

Colors/Themes + Inclusive Design Principles

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Week 4

Agenda

- Learn about C.R.A.P
- Color basics
 - 3 basic ways color is used
- Color blindness
- Some common problems with color use
- Quick discussion of palettes
- Quick discussion of themes
- Lab 4
- Q & A with Dr. Daniel Anderson, who pioneered this EDS specialization and class!

Learning Objectives

- Learn a bit more about good/inclusive design principles
- Understand different types of color palettes
 - ...and when you should use one versus another
- Understand and be able to effectively evaluate concerns related to color blindness
- Be able to fluently change colors/fills within ggplot

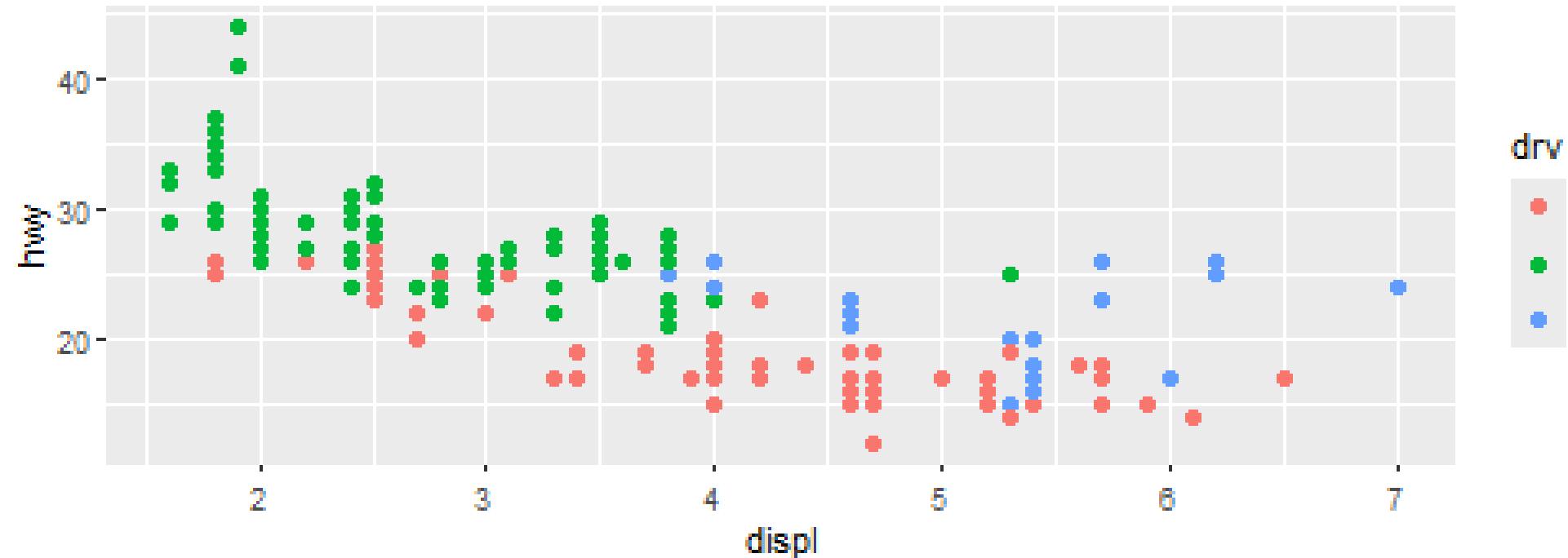
Universal principles

- Contrast
- Repetition
- Alignment
- Proximity
 - Pioneered by Robin Patricia Williams

- These design principles apply everywhere!
- Graphic design, art, music, architecture... and graphs!

Is that gray background okay?

```
theme_set(theme_gray())
ggplot(mpg, aes(x = displ, y = hwy, color = drv)) +
  geom_point(size = 2)
```



```
theme(gray)
```

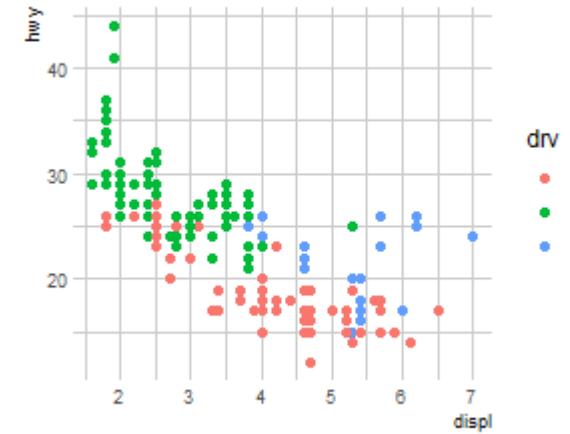
Applying CRAP to ggplot

- We can follow CRAP principles to make big improvements to our plots
- Claus Wilke's chapter covers lots of these graph-specific principles
- We can apply these principles to ggplot plots

Like this!

```
library(hrbrthemes)

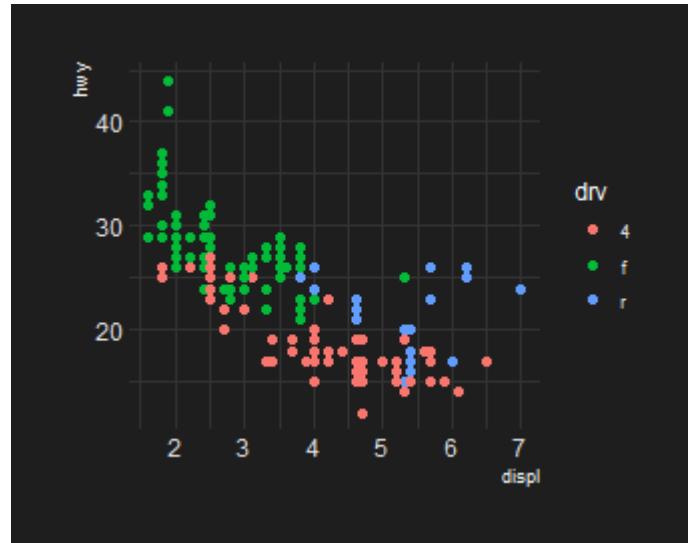
ggplot(mpg, aes(x = displ, y = hwy,
                 color = drv)) +
  geom_point(size = 2) +
  theme_ipsum_ps()
```



And this!

```
library(hrbrthemes)

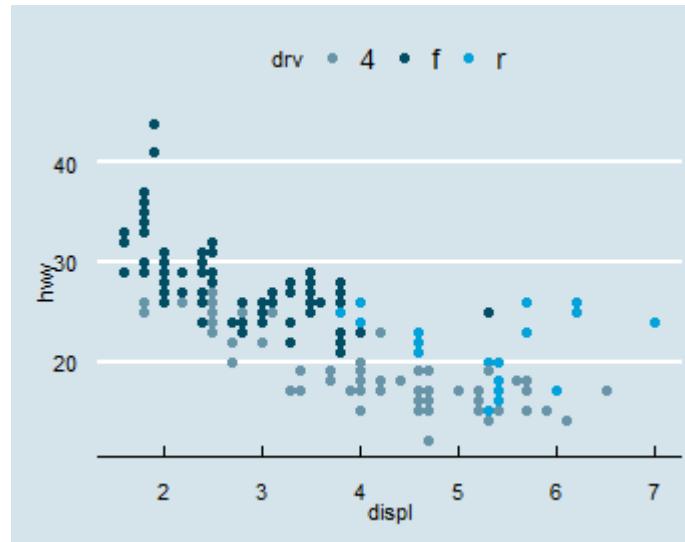
ggplot(mpg, aes(x = displ, y = hwy,
                 color = drv)) +
  geom_point(size = 2) +
  theme_modern_rc()
```



Or this!

```
library(ggthemes)

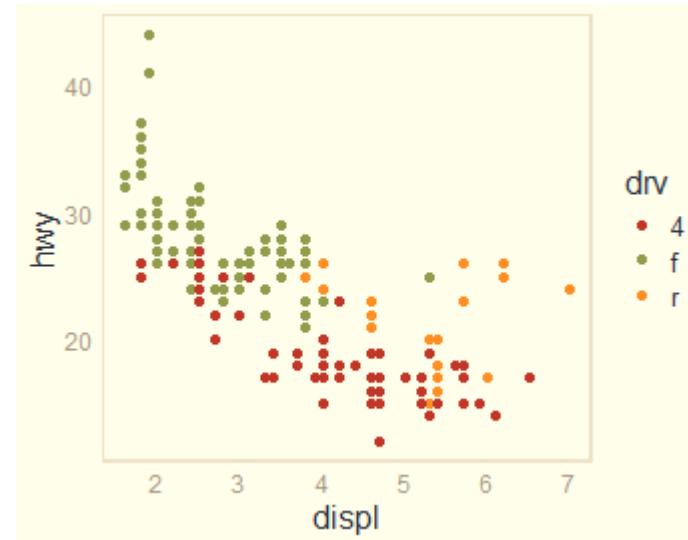
ggplot(mpg, aes(x = displ, y = hwy,
                 color = drv)) +
  geom_point(size = 2) +
  scale_color_economist() +
  theme_economist()
```



And even this!

```
#remotes::install_github("gadenbuie/ggpomological")
library(ggpomological)

ggplot(mpg, aes(x = displ, y = hwy,
                 color = drv)) +
  geom_point(size = 2) +
  scale_color_pomological() +
  theme_pomological_fancy()
```



One magic, powerful function

`theme()`

- Depending on time we come back to an anatomy of `theme` the end today or next week!

Before we get too deep into themes of which colors is an essential element...

Some very practical advice about colors

- Keep straight when color is mapped to a variable through `aes` and when it's modifying an element overall
 - Former requires `scale_color_*` or `scale_fill_*` while the latter does not
- Keep straight colors and fills (see former bullet)
- Use advice of others to your advantage (e.g., <http://colorbrewer2.org/>)



Why color choice matters



Why color choice matters

Another quick example

{rayshader}

3 fundamental uses of color

1. Distinguish groups from each other
2. Represent data values
3. Highlight

Color as a tool to distinguish

Discrete items

- Often no intrinsic order

Qualitative color scale

- Finite number of colors
 - Chosen to maximize distinctness, while also be equivalent
 - Equivalent
 - No color should stand out
 - No impression of order

Some Examples

Okabe Ito



ColorBrewer Dark2



ggplot2 hue



See more about the Okabe Ito palette origins [here](#)

How do we use them?

Imagine we have data like this

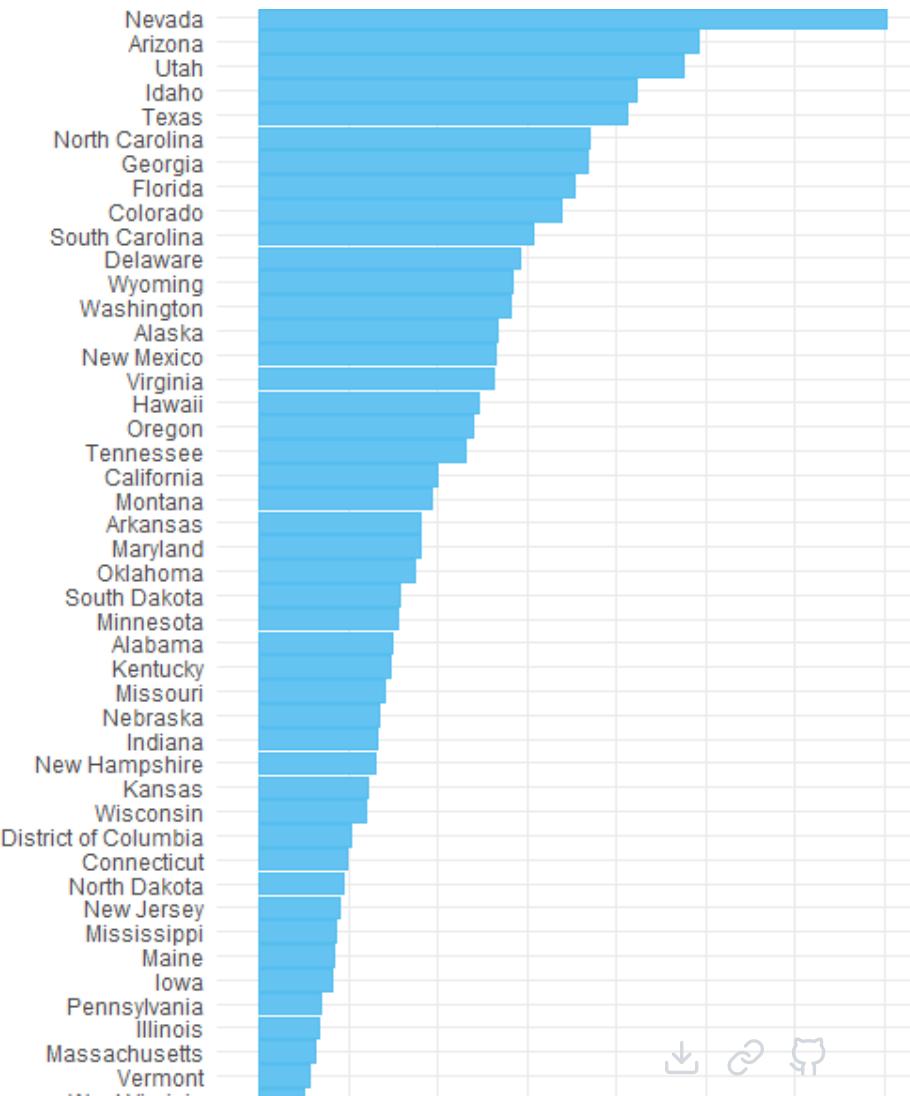
```
popgrowth_df
```



```
## # A tibble: 51 × 8
##   region division           state    pop2000    pop2010  popgrowth     area fill_color
##   <fct>  <chr>            <fct>    <dbl>     <dbl>      <dbl>    <dbl> <chr>
## 1 West    Mountain         Nevada    1998257   2700551     0.351 109781. West
## 2 West    Mountain         Arizona   5130632   6392017     0.246 113594. West
## 3 West    Mountain         Utah      2233169   2763885     0.238  82170. West
## 4 West    Mountain         Idaho     1293953   1567582     0.211  82643. West
## 5 South   West South Central Texas    20851820  25145561     0.206 261232. South
## 6 South   South Atlantic   North Carolina 8049313   9535483     0.185  48618. South
## # i 45 more rows
```

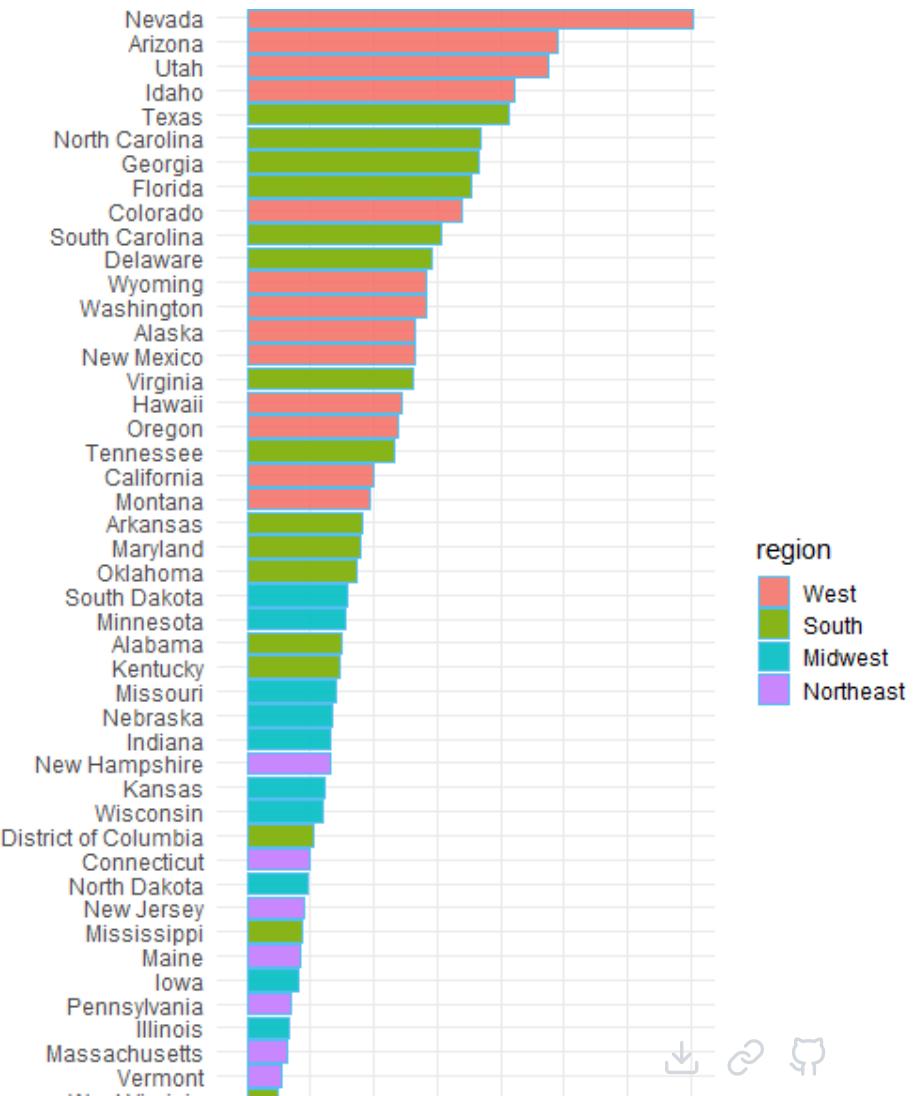
Maybe a plot like this

```
ggplot(popgrowth_df,  
       aes(x = popgrowth,  
            y = state)) +  
  geom_col(alpha = 0.9)
```

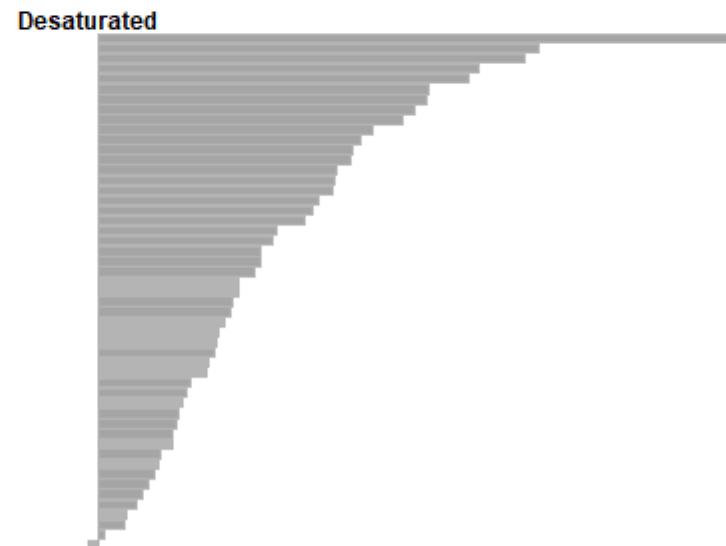
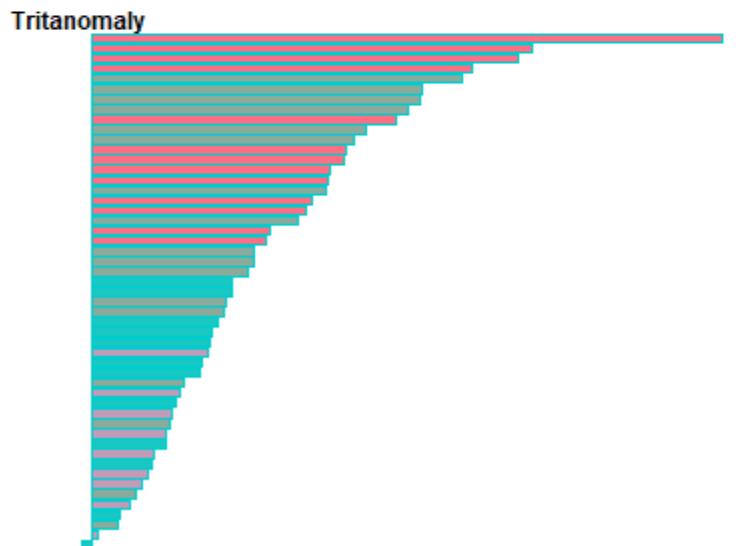
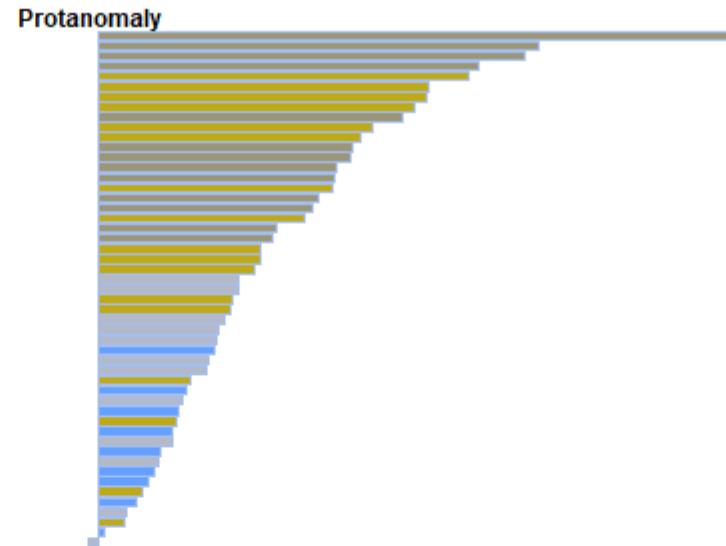
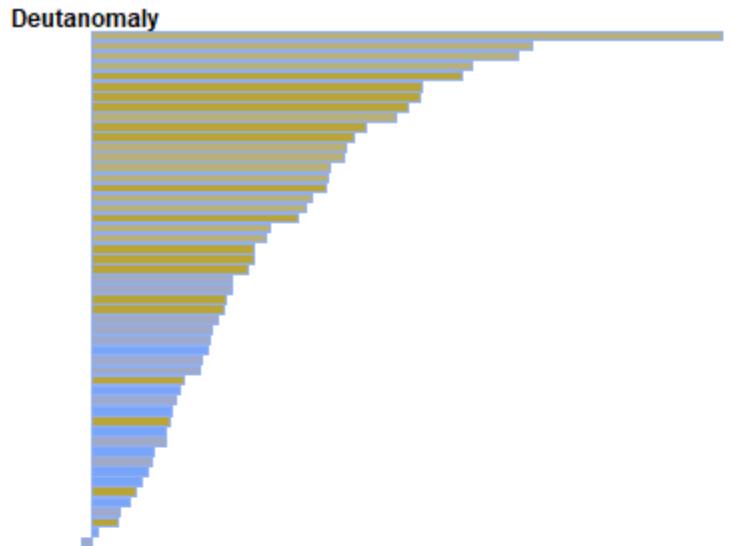


Alternatively, fill by region

```
ggplot(popgrowth_df,  
       aes(x = popgrowth,  
            y = state)) +  
  geom_col(aes(fill = region),  
           alpha = 0.9)
```

