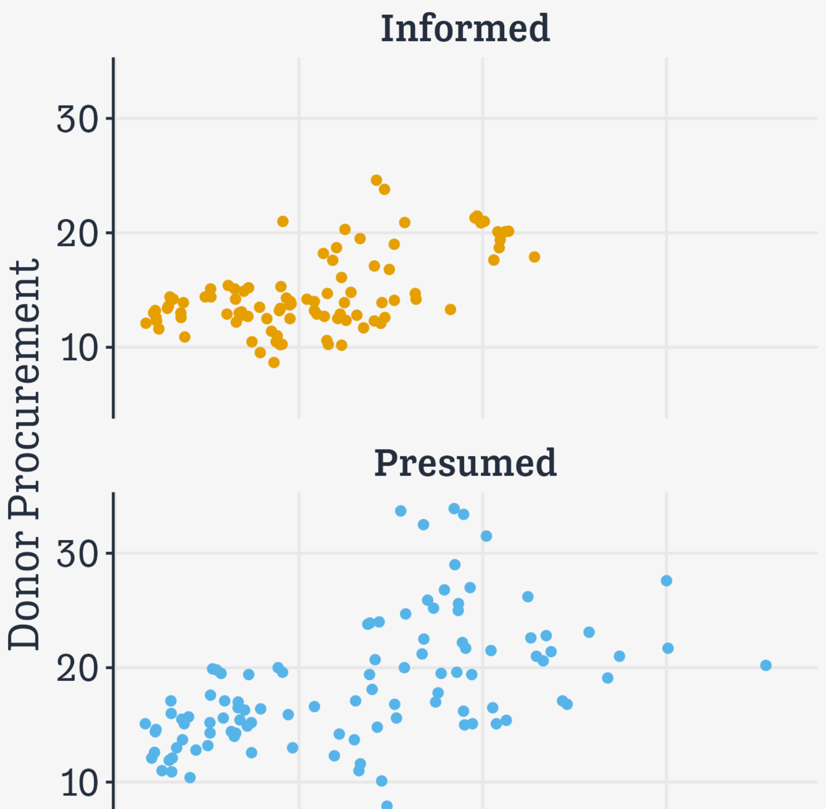
The `guides()` function

Control overall properties of the guide labels.

Common use: turning it off.

We'll see more advanced uses later.

```r
organdata |>
  ggplot(mapping = aes(x = roads,
                       y = donors,
                       color = consent_law)) +
  geom_point() +
  facet_wrap(~ consent_law, ncol = 1) +
  guides(color = "none") +
  labs(x = "Road Deaths",
       y = "Donor Procurement")
```



The `theme()` function

`theme()` styles parts of your plot that are *not* directly representing your data. Often the first thing people want to adjust; but logically it's the *last* thing.

```
## Using the "classic" ggplot theme here
organdata ▷
  ggplot(mapping = aes(x = roads,
                       y = donors,
                       color = consent_law)) +
  geom_point() +
  labs(title = "By Consent Law",
    x = "Road Deaths",
    y = "Donor Procurement",
    color = "Legal Regime:") +
  theme(legend.position = "bottom",
        plot.title = element_text(color = "dark
                                  face = "bold"
```