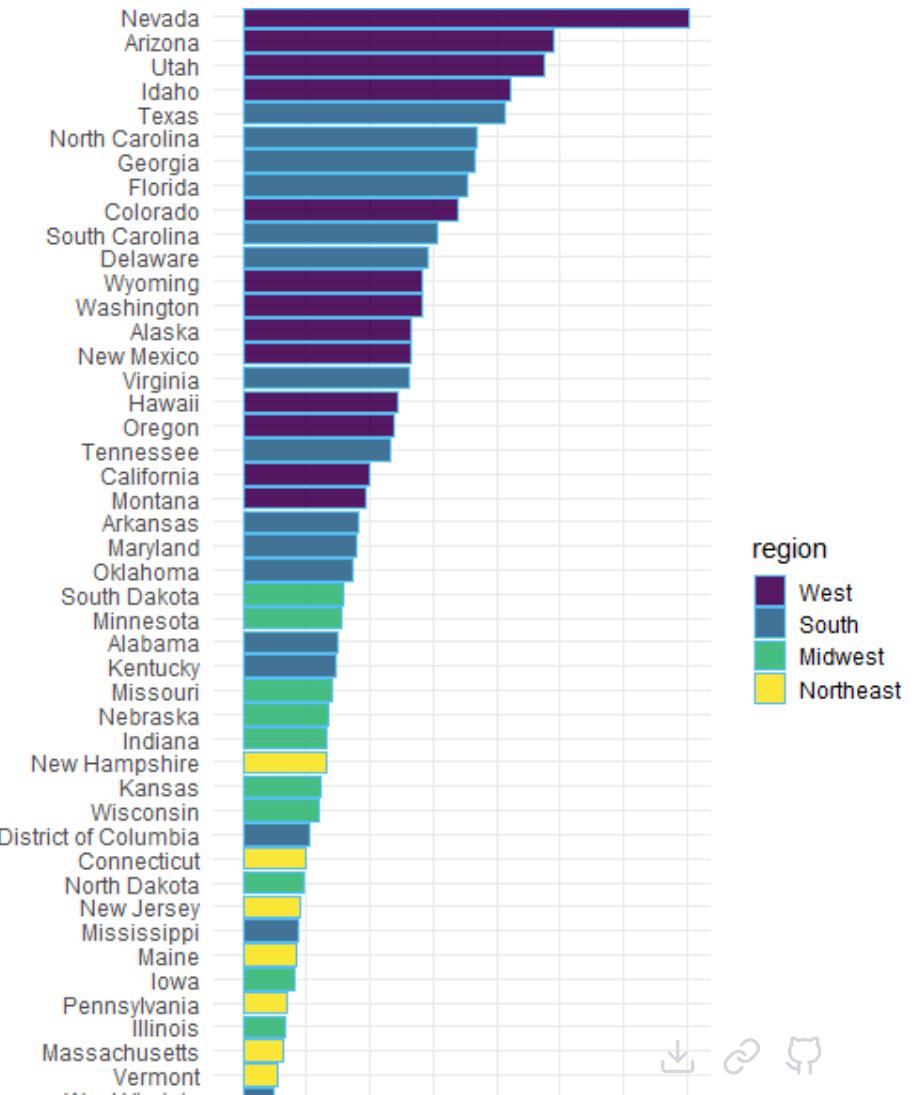


Problem with default palette

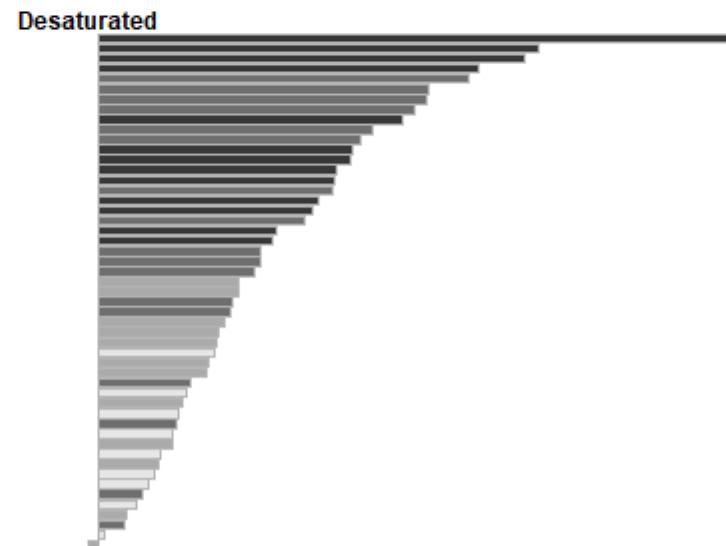
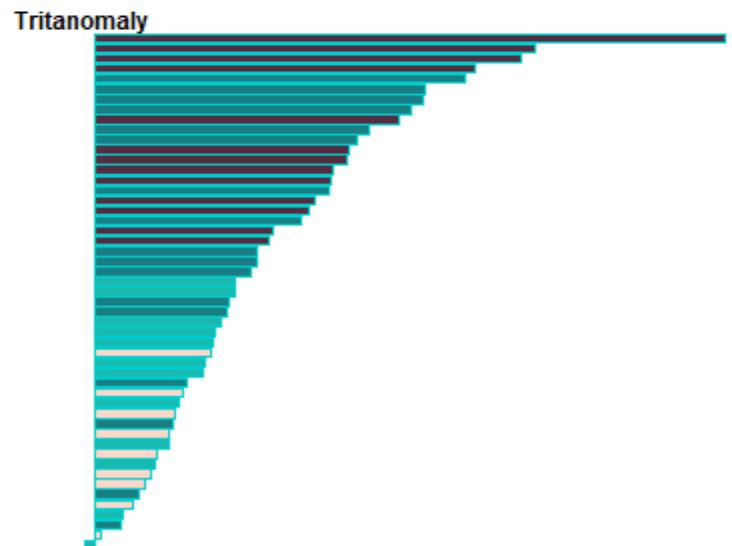
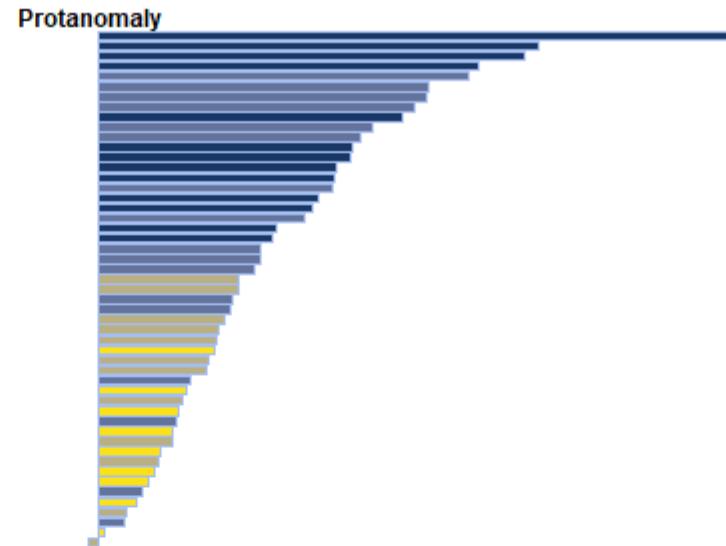
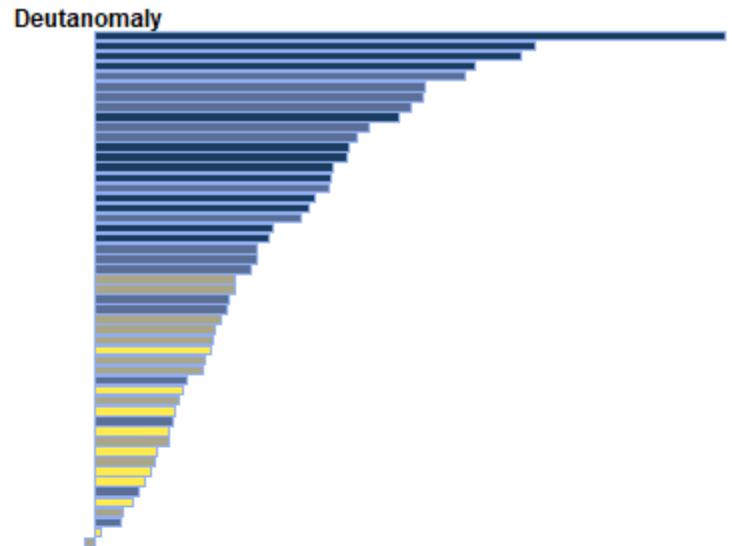


Alternative: viridis

```
ggplot(popgrowth_df,  
       aes(x = popgrowth,  
            y = state)) +  
  geom_col(aes(fill = region),  
           alpha = 0.9) +  
  scale_fill_viridis_d()
```



Revised version

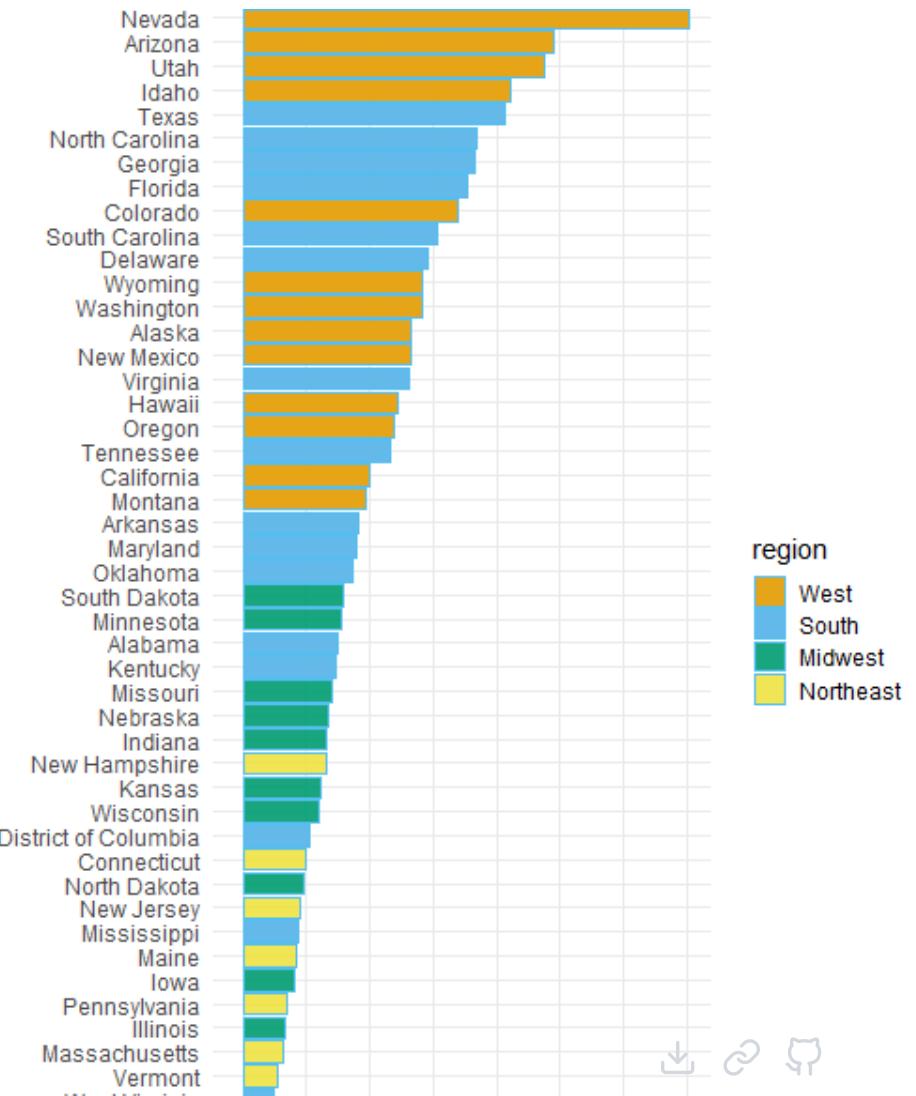


The Okabe Ito palette

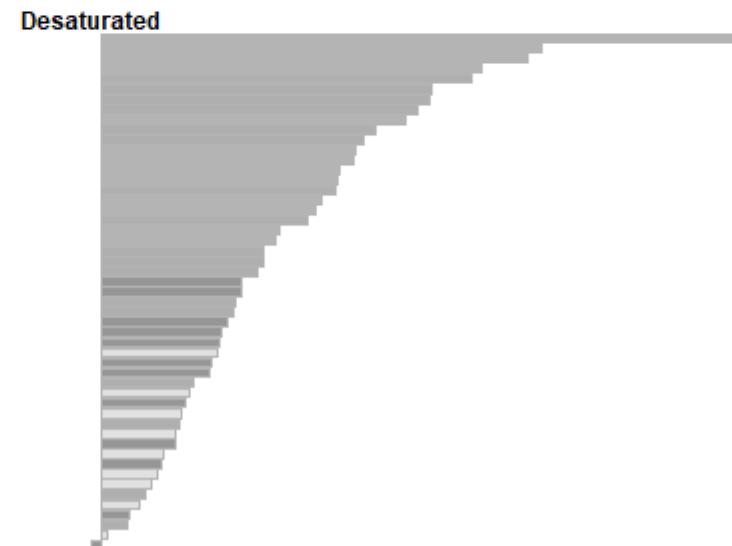
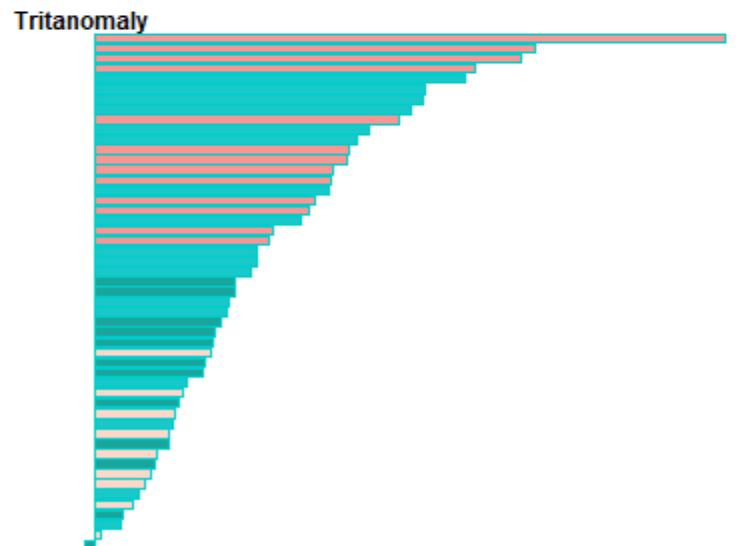
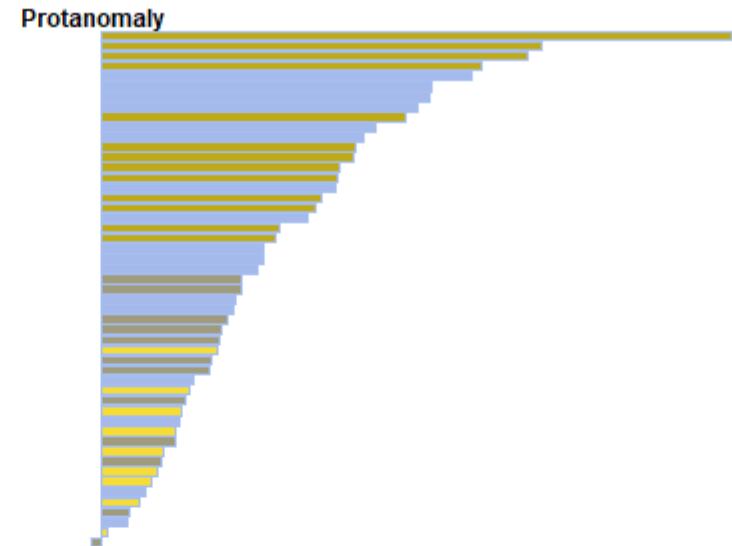
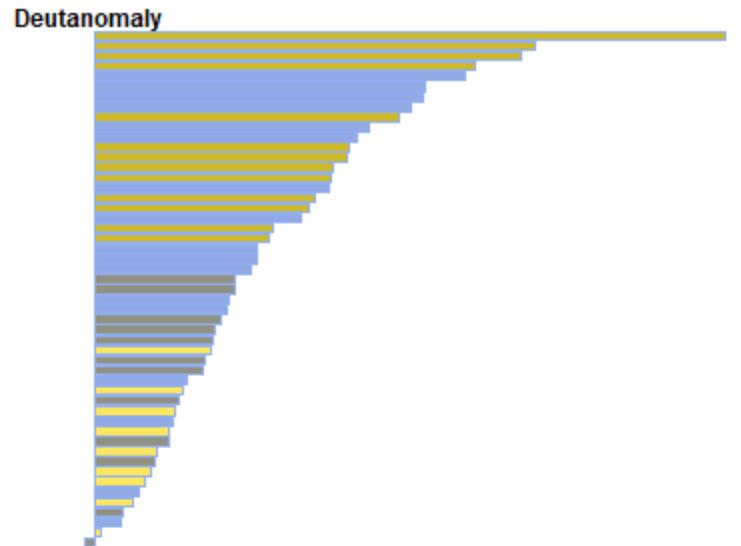
- From Color Universal Design

```
library(colorblindr)
```

```
ggplot(popgrowth_df,  
       aes(x = popgrowth,  
            y = state)) +  
  geom_col(aes(fill = region),  
           alpha = 0.9) +  
  scale_fill_OkabeIto()
```



Okabe Ito for colorblindness



How am I checking for colorblindness?

- Also part of the **{colorblindr}** package ([here](#))
 - depends on the dev versions of **{colorspace}** and **{cowplot}**, which are useful packages in their own right

```
#devtools::install_github("wilkelab/cowplot")
#install.packages("colorspace", repos = "http://R-Forge.R-project.org")

#devtools::install_github("clauswilke/colorblindr")
```

A note on installation

Occasionally people have run into issues with the install on the previous slide.

If that happens, try `install.packages("colorBlindness")` instead, and use the `cvdPlot` function.

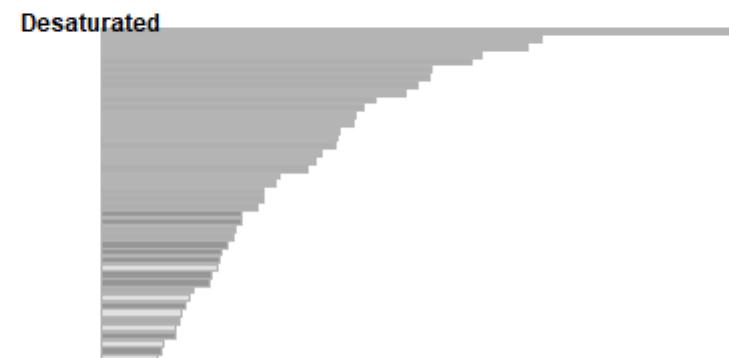
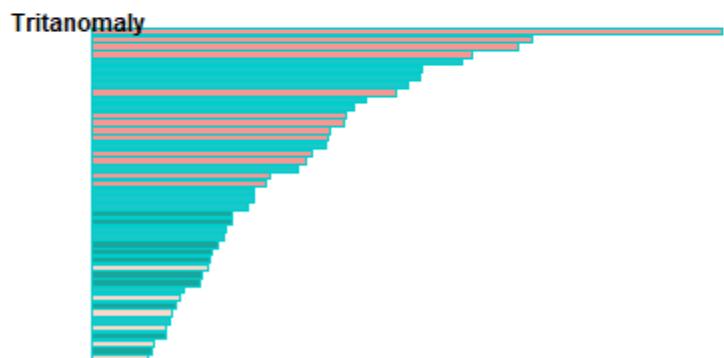
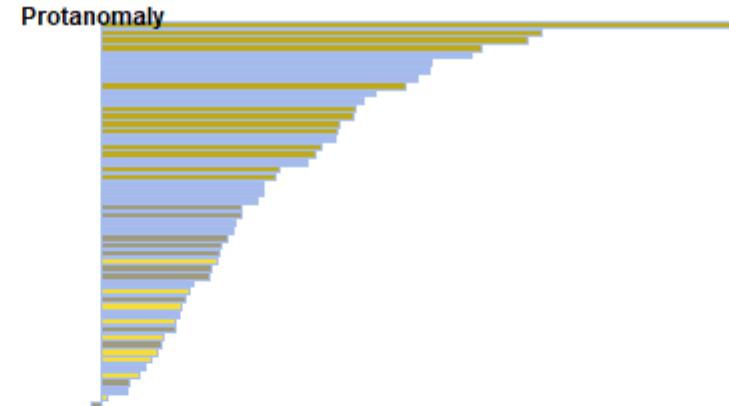
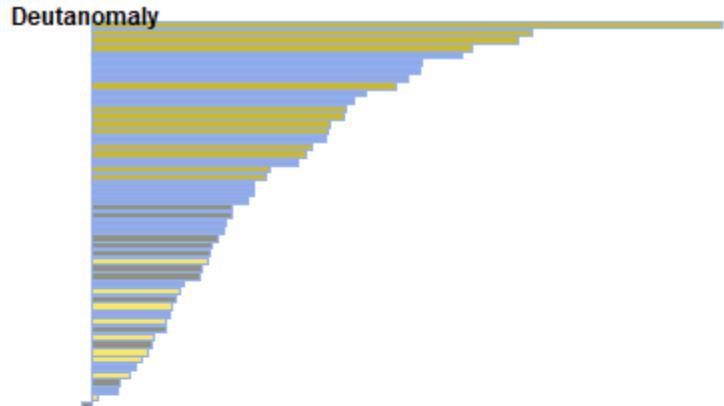
(It's just a modified version of `cvd_grid` from *colorblindr*)

```

p <- ggplot(popgrowth_df,
             aes(x = popgrowth,
                 y = state)) +
  geom_col(aes(fill = region),
           alpha = 0.9) +
  scale_fill_OkabeIto() +
  theme_void() # not necessary but I like it

colorblindr::cvd_grid(p)

```



region
West
South
Midwest
Northeast

region
West
South
Midwest
Northeast



Colors for continuous values

Sequential scale examples

ColorBrewer Blues



Heat



Viridis



Sequential scales

- Which values are larger/smaller
- How distant two values are from each other
 - Scale must be perceptually uniform across its entire range
 - Similar to an interval scale, but for color
- Often based on a single hue
- Multi-hue sequential scales tend to follow gradients in the natural world

Common uses of sequential palettes

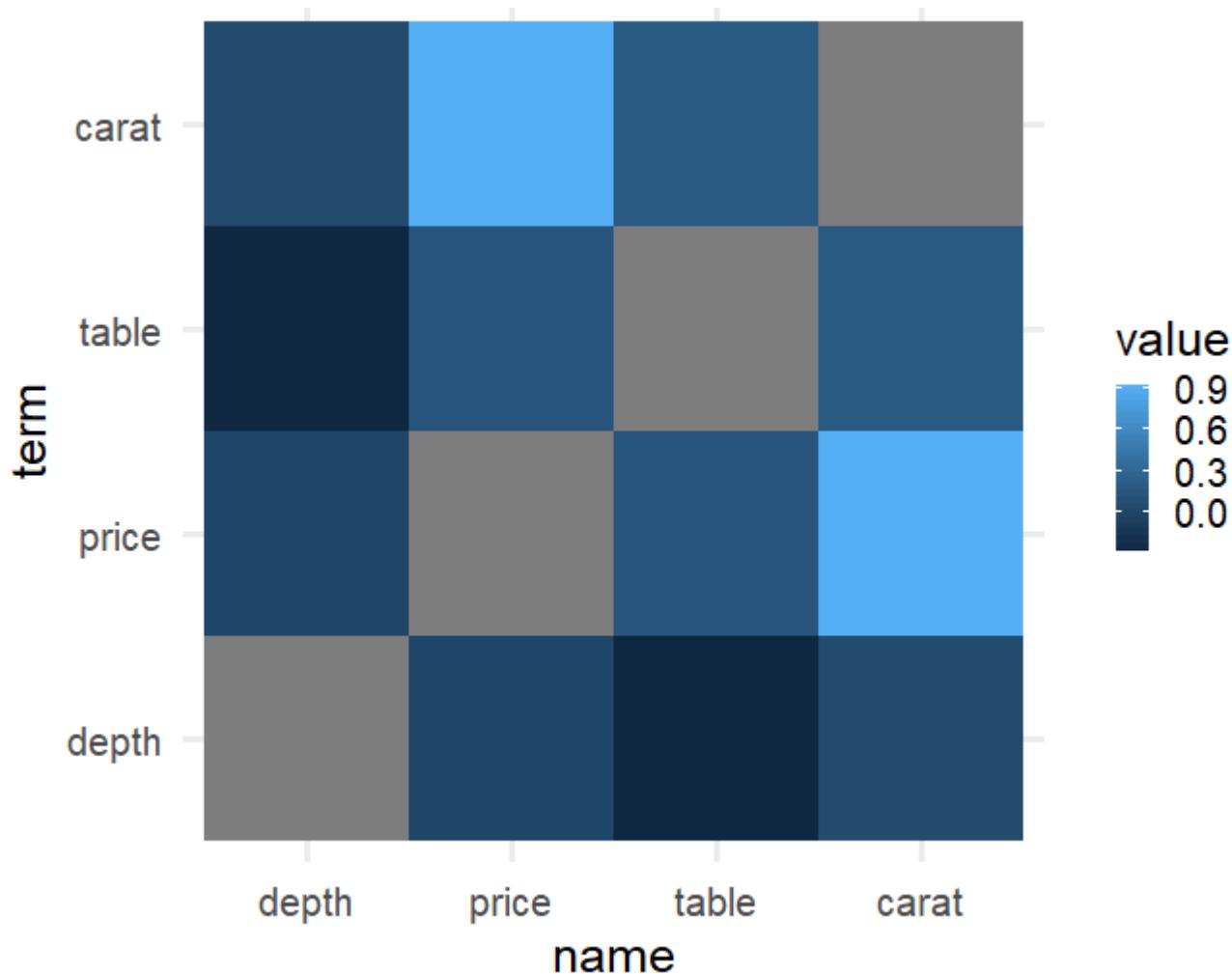
Heatmaps

First the data:

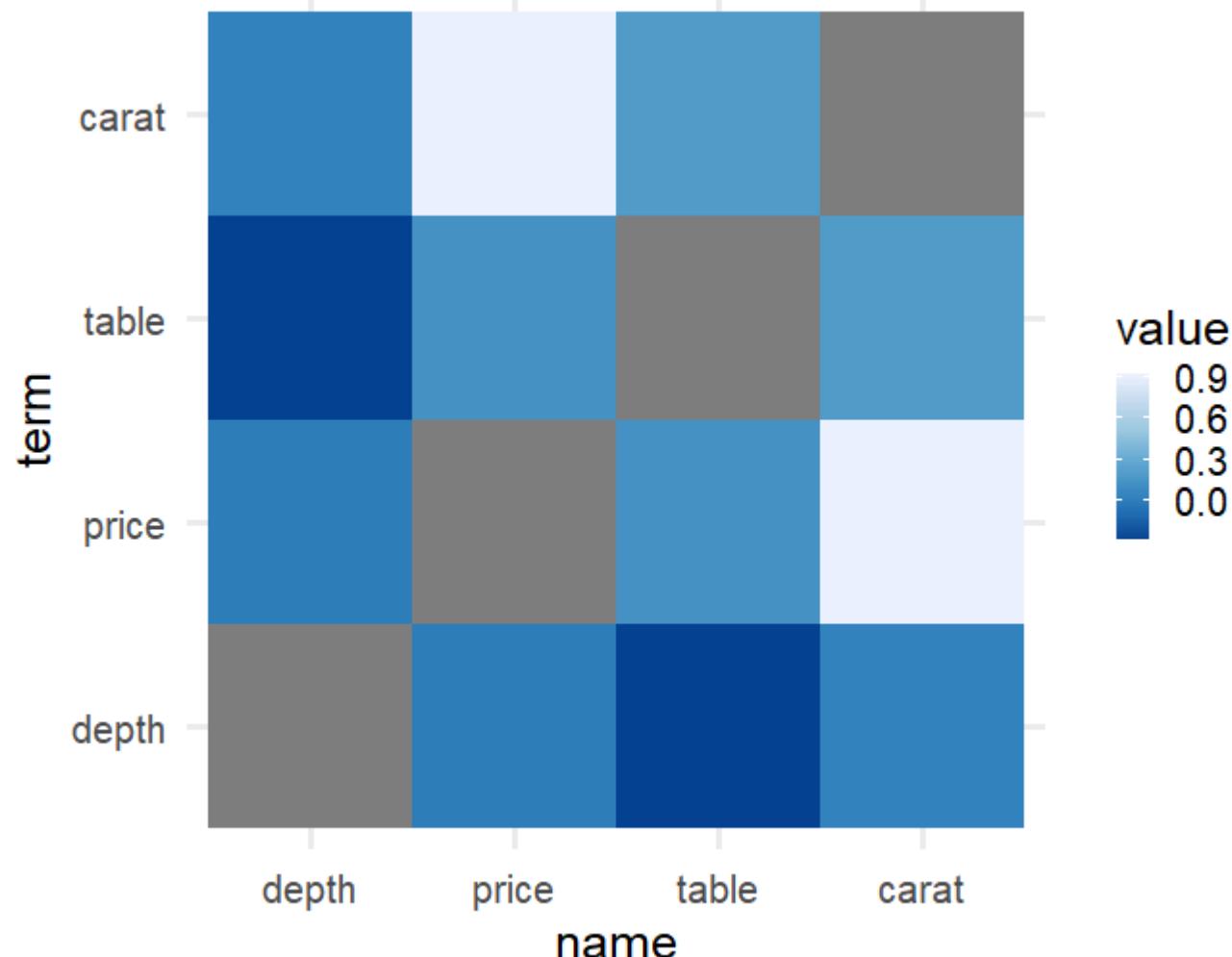
```
hm <- diamonds %>%
  select(table, price, depth, carat) %>%
  corrr::correlate() %>%
  pivot_longer(-term) %>%
  mutate(name = fct_reorder(name, value),
        term = fct_reorder(term, value))
hm

## # A tibble: 16 × 3
##   term    name   value
##   <fct> <fct>   <dbl>
## 1 table  table  NA
## 2 table  price   0.127
## 3 table  depth  -0.296
## 4 table  carat   0.182
## 5 price   table  0.127
## 6 price   price  NA
## # ... with 10 more rows
```

```
ggplot(hm, aes(name, term)) +  
  geom_tile(aes(fill = value)) +  
  coord_fixed()
```



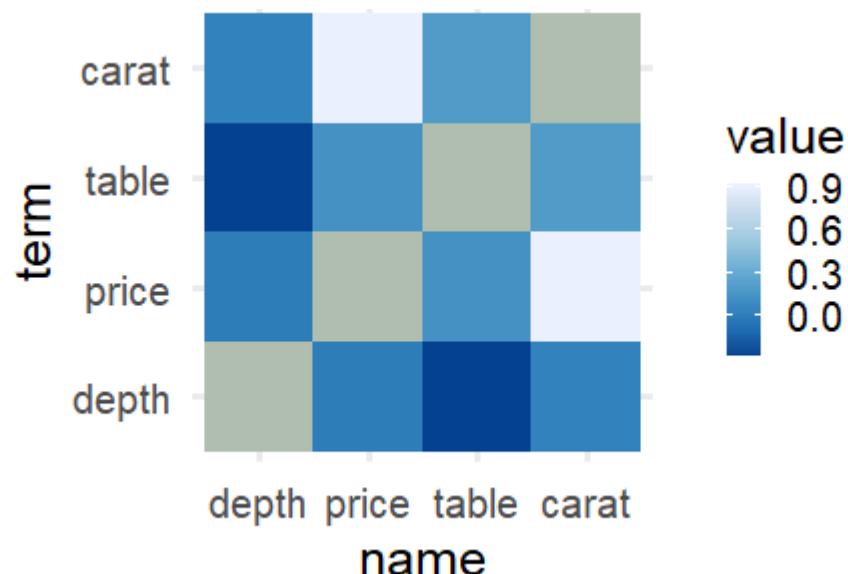
```
ggplot(hm, aes(name, term)) +  
  geom_tile(aes(fill = value)) +  
  coord_fixed() +  
  scale_fill_distiller(palette = "Blues")
```



Change the NA value

In any `scale_*` you can change the NA value, including to "transparent".

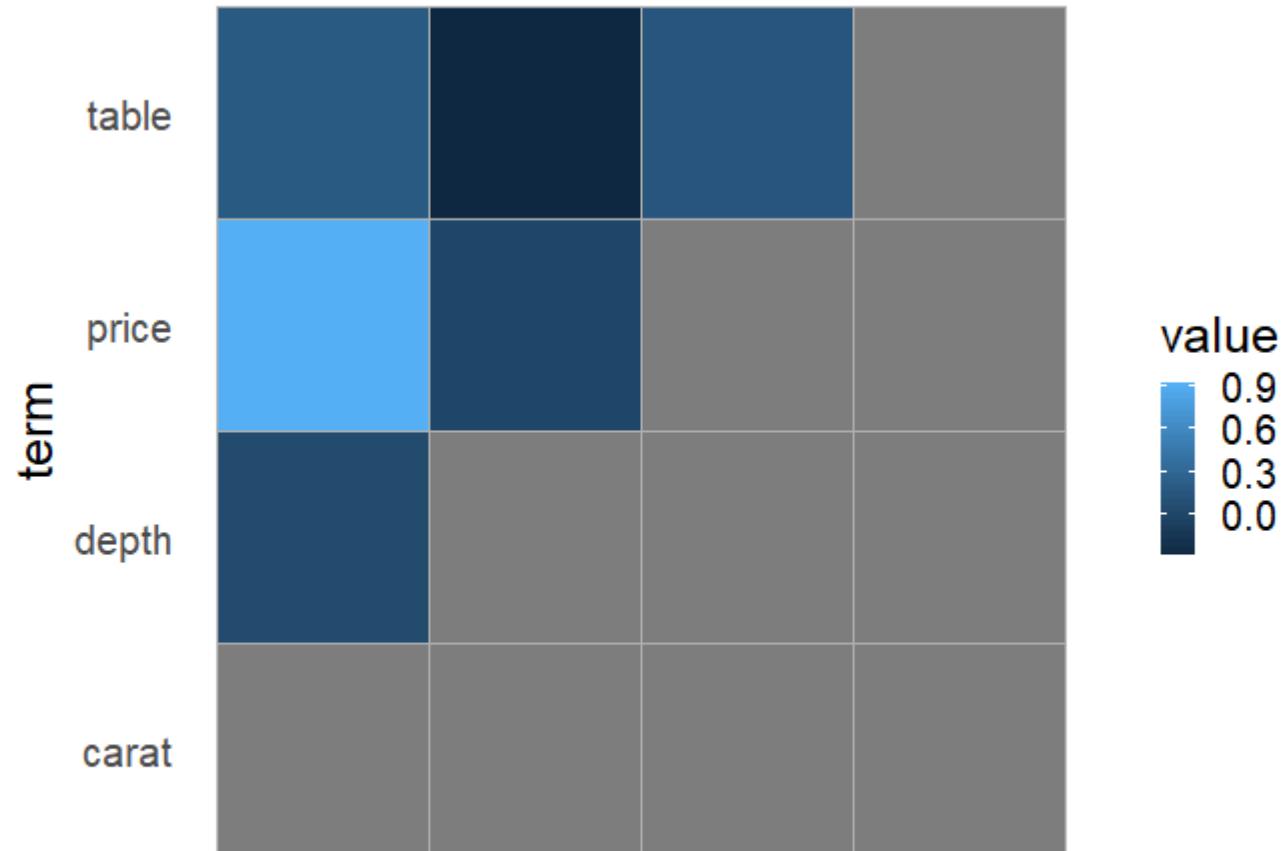
```
ggplot(hm, aes(name, term)) +  
  geom_tile(aes(fill = value)) +  
  coord_fixed() +  
  scale_fill_distiller(palette = "Blues",  
                      na.value = "#b0bfb0")
```



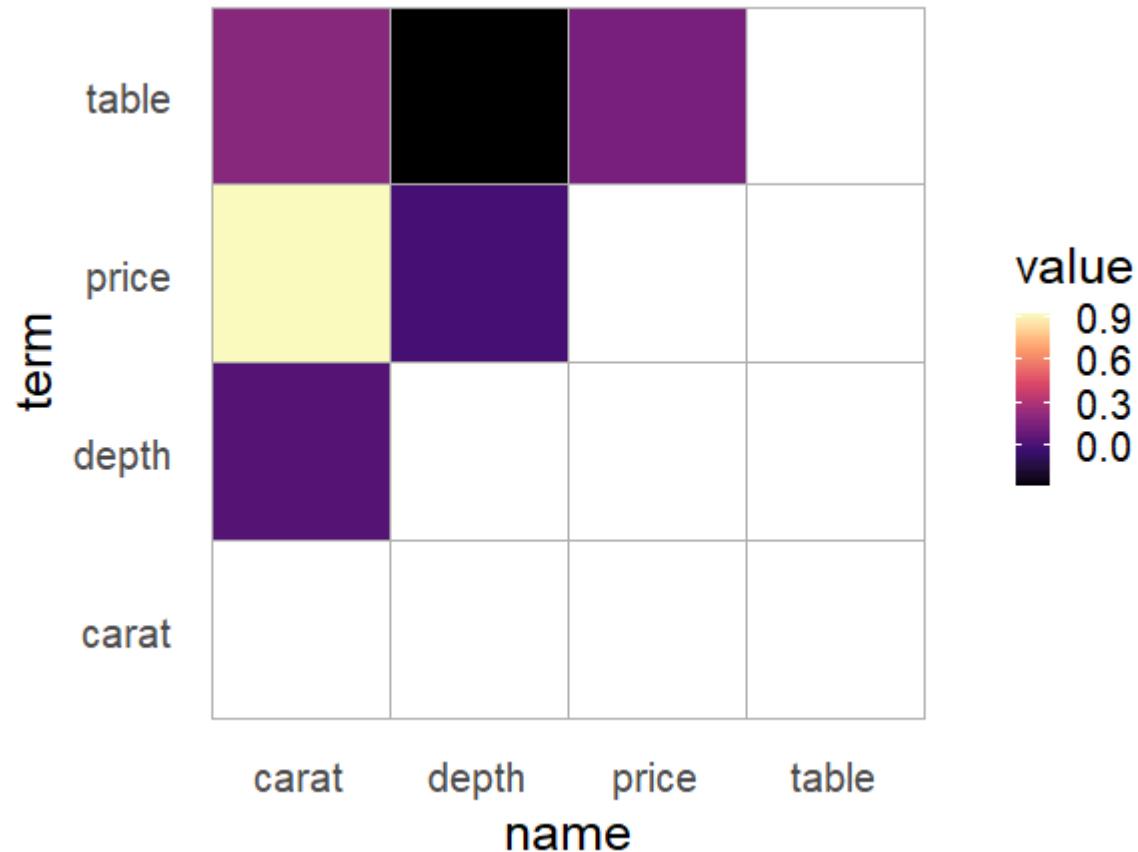
Use this to our advantage

```
hm2 <- diamonds %>%
  select(table, price, depth, carat) %>%
  corrr::correlate() %>%
  corrr::shave(upper = FALSE) %>%
  pivot_longer(-term)
```

```
hm2_default <- ggplot(hm2, aes(name, term)) +  
  geom_tile(aes(fill = value), color = "gray70") +  
  coord_fixed() +  
  theme(panel.grid.major = element_blank(),  
        panel.grid.minor = element_blank())  
hm2_default
```



```
hm2_default +  
  scale_fill_viridis_c(  
    option = "magma",  
    na.value = "transparent"  
)
```



Choropleths

```
library(tidycensus)
library(tigris)
options(tigris_use_cache = TRUE)
options(tigris_class="sf")

#census_api_key("Insert your key", install=TRUE)

vars_pl_2020 <- load_variables(2020, "p1")
#View(vars)
```

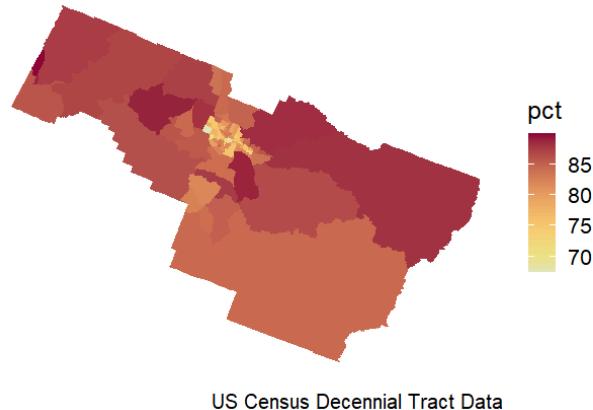
Percentage of people identifying as White
Lane County



Heat palette

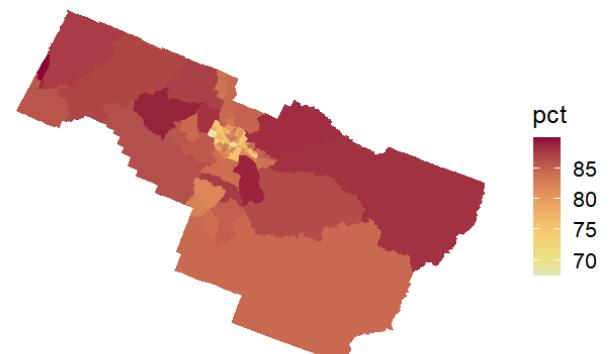
```
lane %>%
  mutate(pct = 100 * (value / summary_value)) %
  ggplot(aes(fill = pct, color = pct)) +
  geom_sf() +
  coord_sf(crs = 26915) +
  theme_dviz_map(font_size = 25, font_family =
  scale_fill_continuous_sequential("Heat") +
  scale_color_continuous_sequential("Heat") +
  labs(title = "Percentage of people identifying as White",
       subtitle = "Lane County",
       caption = "US Census Decennial Tract Data")
```

Percentage of people identifying as White
Lane County



Percentage of people identifying as White

Lane County



US Census Decennial Tract Data

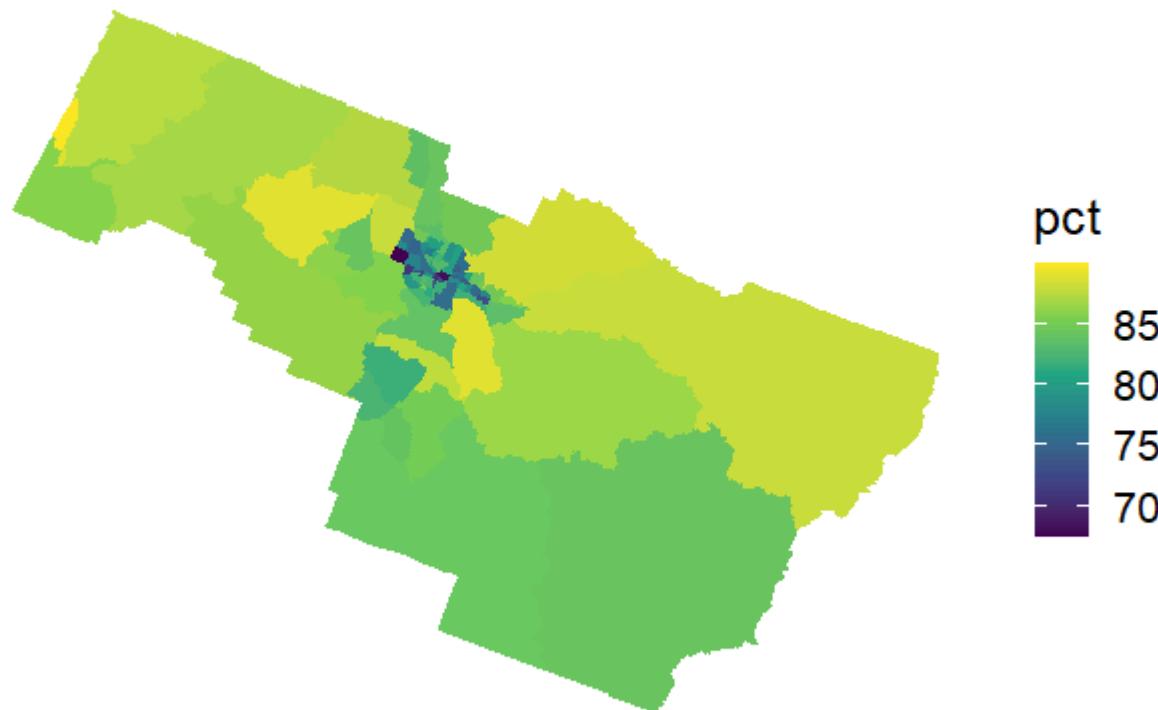
Options

- `scale_fill_continuous_sequential("Heat")`
- `scale_color_continuous_sequential("Heat")`
- `scale_fill_discrete_sequential("Heat")`
- `scale_color_discrete_sequential("Heat")`

viridis palette

Percentage of people identifying as White

Lane County



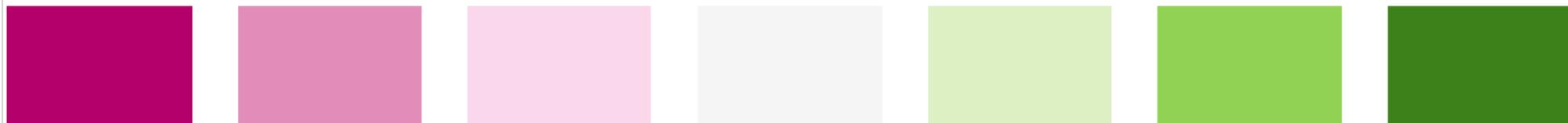
US Census Decennial Tract Data

Diverging palettes

CARTO Earth



ColorBrewer PiYG



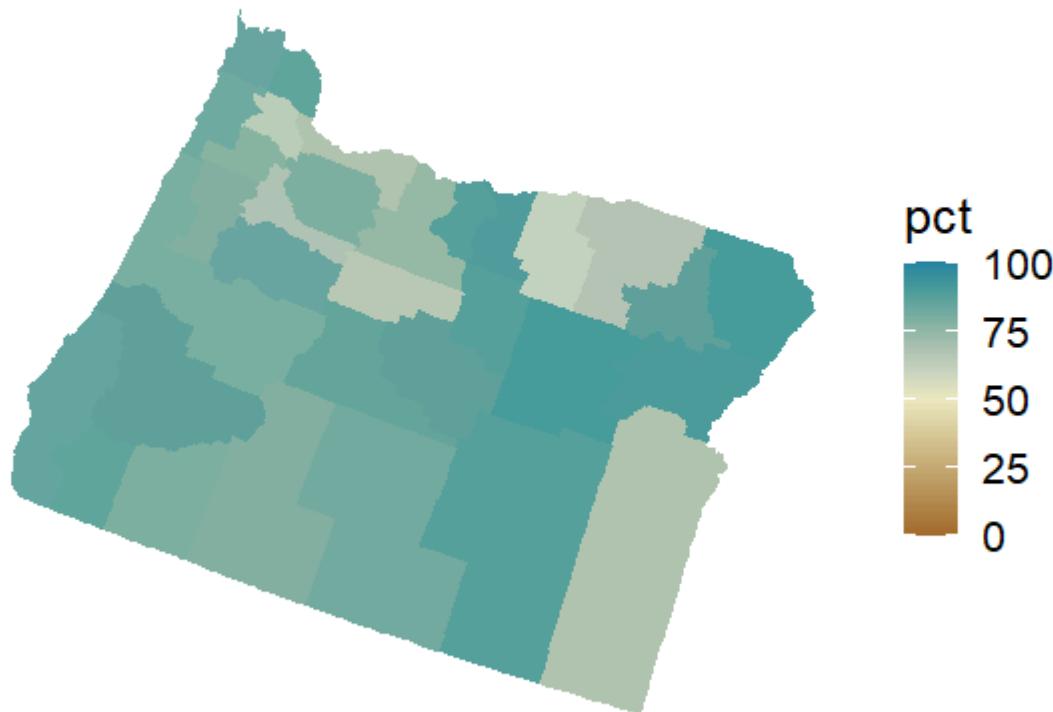
Blue-Red



Earth palette

Percentage of people identifying as White

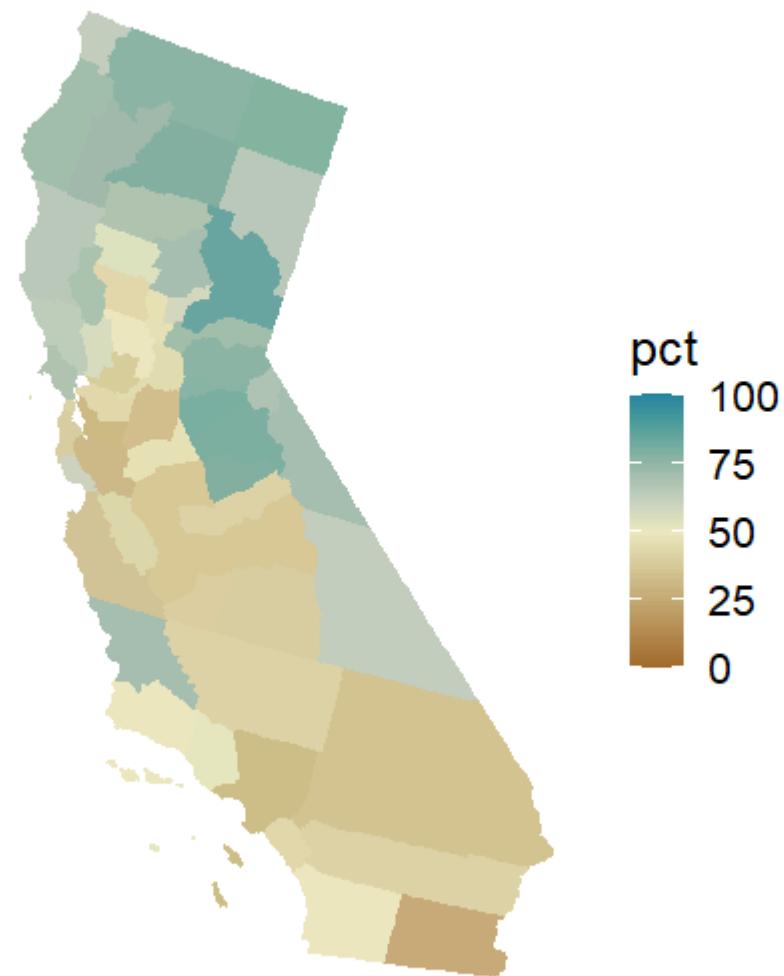
Oregon



US Census Decennial Tract Data

Percentage of people identifying as White

California

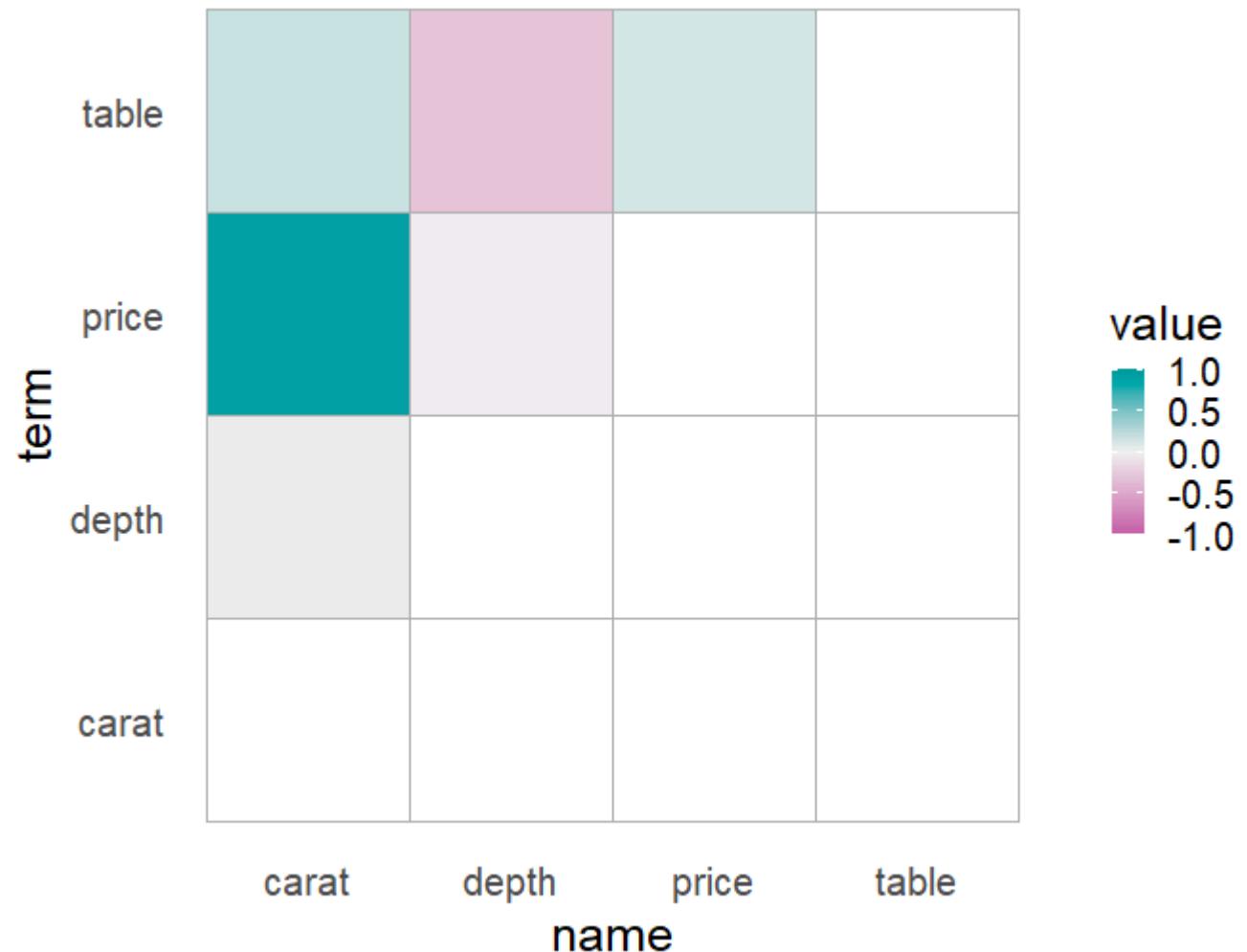


Back to our heatmap

Use a diverging palette that balances at zero.

Notice the transparency is now a bit problematic...

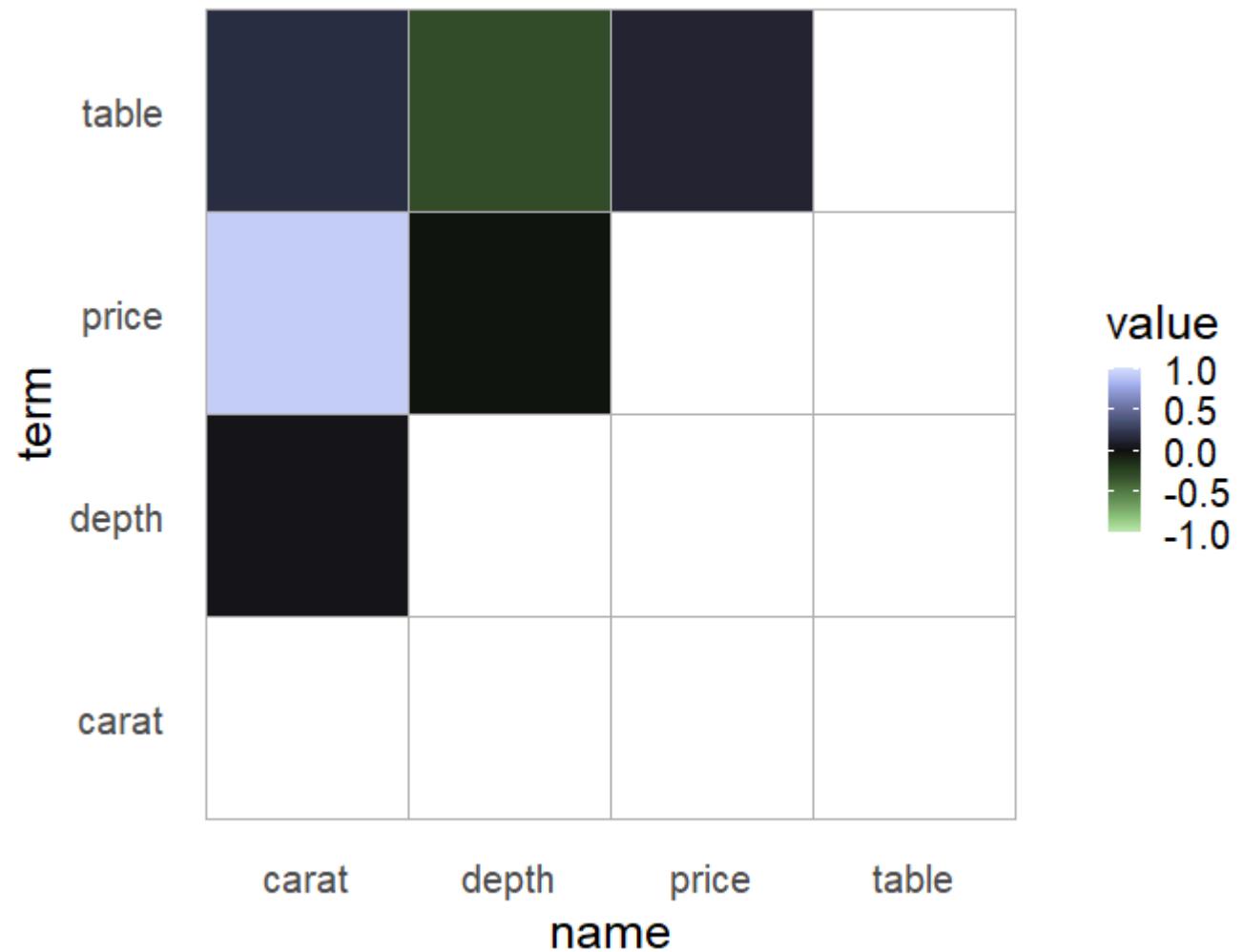
```
library(colorspace)
hm2_default +
  scale_fill_continuous_divergingx(
    palette = "Tropic",
    mid = 0,
    limits = c(-1, 1),
    rev = TRUE,
    na.value = "transparent"
)
```



Try a different palette

Most diverging palettes have a light gray or white center point. Some have black. Let's try one of those.

```
hm2_default +  
  scale_fill_continuous_diverging(  
    palette = "Tofino",  
    mid = 0,  
    limits = c(-1, 1),  
    rev = TRUE,  
    na.value = "transparent"  
)
```

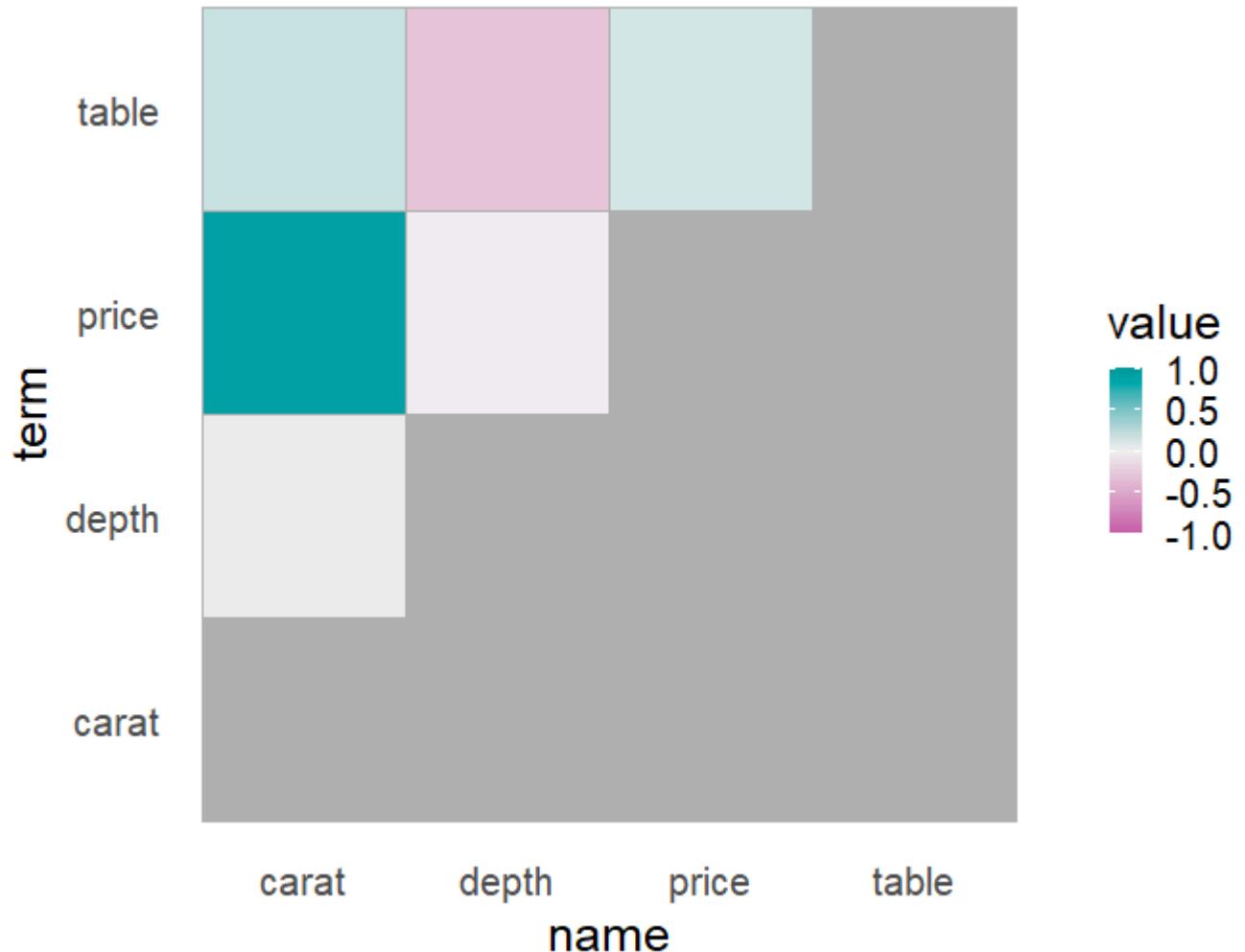


don't love it

One more try!

Back to our last palette, but change the NA value to a light gray

```
hm2_default +  
  scale_fill_continuous_divergingx(  
    palette = "Tropic",  
    mid = 0,  
    limits = c(-1, 1),  
    rev = TRUE,  
    na.value = "gray70"  
)
```



Maybe better? I give up for now...

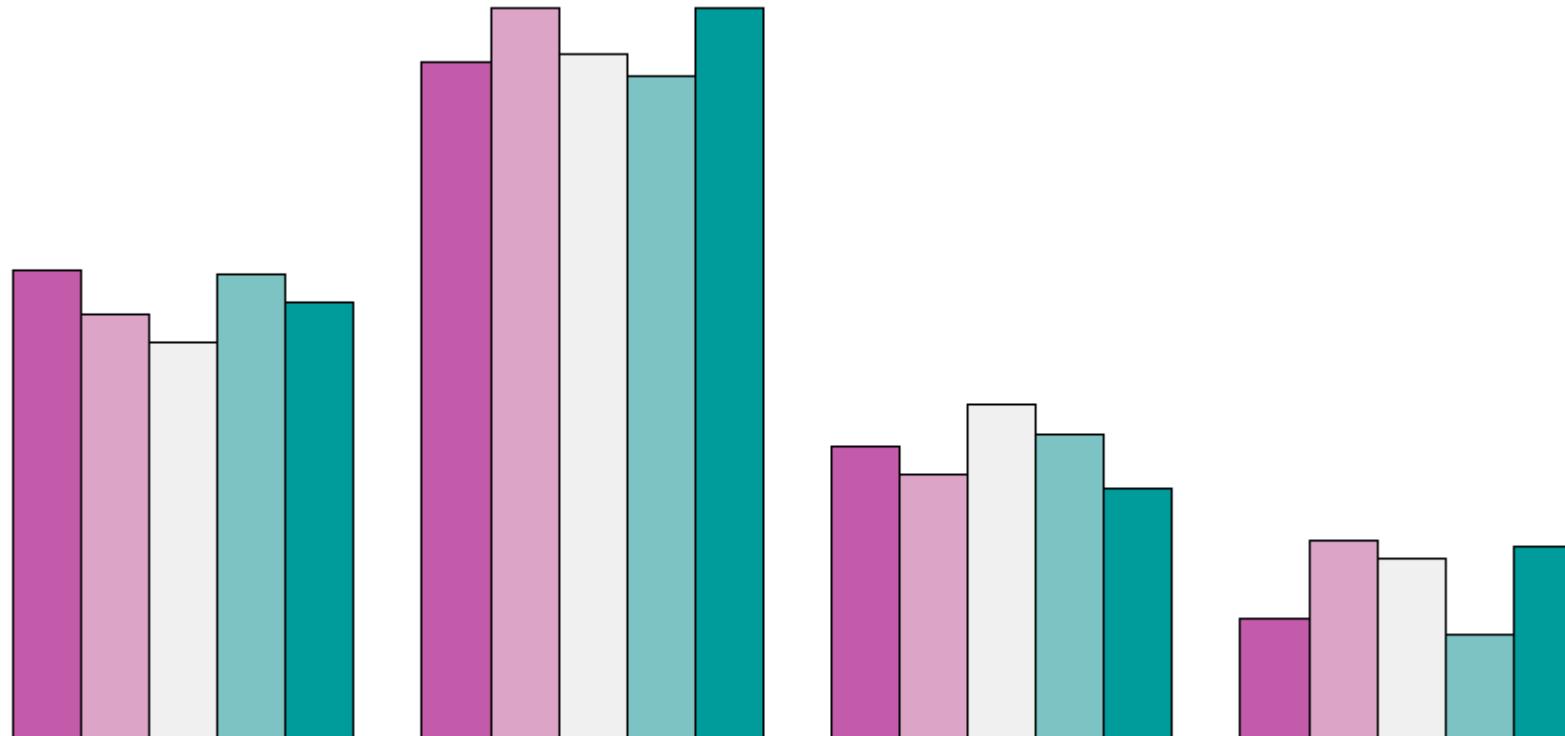
More on colorspace

Great package with great documentation. Let's go look!

colorspace

More diagnostics

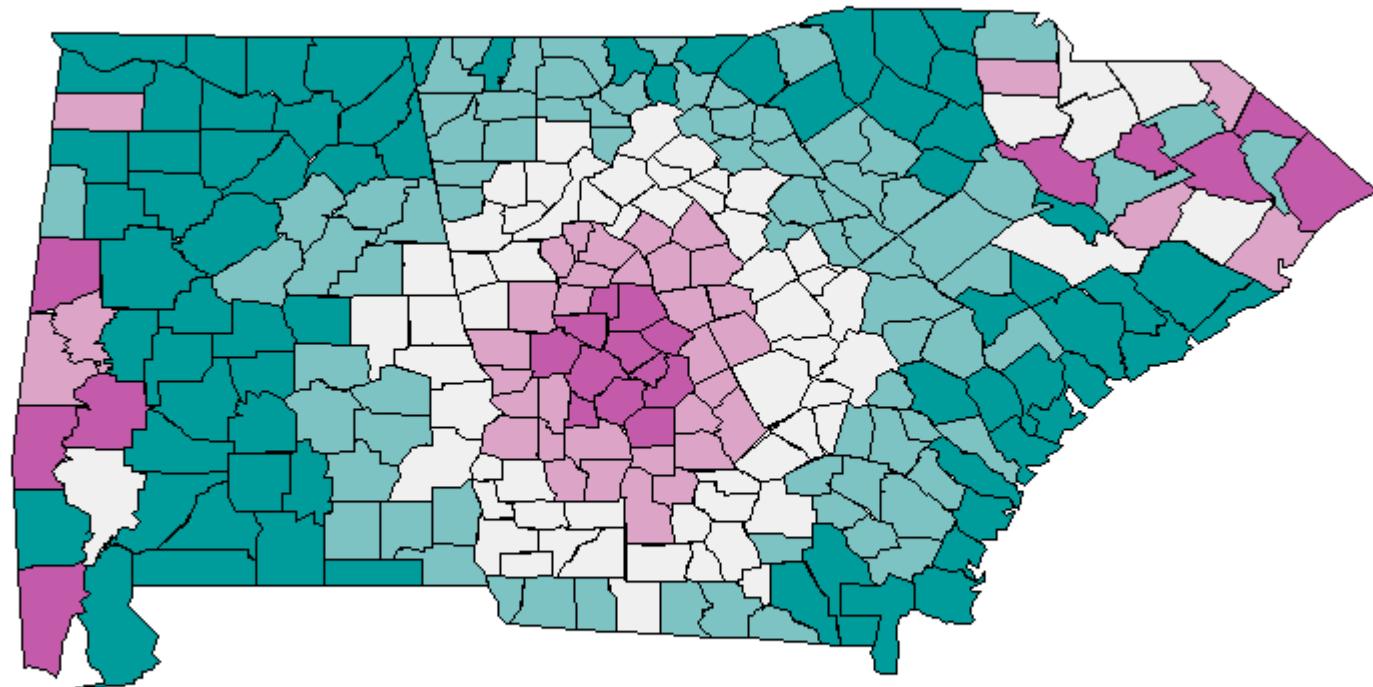
```
our_scale <- diverge_hcl(5, palette = "Tropic", rev = TRUE)
demoplot(our_scale, "bar")
```



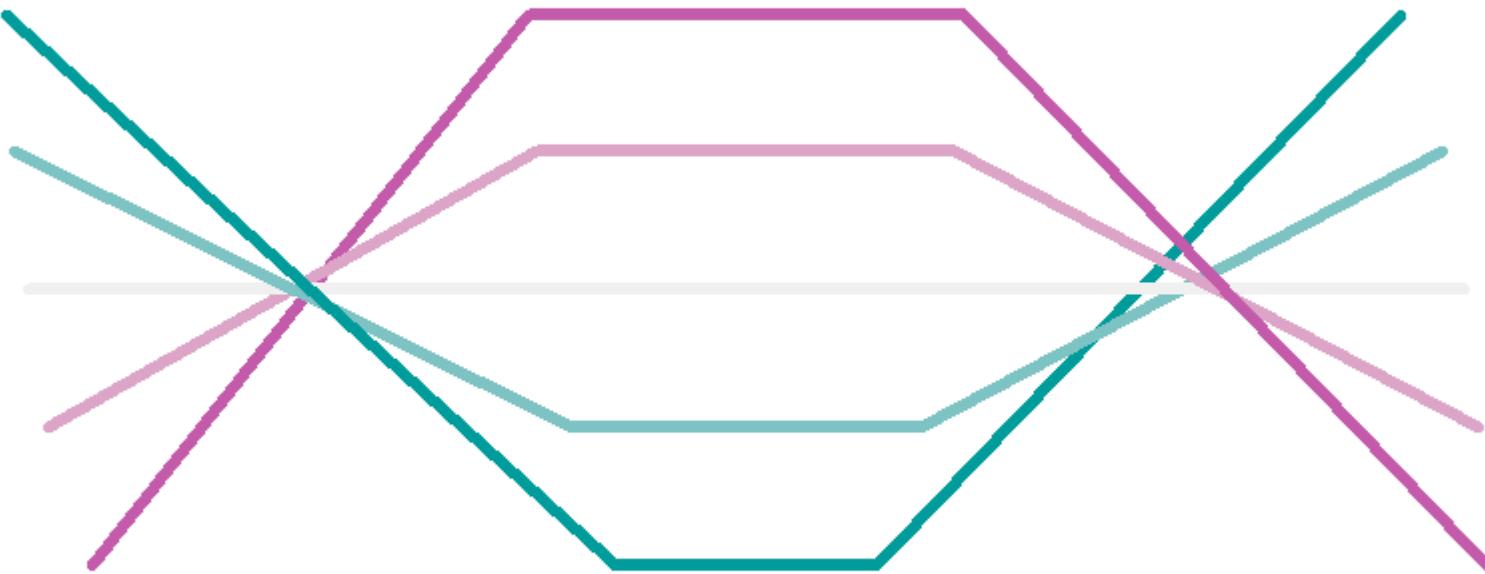
```
demoplot(our_scale, "heatmap")
```



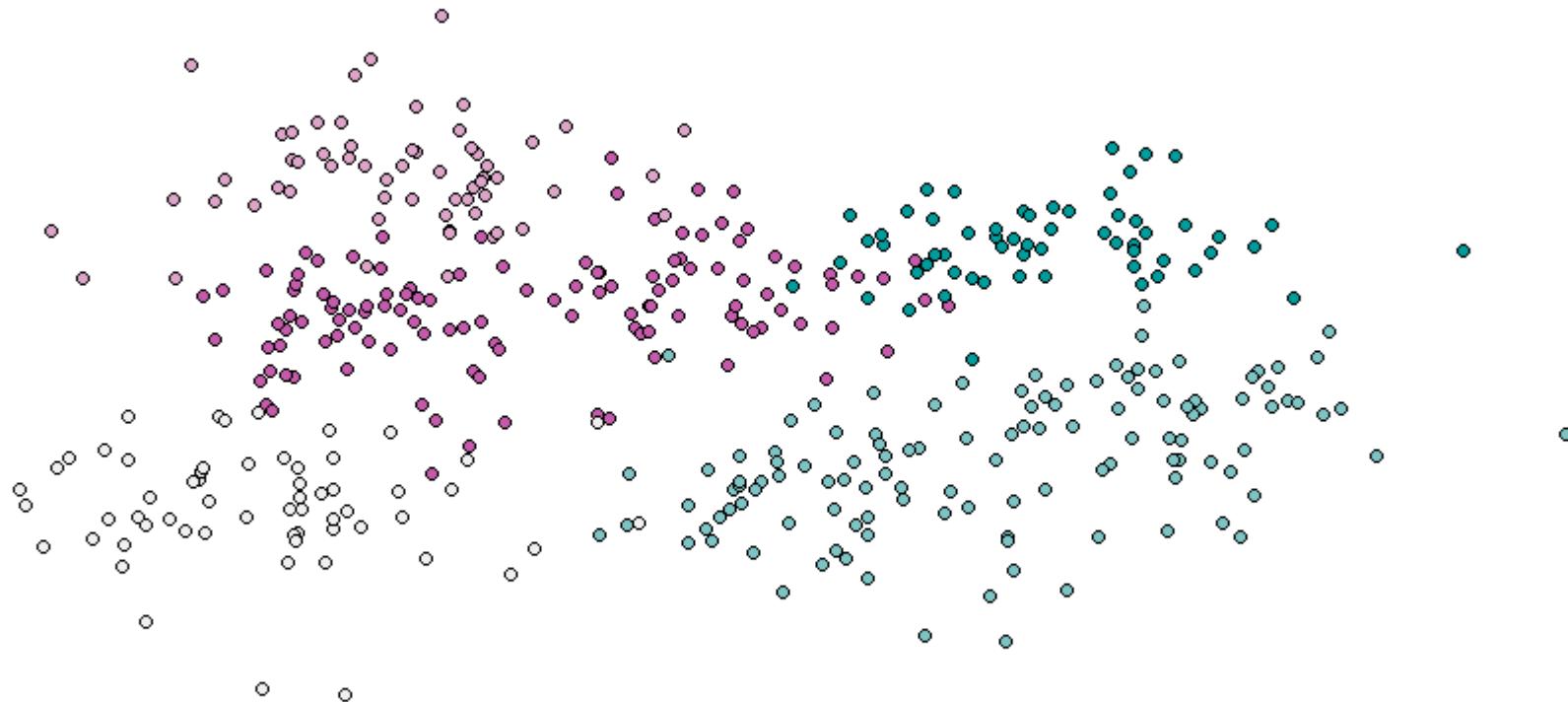
```
demoplot(our_scale, "map")
```



```
demoplot(our_scale, "lines")
```



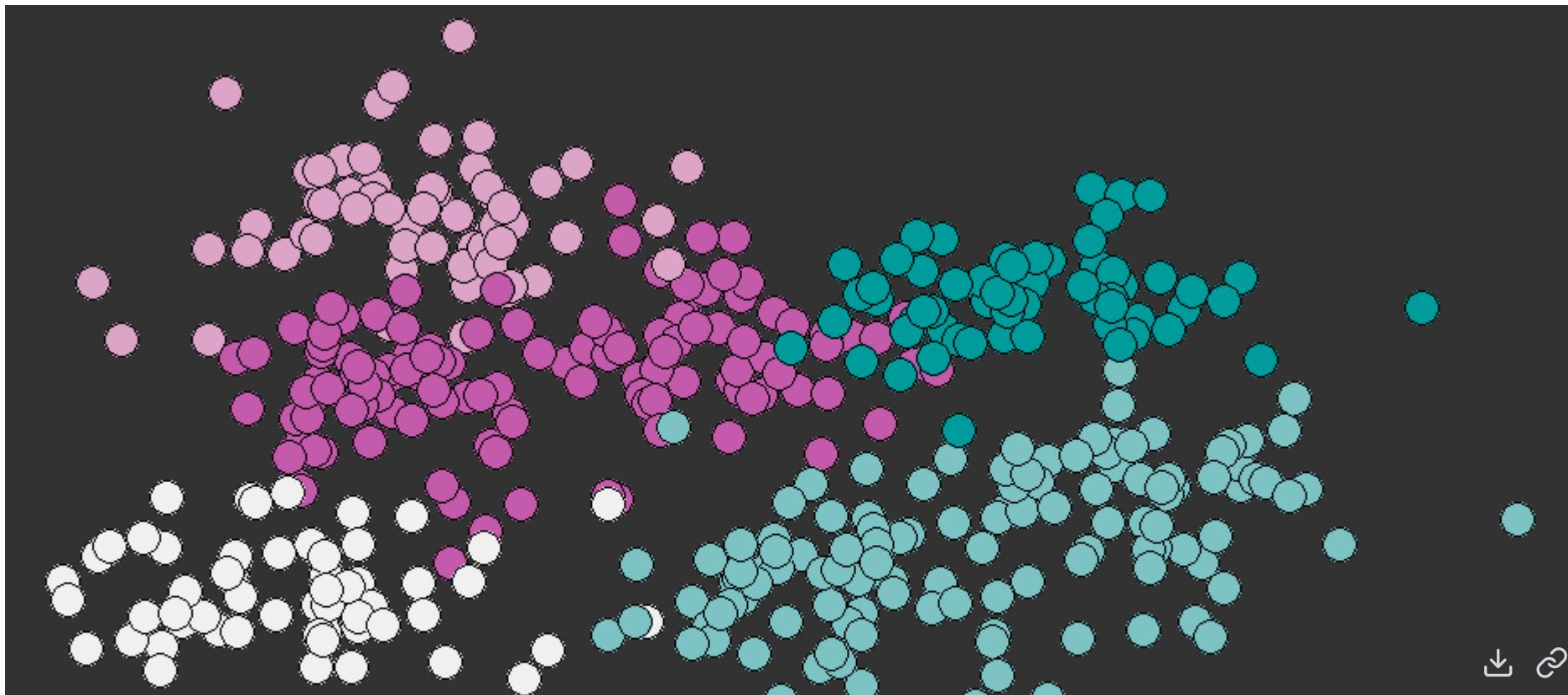
```
demoplot(our_scale, "scatter")
```



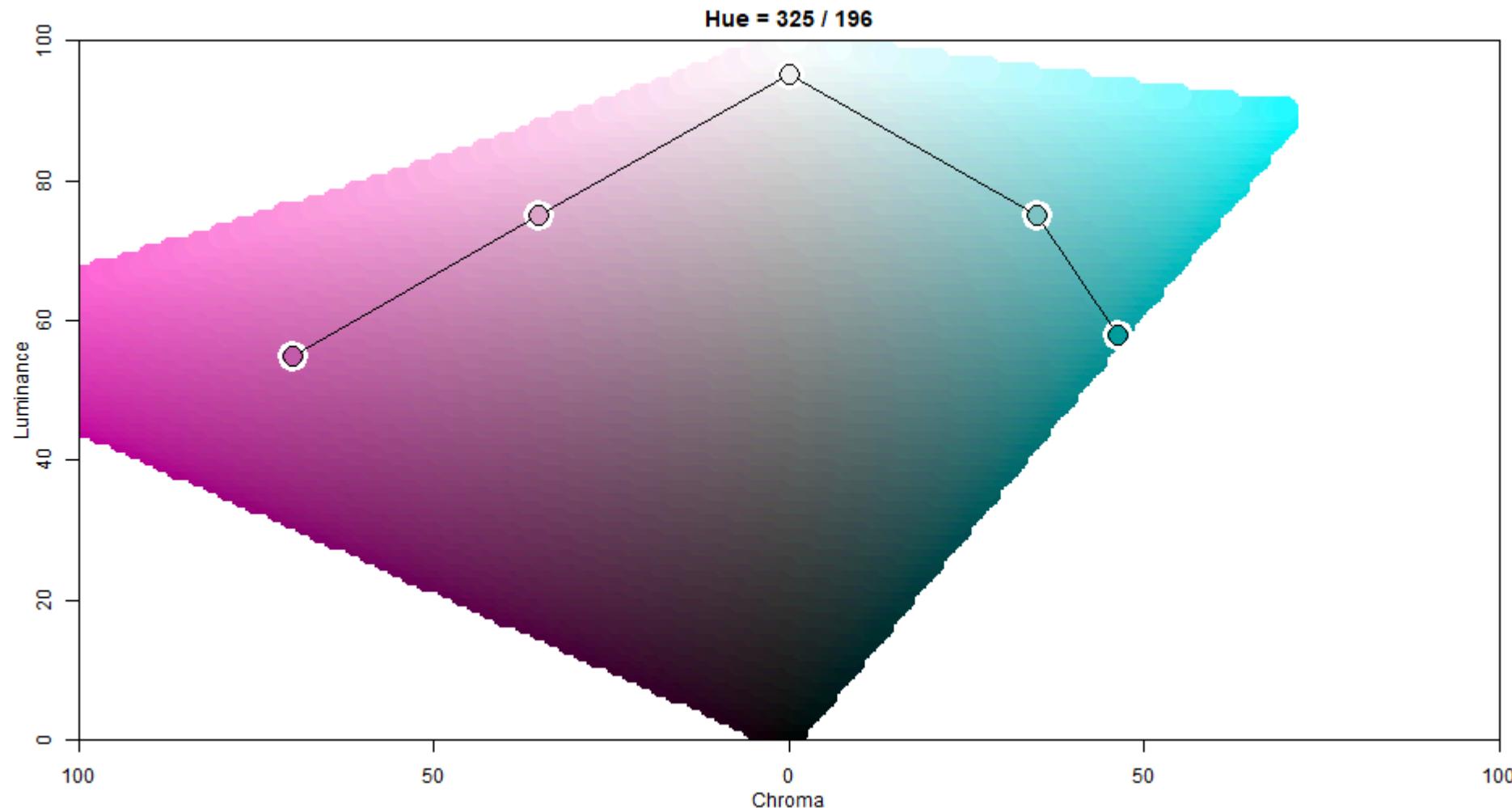
Diferent background

Note - this is base graphics

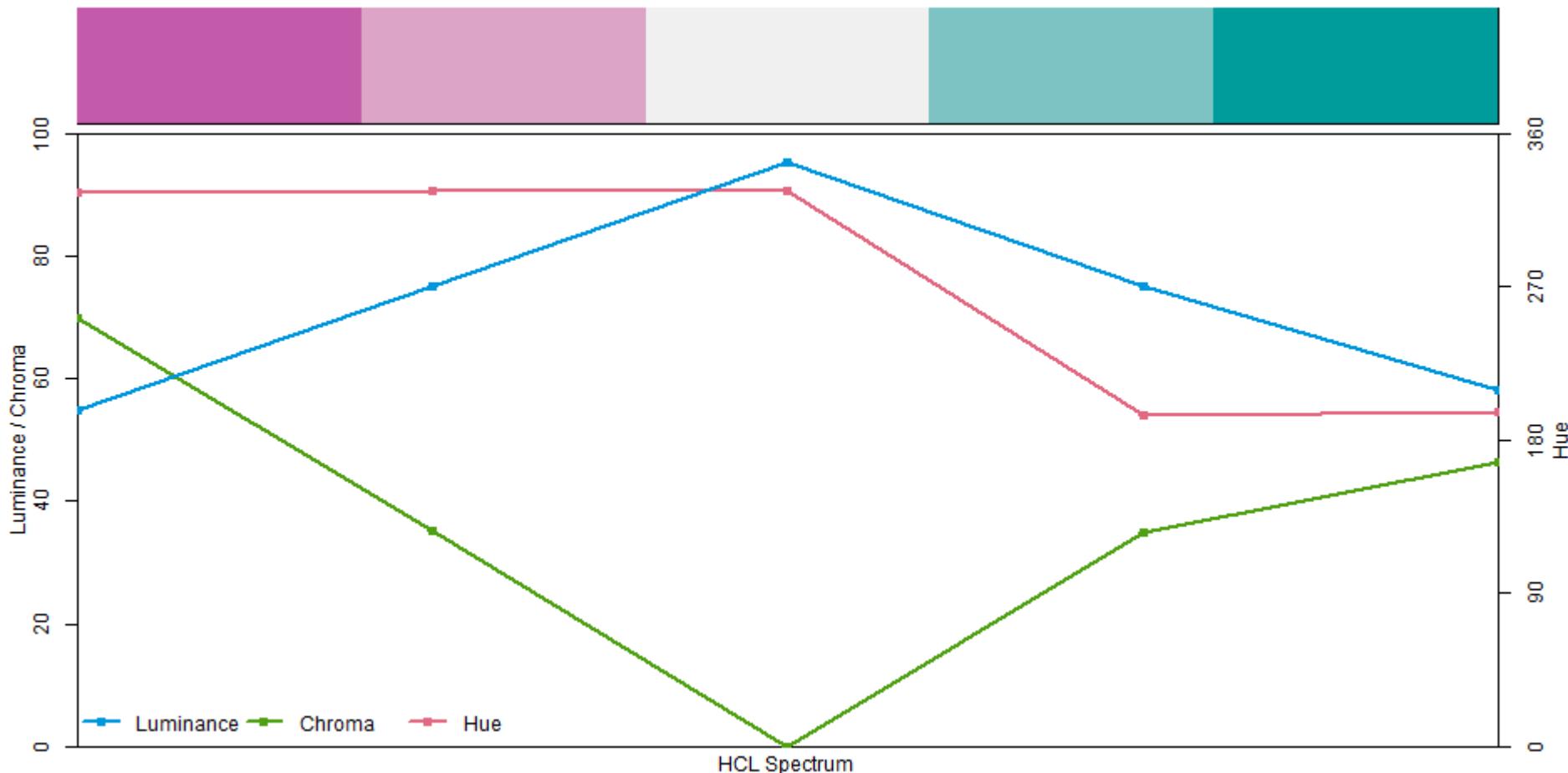
```
par(bg = "gray20", cex = 3, mar = rep(0, 4))
demoplot(our_scale, "scatter")
```



```
hclplot(our_scale)
```



```
specplot(our_scale, type = "o")
```

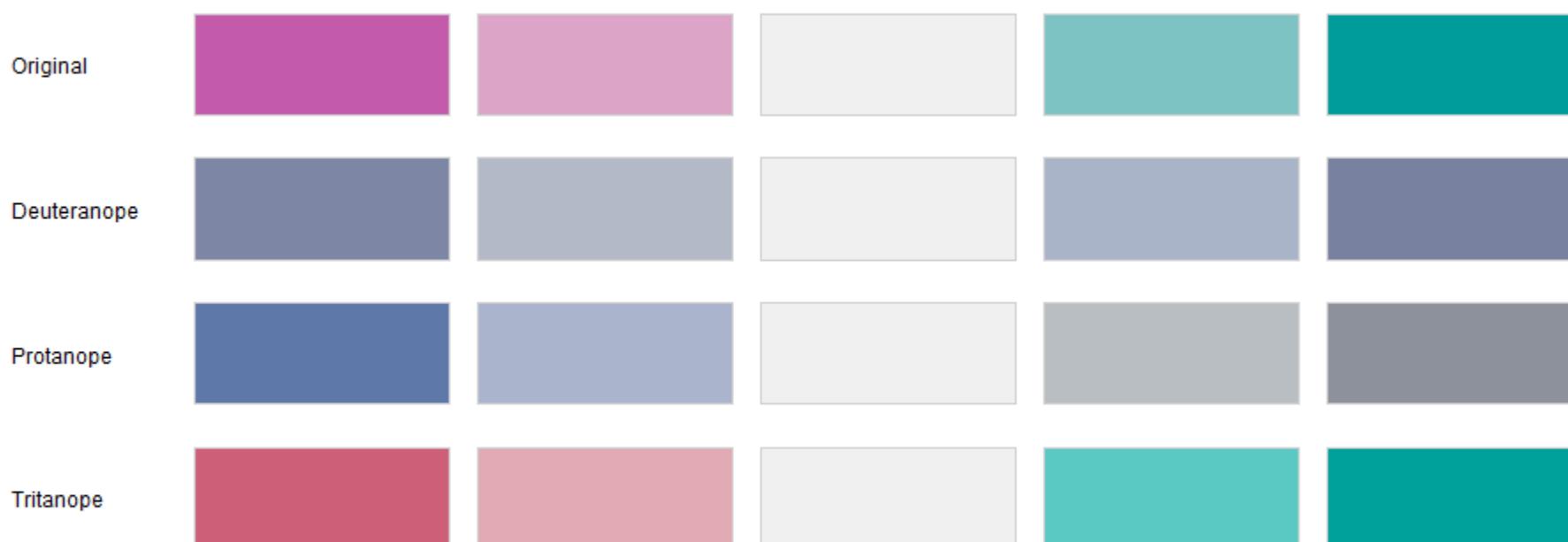


Check colorblindness

We can check for our palette, in addition to our figure overall

```
swatchplot("Our palette" = our_scale, cvd = TRUE)
```

Our palette



Interactivity

You can do this all online interactively with the demo plots

<http://hclwizard.org:3000/hclwizard/>

CVD Emulator online

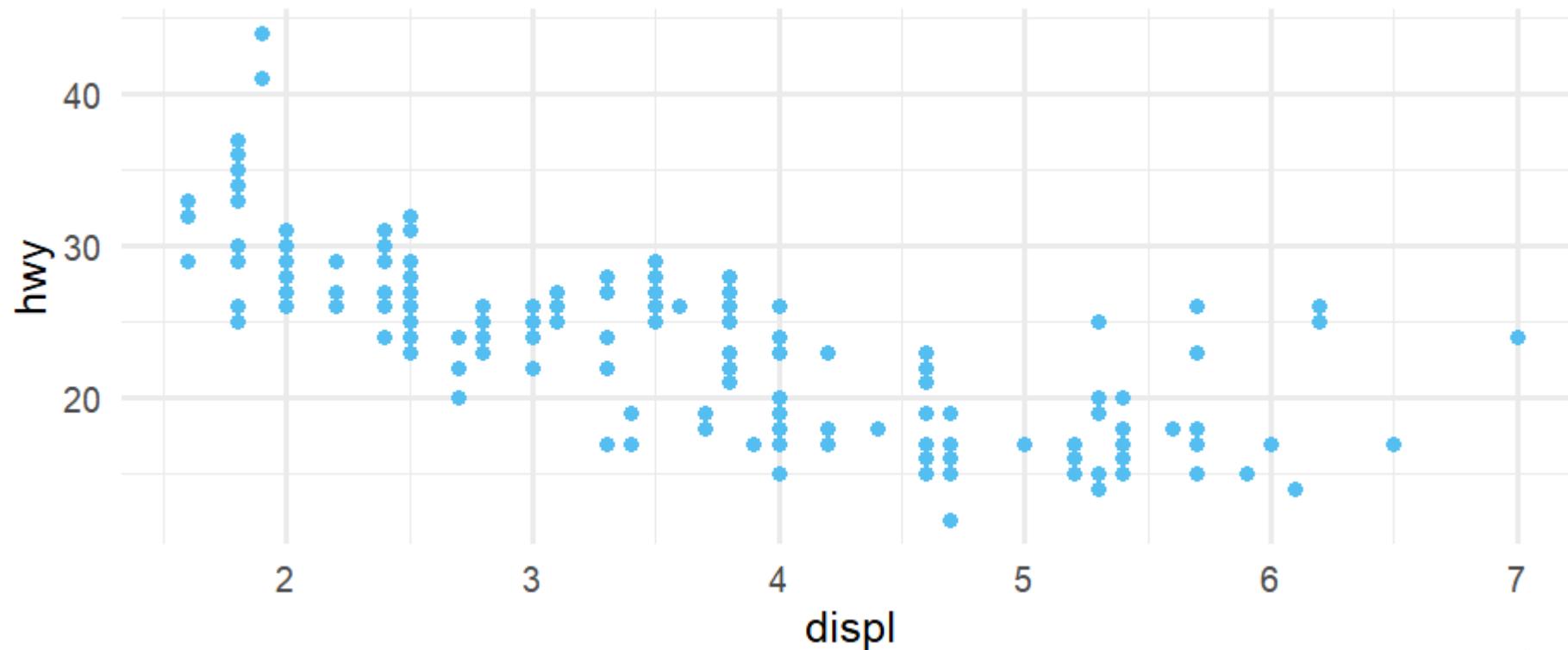
<http://hclwizard.org:3000/cvdemulator/>

Color as a tool to highlight

MPG data

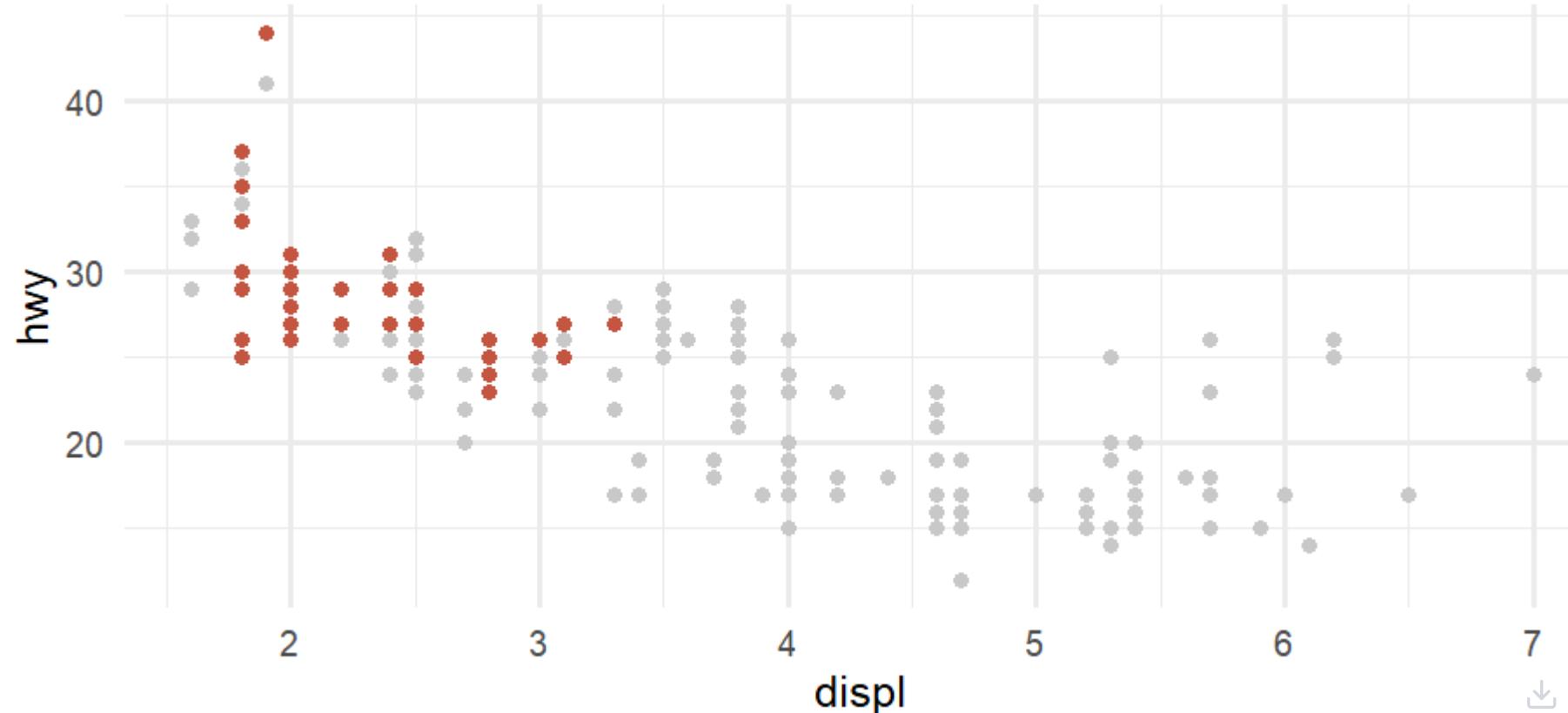
Basic scatterplot of weight to highway mpg

```
ggplot(mpg, aes(displ, hwy)) +  
  geom_point()
```



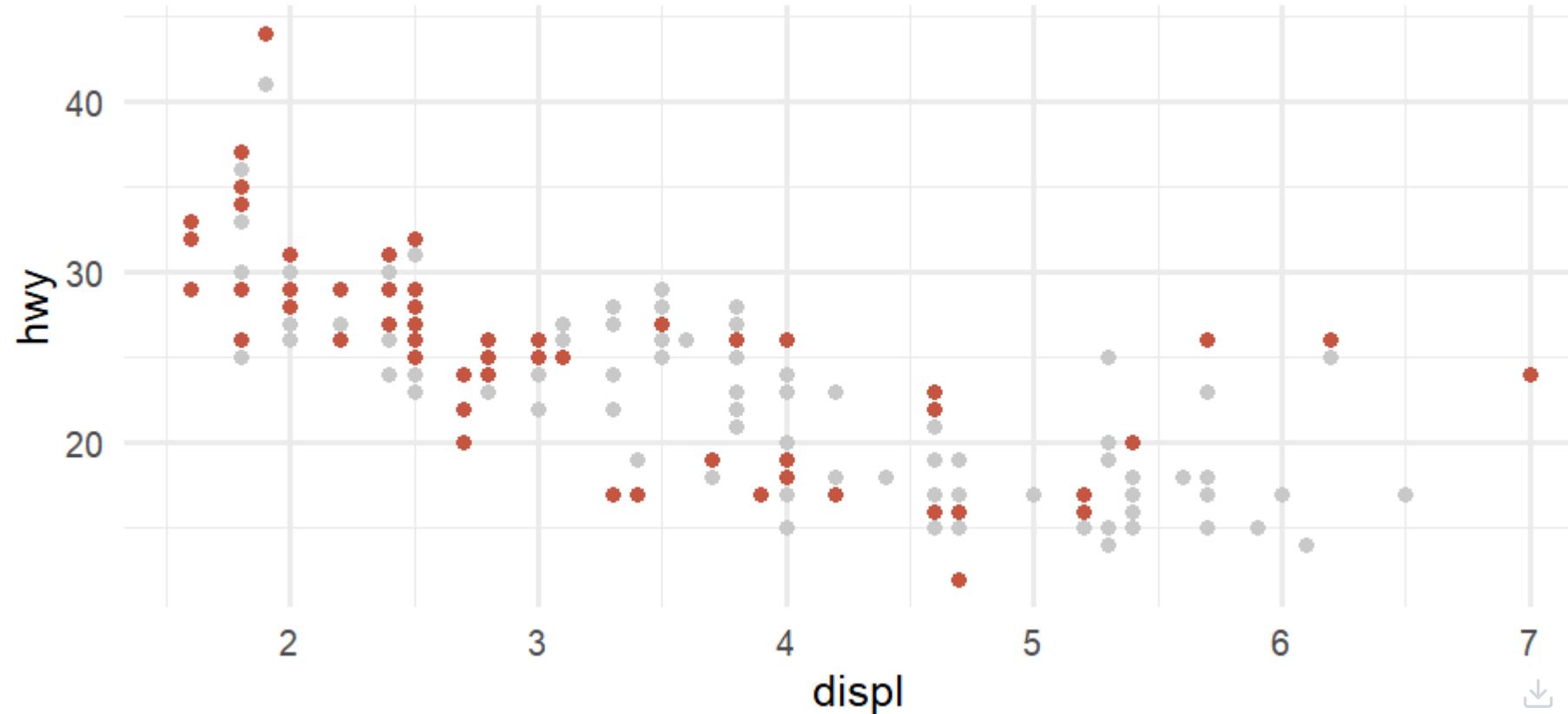
Highlight compact cars

```
ggplot(mpg, aes(displ, hwy)) +  
  geom_point(color = "gray80") +  
  geom_point(data = filter(mpg, class == "compact"),  
             color = "#C55644")
```



Highlight manual cars

```
ggplot(mpg, aes(displ, hwy)) +  
  geom_point(color = "gray80") +  
  geom_point(data = filter(mpg, str_detect(trans, "manual")),  
             color = "#C55644")
```

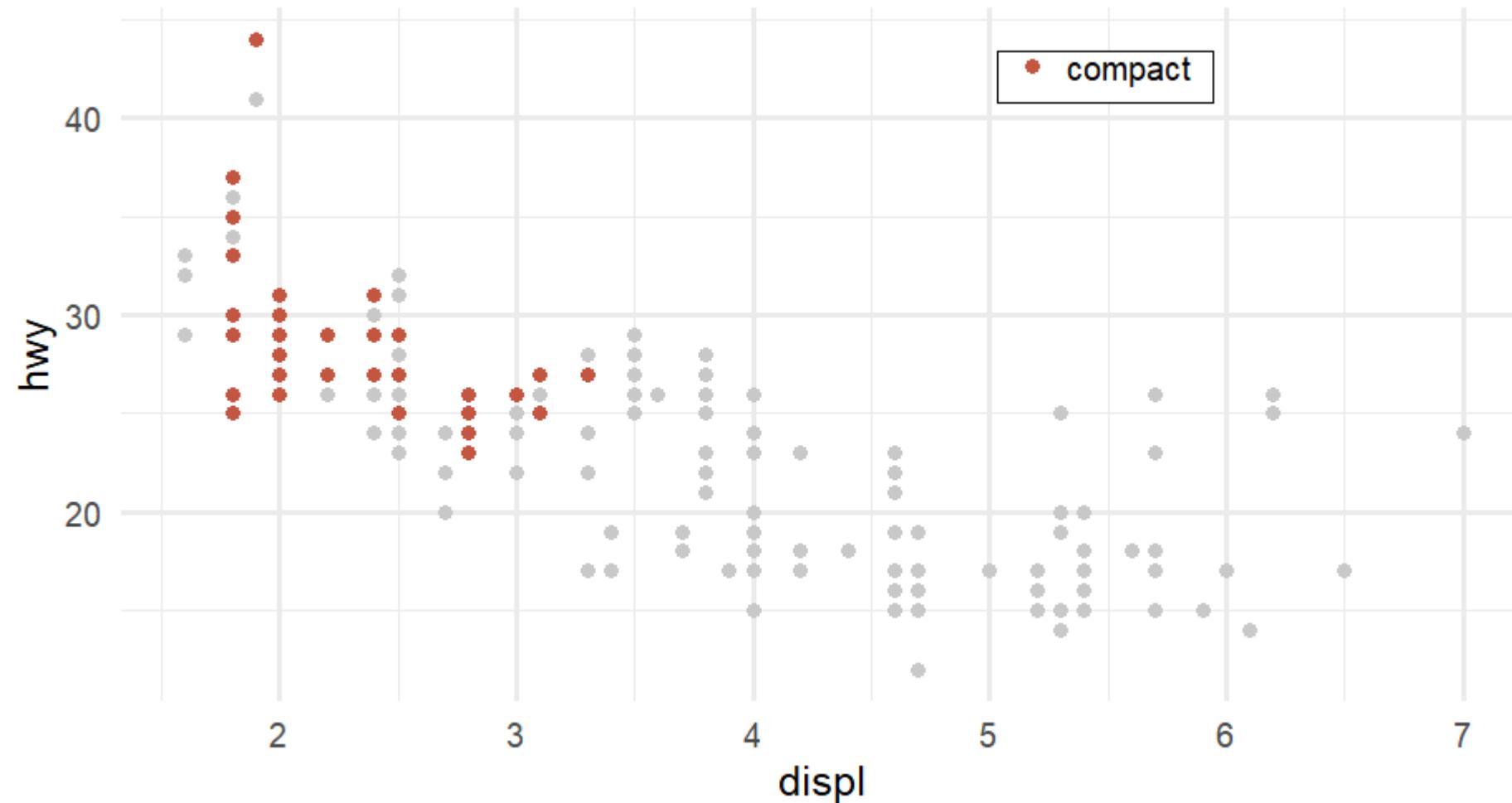


Add a legend

Couple of different ways - mostly hacky

- Pretend you have a column called "compact"

```
ggplot(mpg, aes(displ, hwy)) +  
  geom_point(color = "gray80") +  
  geom_point(data = filter(mpg, class == "compact"),  
             aes(color = "compact")) +  
  scale_color_manual(values = "#C55644") +  
  theme(  
    legend.title = element_blank(),  
    legend.position = c(0.7, 0.9),  
    legend.box.background = element_rect(colour = "black"),  
    legend.box.margin = margin(t = -0.5, unit = "cm"))  
)
```

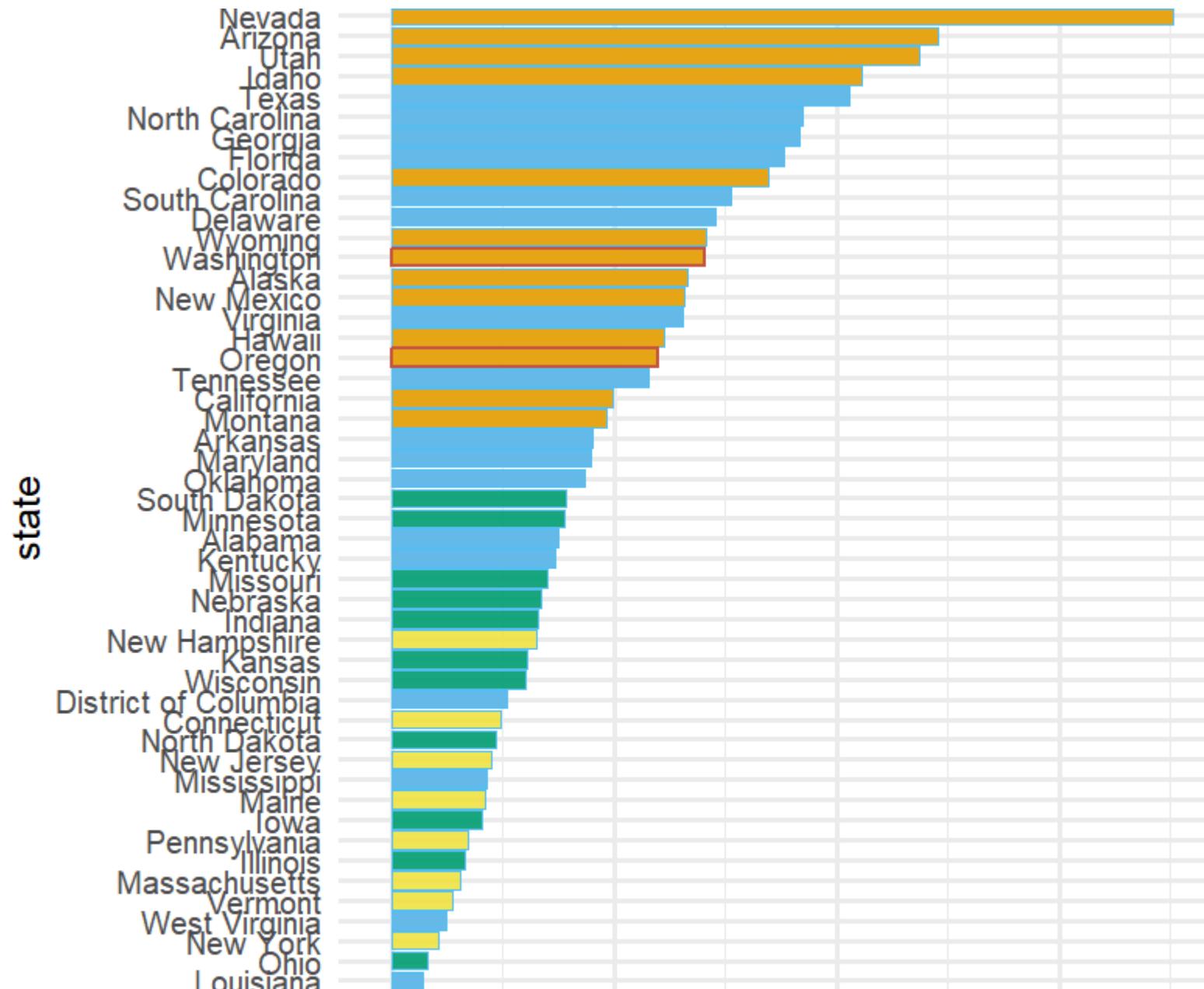


Back to our states plot

Highlight Oregon and Washington

```
#popgrowth_df <- popgrowth_df %>%
  # arrange(desc(popgrowth)) %>%
  # mutate(state = factor(state, levels = state))

ggplot(popprowth_df, aes(x=popgrowth, y=state)) +
  geom_col(aes(fill = region), alpha = 0.9) +
  geom_col(data=filter(popprowth_df, state == 'Oregon' | state == 'Washington'),
            fill = NA,
            color = '#C55644',
            linewidth = 1,
            width = 0.9) + # Added width parameter to make it slightly narrower
  scale_fill_OkabeIto()
```



region

- West
- South
- Midwest
- Northeast

Color labels

```
states <- unique(popgrowth_df$state)

label_color <- ifelse(states == "Oregon" | states == "Washington",
                      "#C55644",
                      "gray30")

label_color
```

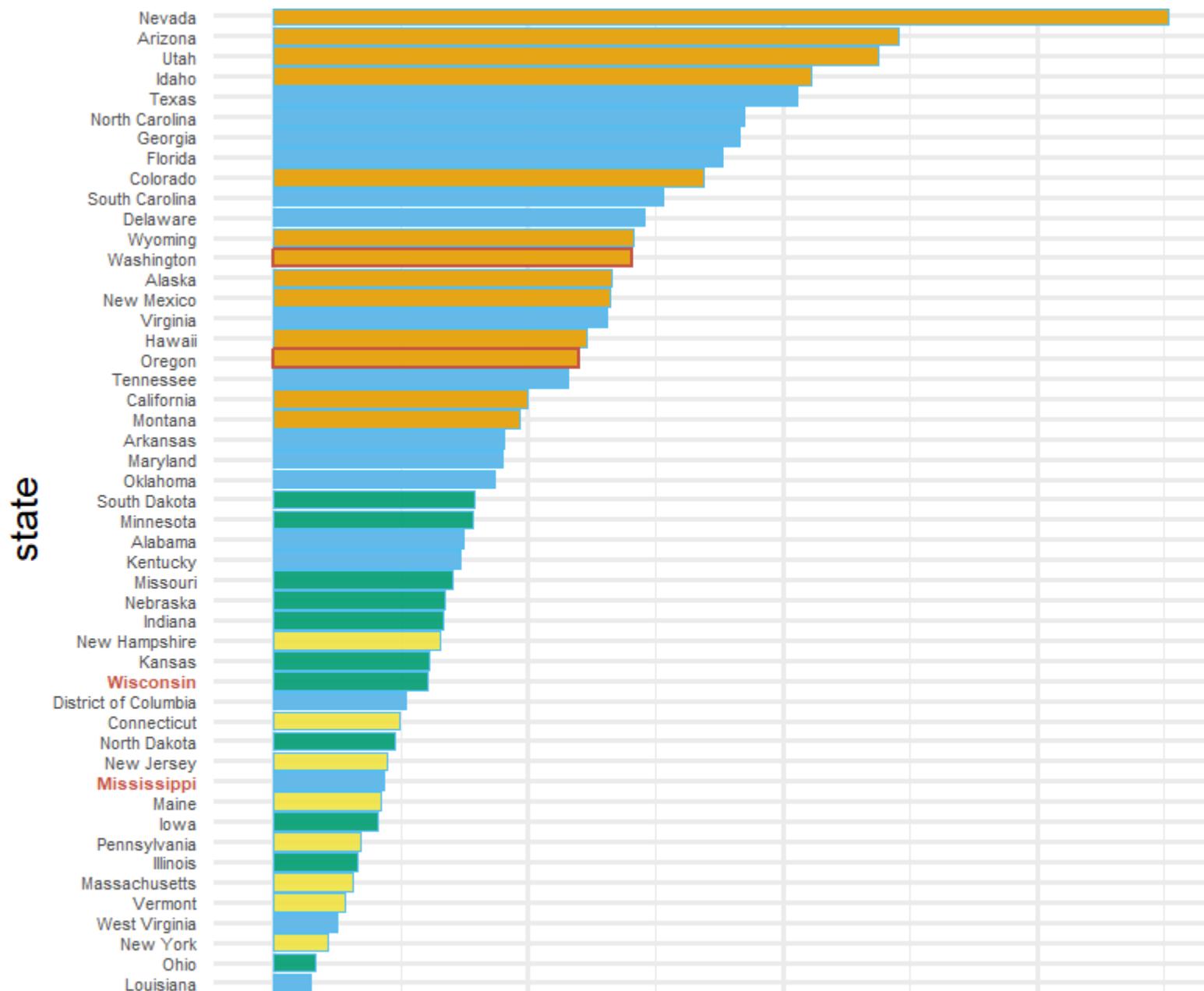
```
## [1] "gray30"  "gray30"
## [15] "gray30"  "gray30"  "gray30"  "#C55644" "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"
## [29] "gray30"  "gray30"
## [43] "gray30"  "gray30"
```

```
label_face <- ifelse(states == "Oregon" | states == "Washington",
                      "bold",
                      "plain")

label_face
```

```
## [1] "plain"  "bold"
## [19] "plain"  "plain"
## [37] "plain"  "plain"
```

```
ggplot(popgrowth_df, aes(x=popgrowth, y=state)) +  
  geom_col(aes(fill = region), alpha = 0.9) +  
  geom_col(data=filter(popgrowth_df, state == 'Oregon' | state == 'Washington'),  
           fill = NA,  
           color = '#C55644',  
           linewidth = 1,  
           width = 0.9) + # Added width parameter to make it slightly narrower  
  scale_fill_OkabeIto() +  
  theme(  
    axis.text.y = element_text(  
      color = label_color,  
      face = label_face  
    )  
  )
```

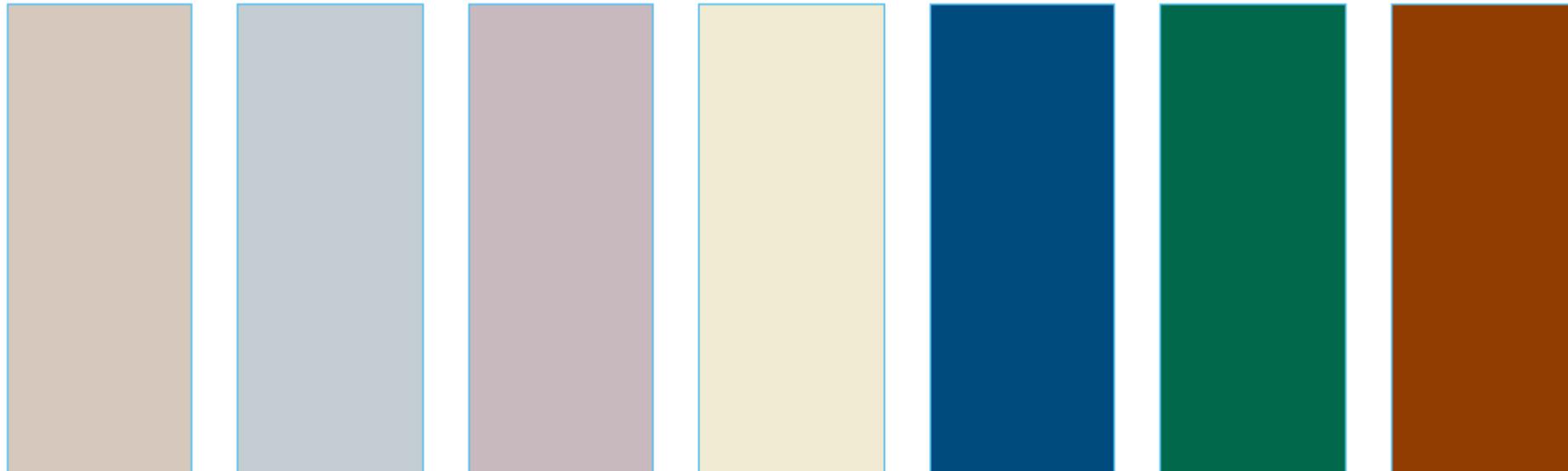


region

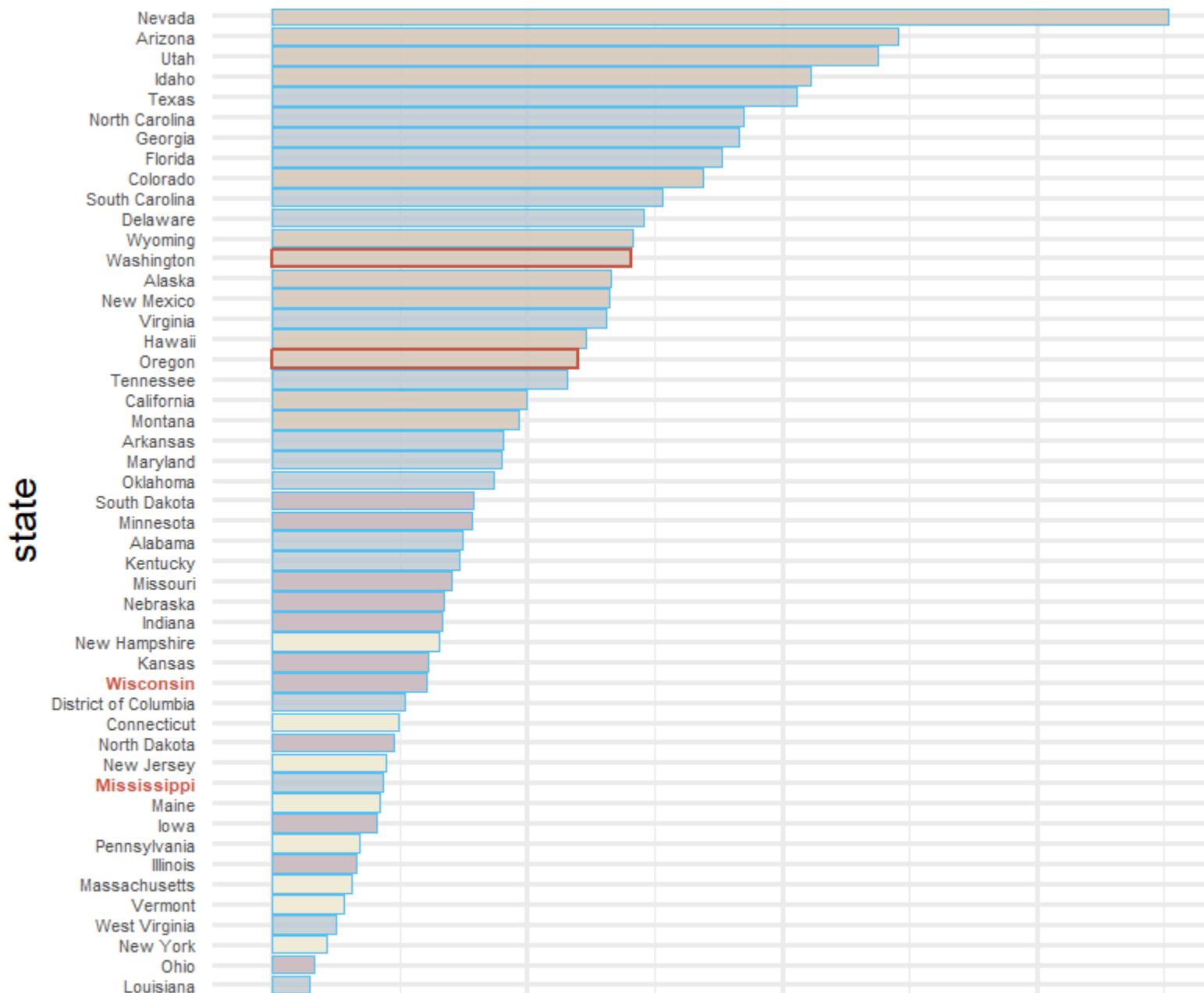
- West
- South
- Midwest
- Northeast

Even better

```
accent_OkabeIto <- palette_OkabeIto[c(1, 2, 7, 4, 5, 3, 6)]
accent_OkabeIto[1:4] <- desaturate(lighten(accent_OkabeIto[1:4], .4), .8)
accent_OkabeIto[5:7] <- darken(accent_OkabeIto[5:7], .3)
gg_color_swatches(7) +
  scale_fill_manual(values = accent_OkabeIto)
```



```
ggplot(popgrowth_df, aes(x=popgrowth, y=state)) +  
  geom_col(aes(fill = region), alpha = 0.9) +  
  geom_col(data=filter(popgrowth_df, state == 'Oregon' | state == 'Washington'),  
            fill = NA,  
            color = '#C55644',  
            linewidth = 1,  
            width = 0.9) +  
  scale_fill_manual(values = accent_OkabeIto) +  
  theme(axis.text.y = element_text(color = label_color,  
                                    face = label_face))
```



region

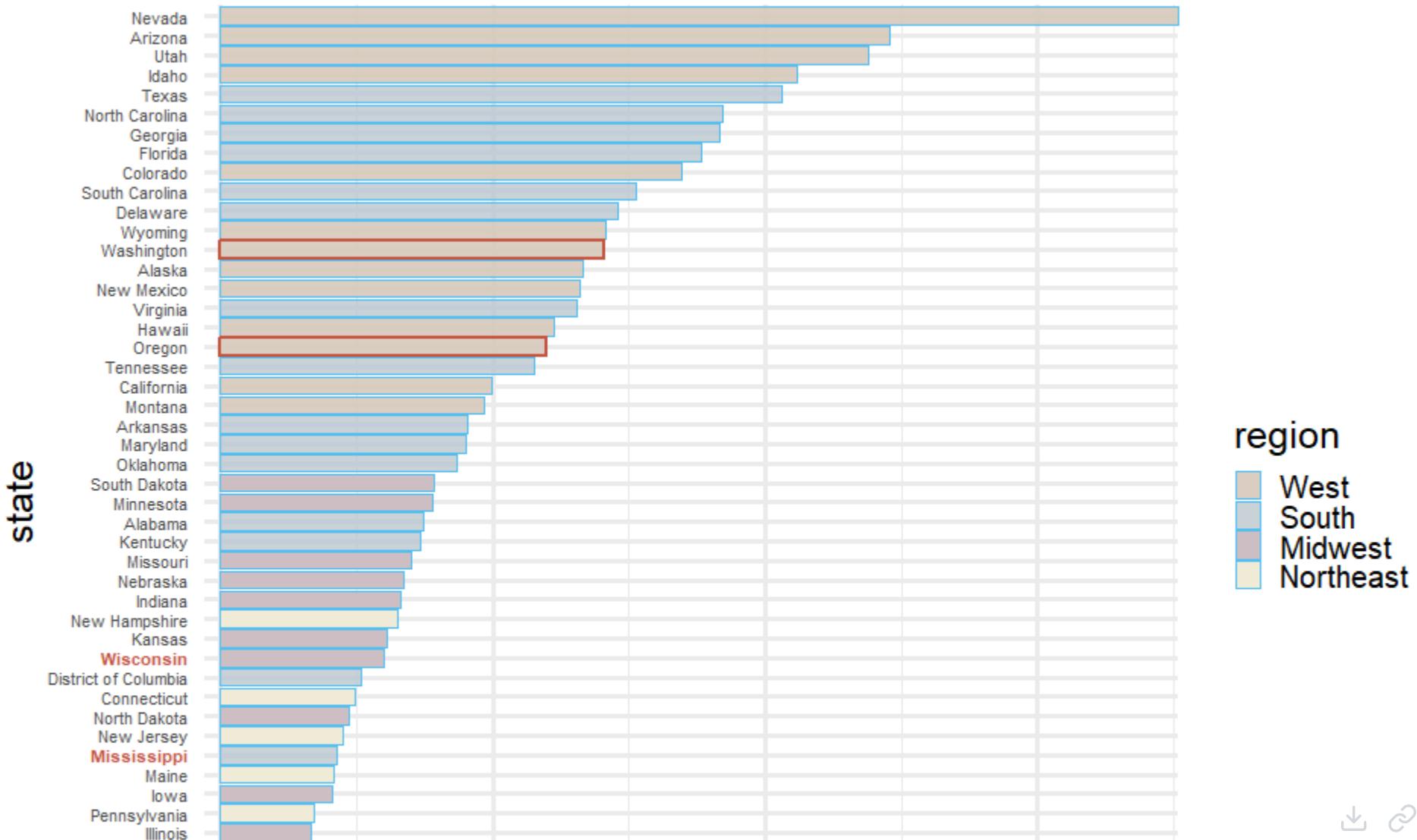
- West
- South
- Midwest
- Northeast

Or even better (imo)

```
library(ggtext)
ggplot(popgrowth_df, aes(x=popgrowth, y=state)) +
  geom_col(aes(fill = region), alpha = 0.9) +
  geom_col(data=filter(popgrowth_df, state == 'Oregon' | state == 'Washington'),
            fill = NA,
            color = '#C55644',
            linewidth = 1,
            width = 0.9) + # Added width parameter to make it slightly narrower
  scale_fill_manual(values = accent_OkabeIto) +
  scale_x_continuous(expand = c(0, 0)) +
  labs(title = "Population growth by region",
       subtitle = "The <span style = 'color: #C55644'>**northwest**</span> is where it's at") +
  theme(axis.text.y = element_text(color = label_color,
                                    face = label_face),
        plot.subtitle = element_markdown())
```

Population growth by region

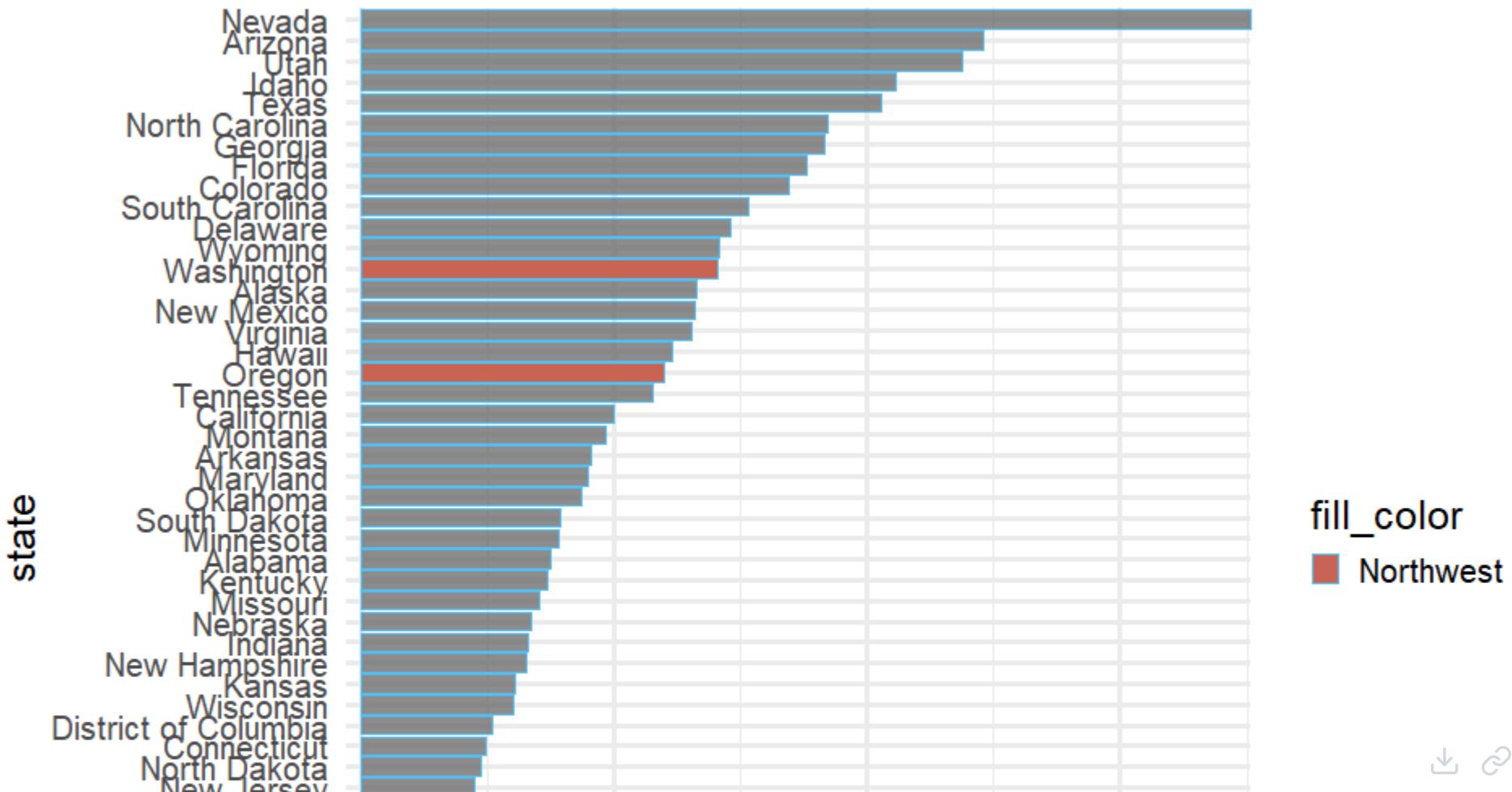
The **northwest** is where it's at



One last attempt

Population growth by region

The **northwest** is where it's at



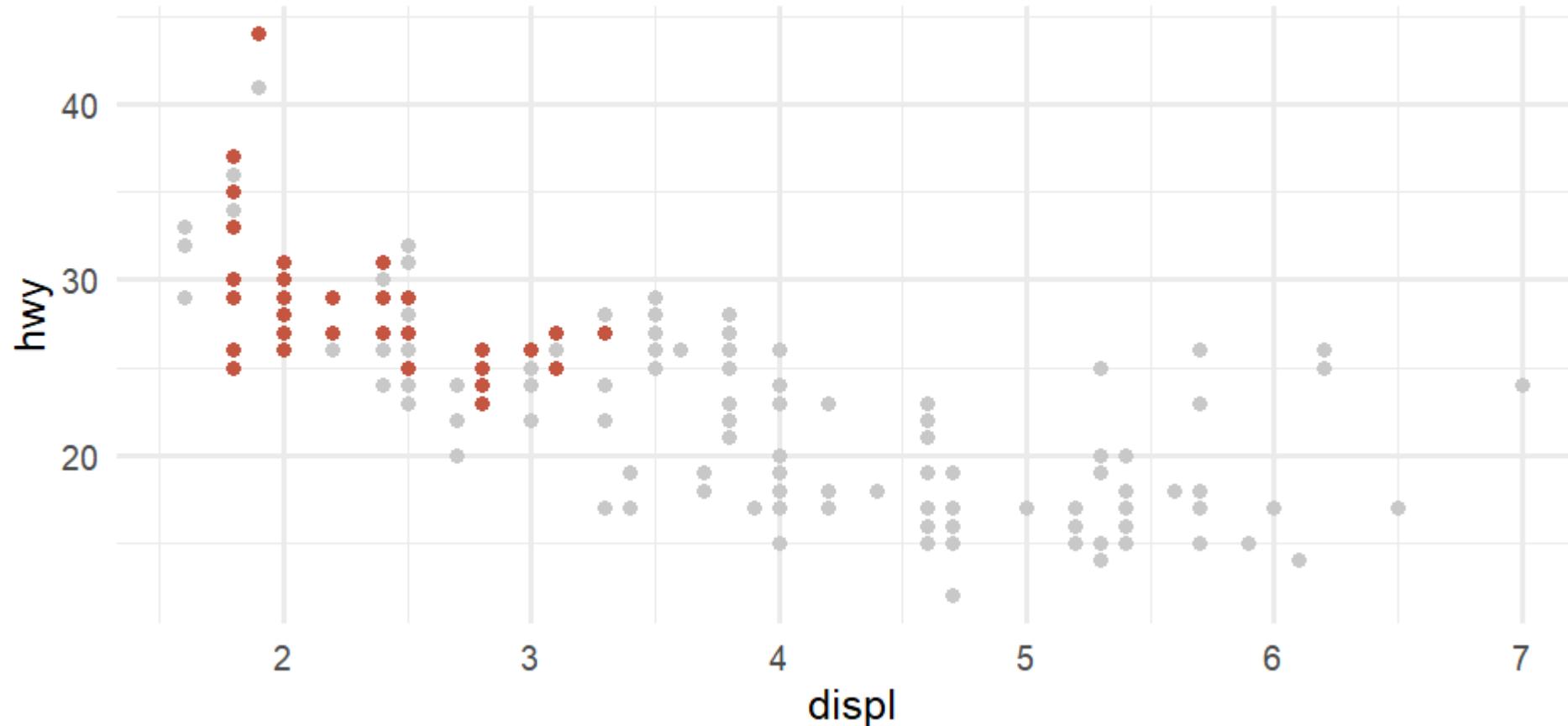
Back to cars plot

We can do this same thing and avoid the legend altogether.

```
ggplot(mpg, aes(displ, hwy)) +  
  geom_point(color = "gray80") +  
  geom_point(data = filter(mpg, class == "compact"),  
             color = "#C55644") +  
  labs(  
    title = "Cars with bigger engines get lower miles per gallon",  
    subtitle = "<span style = 'color: #C55644'>**Compact**</span> cars typically have small engines"  
) +  
  theme(plot.subtitle = element_markdown())
```

Cars with bigger engines get lower miles per gallon

Compact cars typically have small engines



Data viz in Wild

Erick

Emily

Saratessa and Sophia on deck

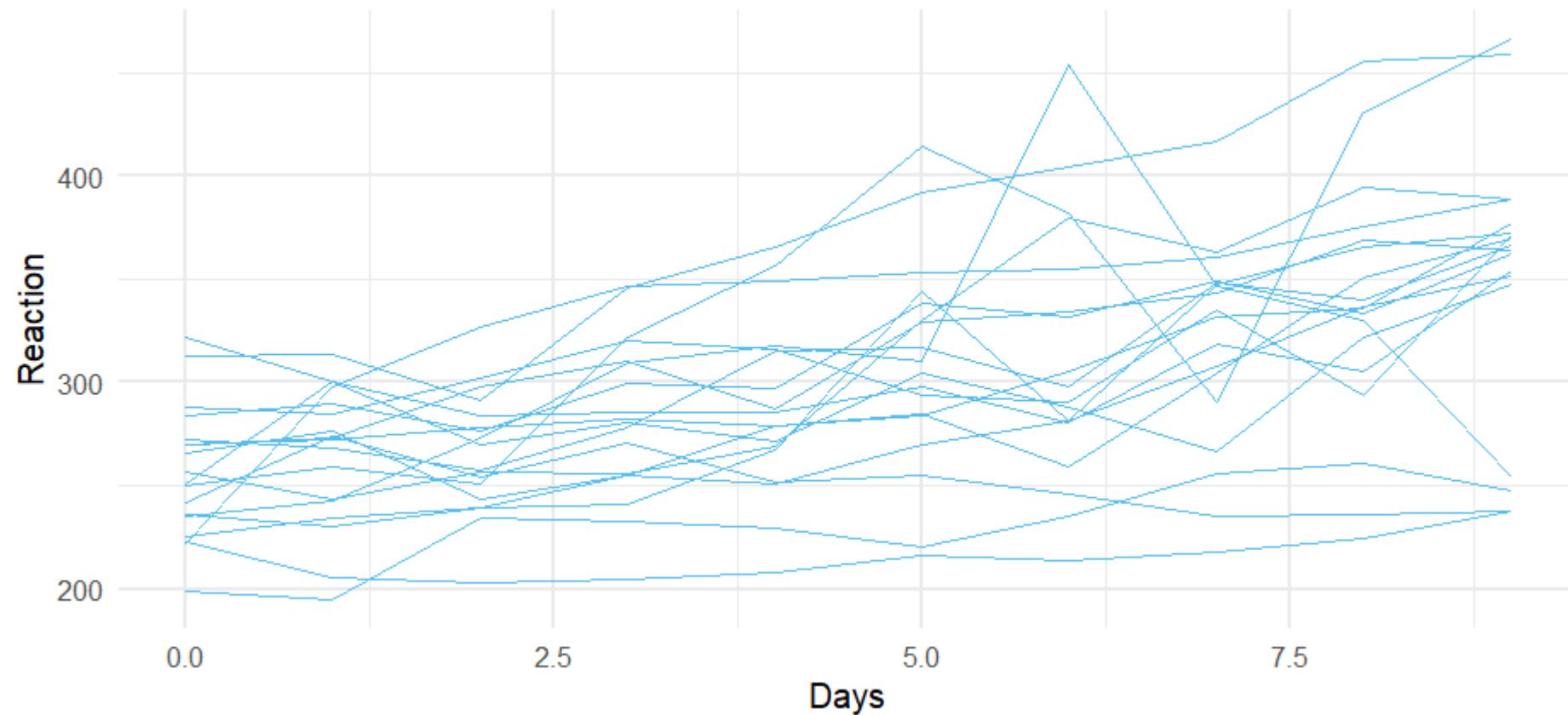
Last example

```
data(sleepstudy, package = "lme4")
head(sleepstudy)
```

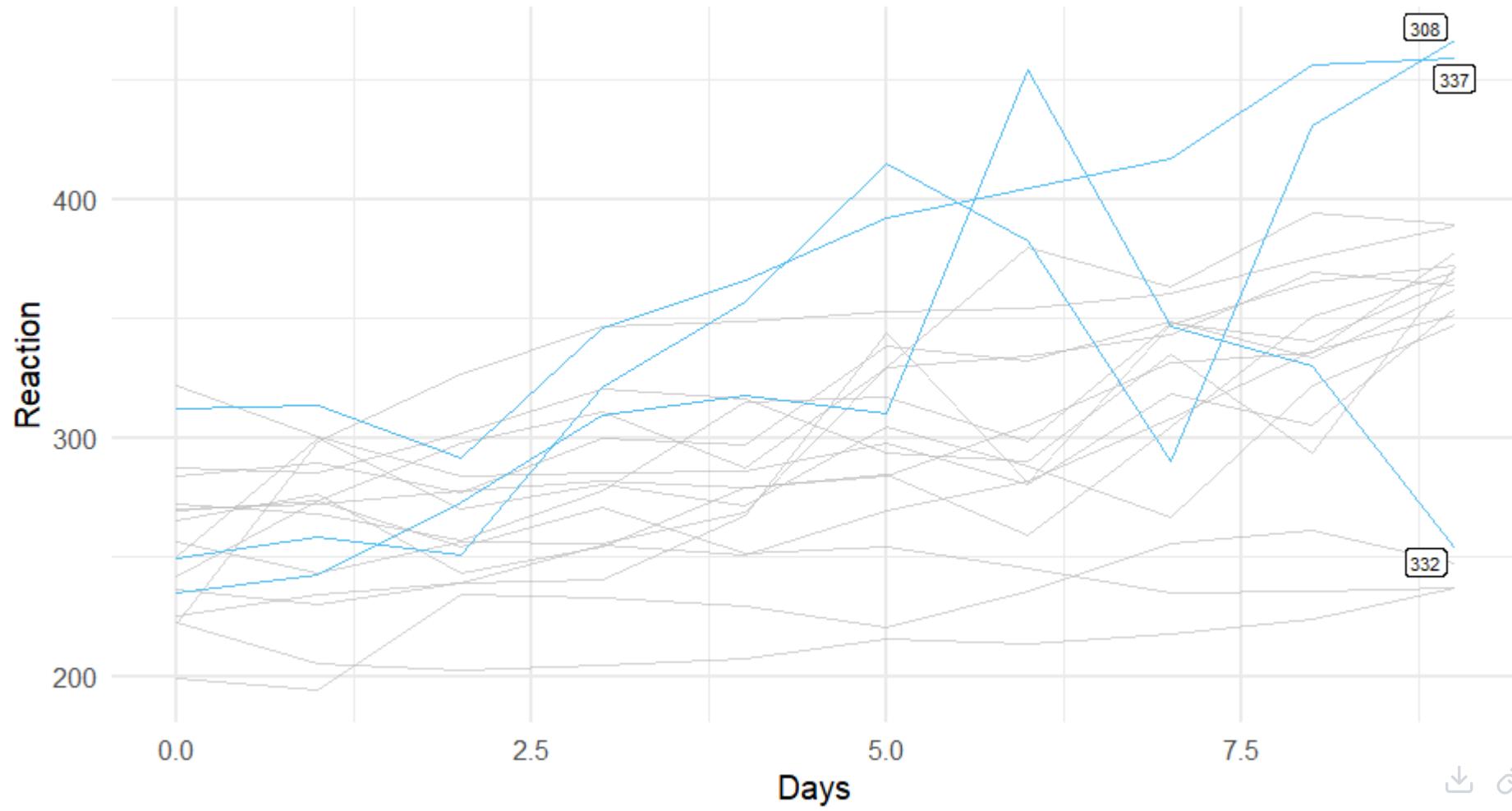
```
##   Reaction Days Subject
## 1      250     0    308
## 2      259     1    308
## 3      251     2    308
## 4      321     3    308
## 5      357     4    308
## 6      415     5    308
```

Plot by subject

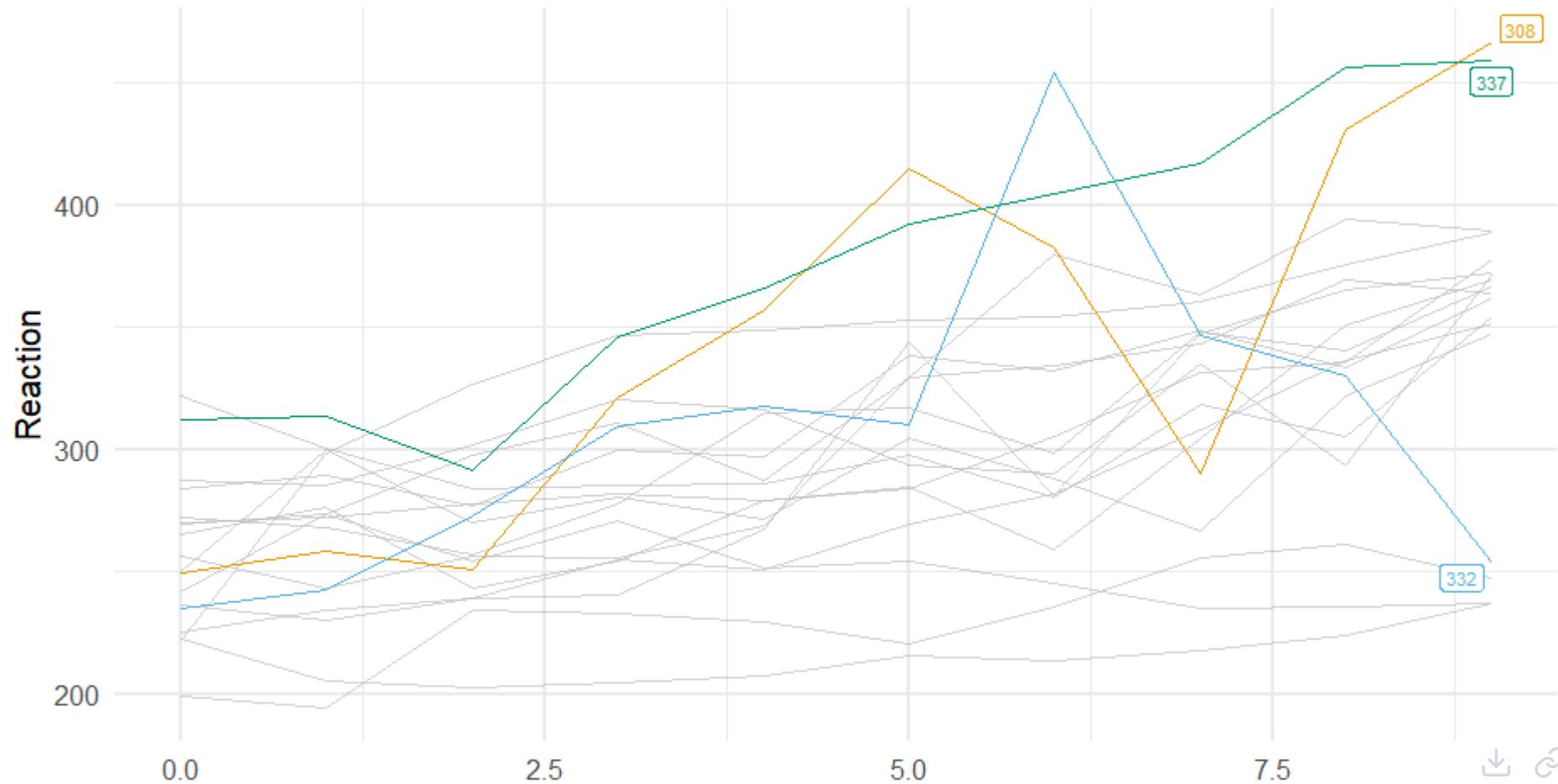
```
ggplot(sleepstudy, aes(Days, Reaction, group = Subject)) +  
  geom_line()
```



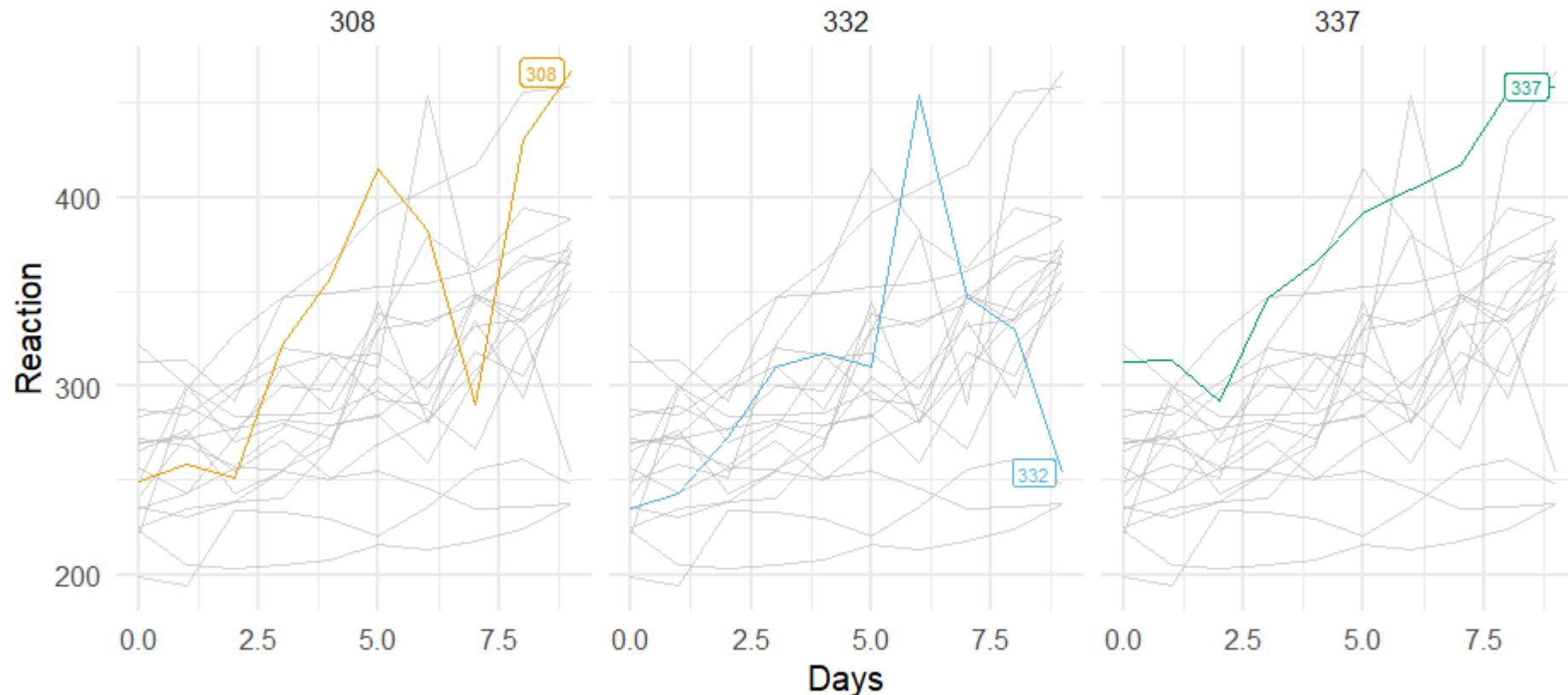
```
library(gghighlight)
ggplot(sleepstudy, aes(Days, Reaction, group = Subject)) +
  geom_line() +
  gghighlight(max(Reaction) > 400)
```



```
library(gghighlight)
ggplot(sleepstudy, aes(Days, Reaction, color = Subject)) +
  geom_line() +
  gghighlight(max(Reaction) > 400) +
  scale_color_OkabeIto()
```



```
library(gghighlight)
ggplot(sleepstudy, aes(Days, Reaction, color = Subject)) +
  geom_line() +
  facet_wrap(~Subject) +
  gghighlight(max(Reaction) > 400) +
  scale_color_OkabeIto()
```



Highlighting

The `gghighlight` package is really neat, but I have had issues with it in the past.

If you have troubles, you can always create them more manually has we did previously

Example

This is a little tricky. First, create a subset of the data that only has the cases you want to highlight

```
# filter
high <- sleepstudy %>%
  filter(Reaction > 400) %>%
  select(Subject)

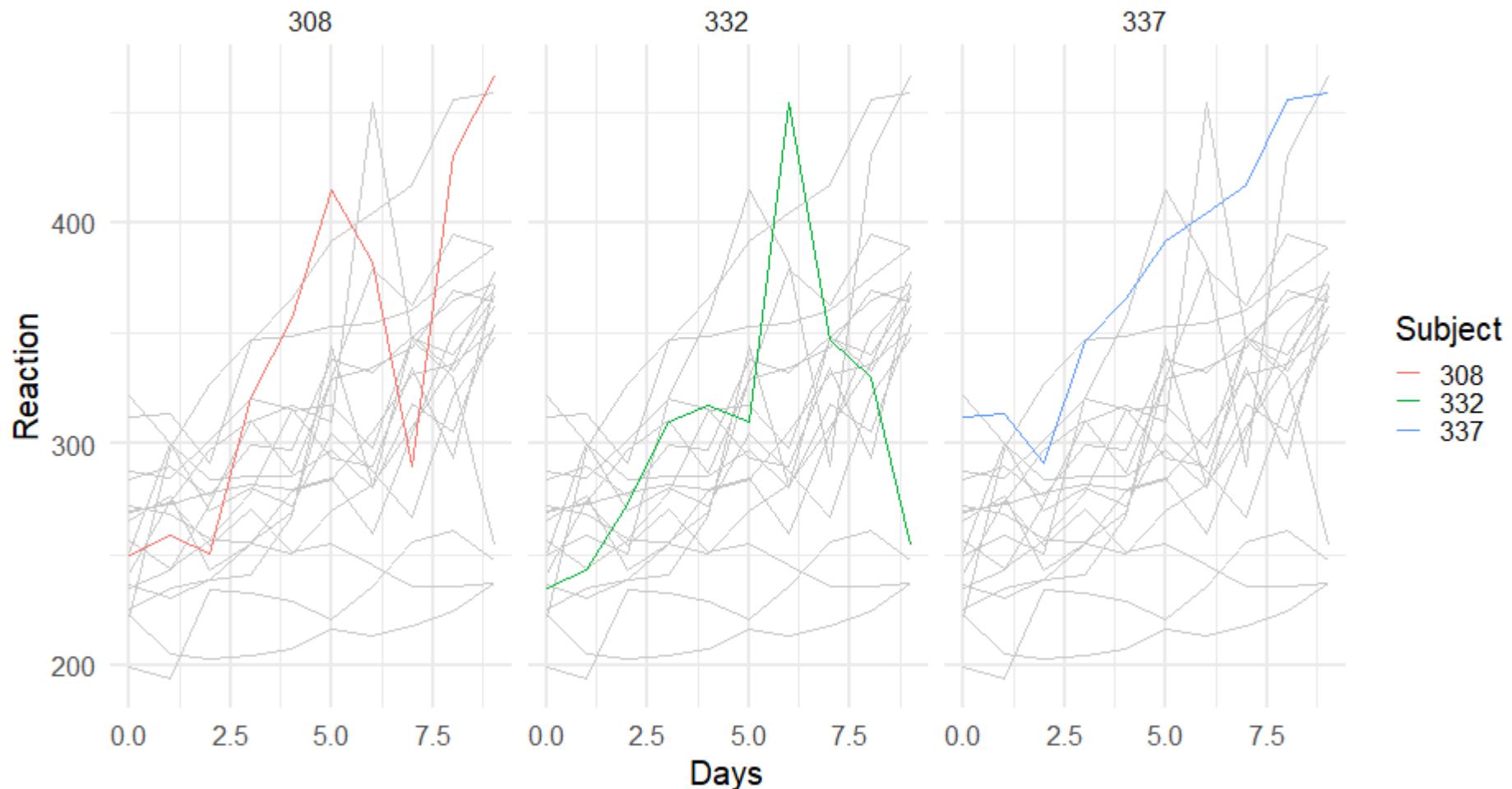
# highlight dataframe
hl <- semi_join(sleepstudy, high)
```

Next, create the background data. We usually just *drop* the faceting column, but in this case we still need the id, so we'll rename it.

```
# rename facet column; use for grouping
bg_data <- sleepstudy %>%
  rename(id = Subject)
```

Finally, plot the highlight data, adding a layer with the background data

```
ggplot(hl, aes(Days, Reaction)) +  
  geom_line(  
    aes(group = id),  
    color = "gray80",  
    data = bg_data  
) +  
  geom_line(  
    aes(color = Subject)  
) +  
  facet_wrap(~Subject)
```

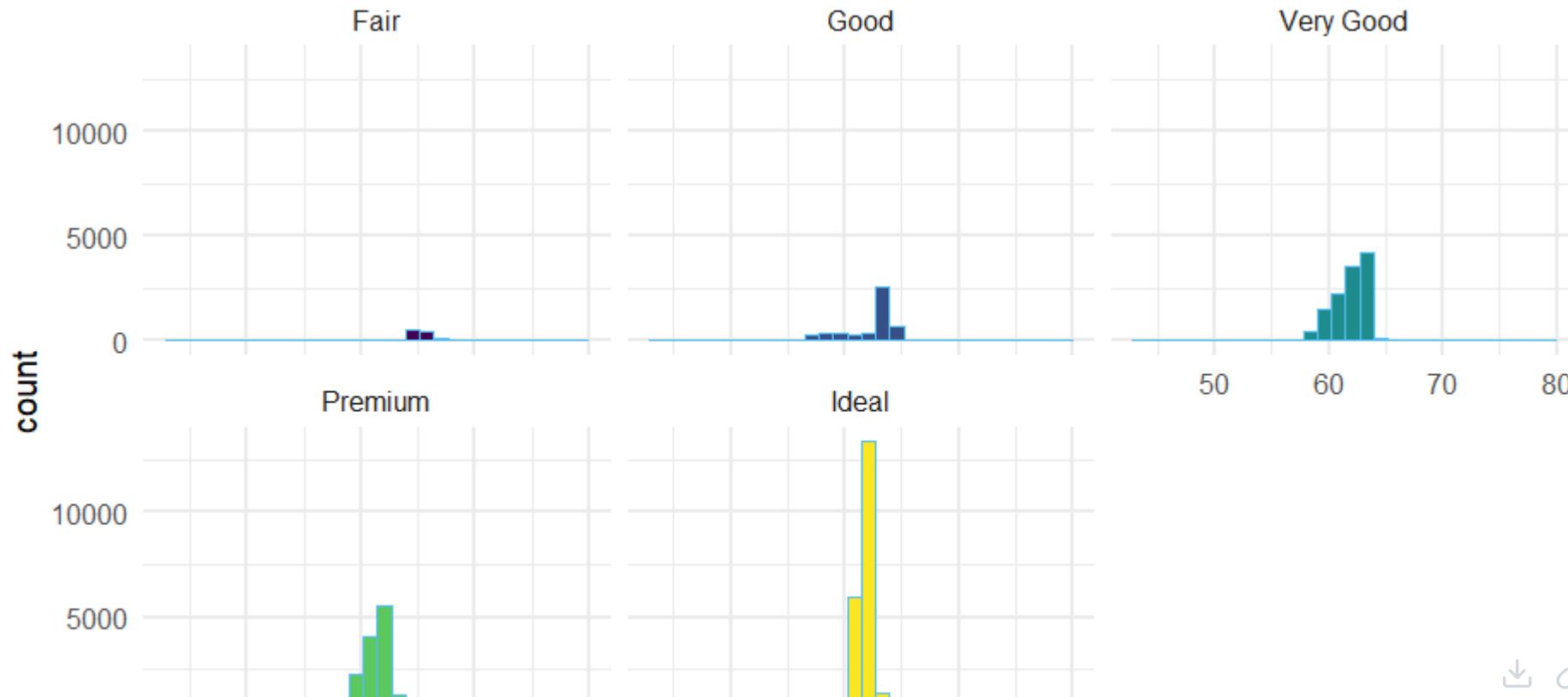


Very quick aside

You can do this with any data - a great alternative to stacked histograms.

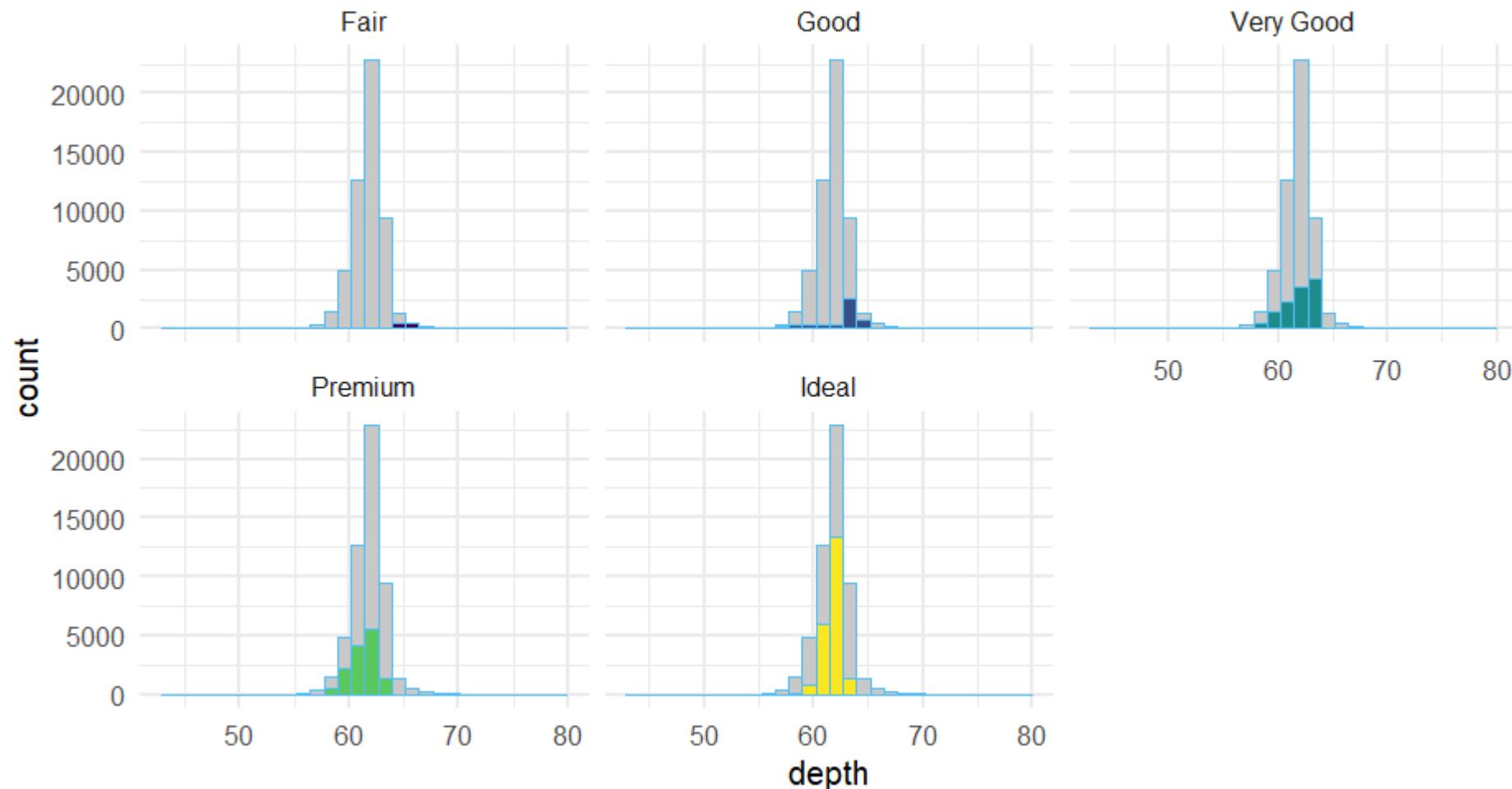
Standard faceted histogram

```
ggplot(diamonds, aes(depth)) +  
  geom_histogram(aes(fill = cut)) +  
  facet_wrap(~cut) +  
  guides(fill = "none") # drop Legend
```



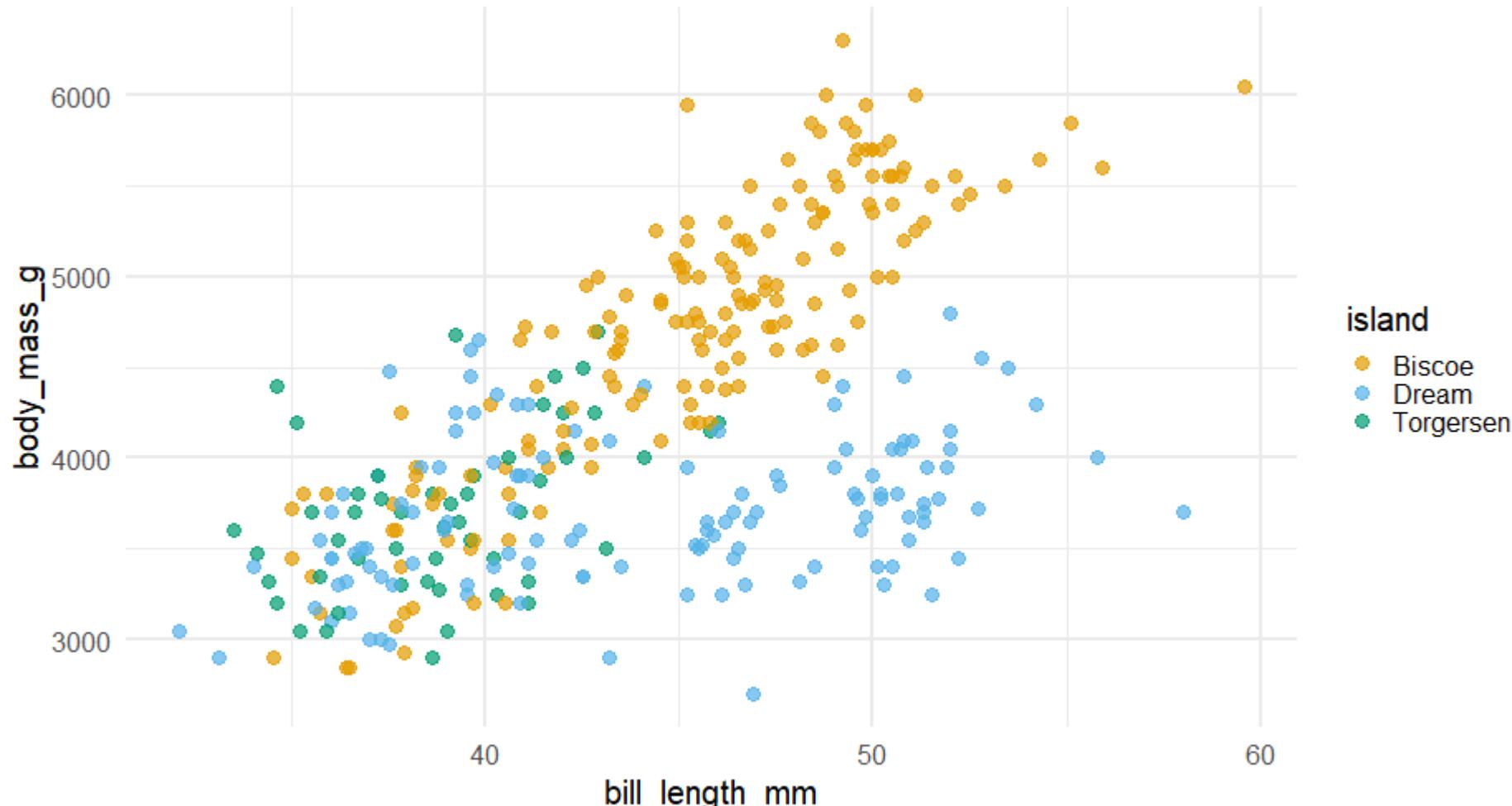
Add a background layer

```
ggplot(diamonds, aes(depth)) +  
  geom_histogram(  
    data = select(diamonds, -cut),  
    fill = "gray80"  
  ) +  
  geom_histogram(aes(fill = cut)) +  
  facet_wrap(~cut) +  
  guides(fill = "none") # drop Legend
```



A few other
things to
consider

Double encodings

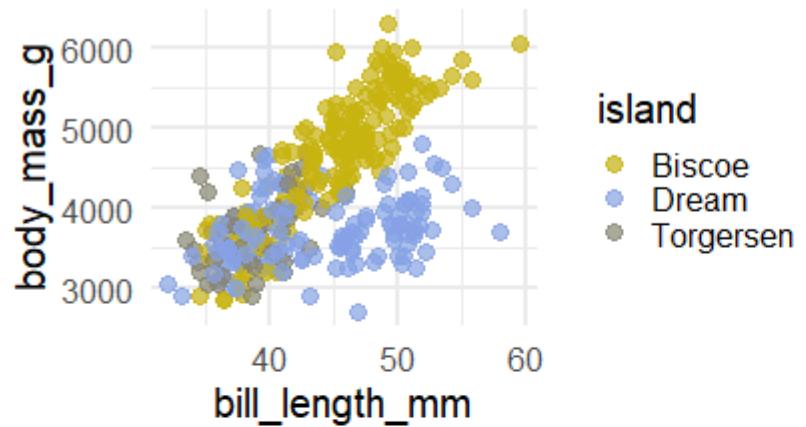


plot is less than ideal. Why?

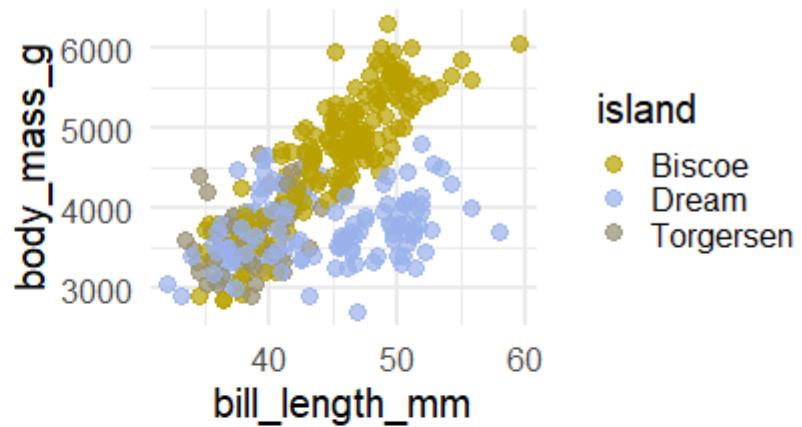
This

Color blindness

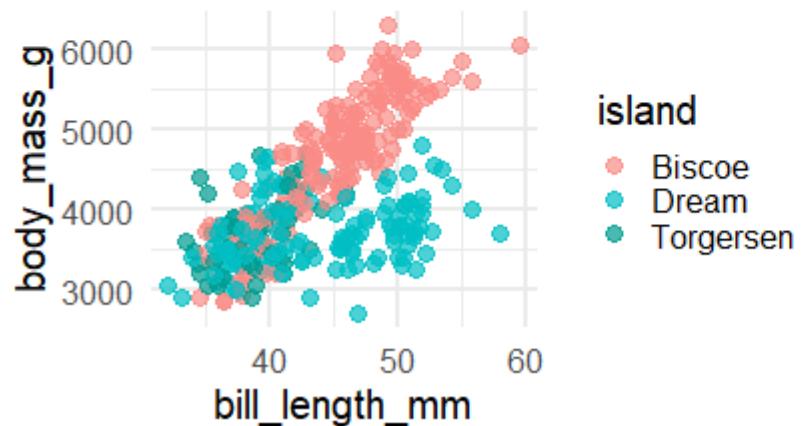
Deutanomaly



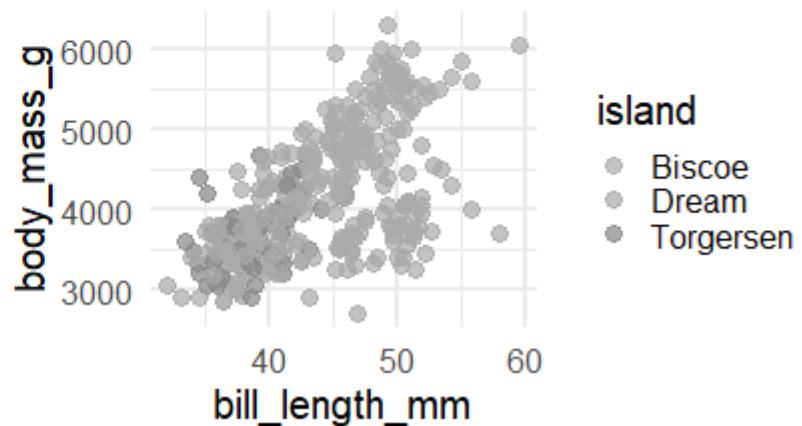
Protanomaly



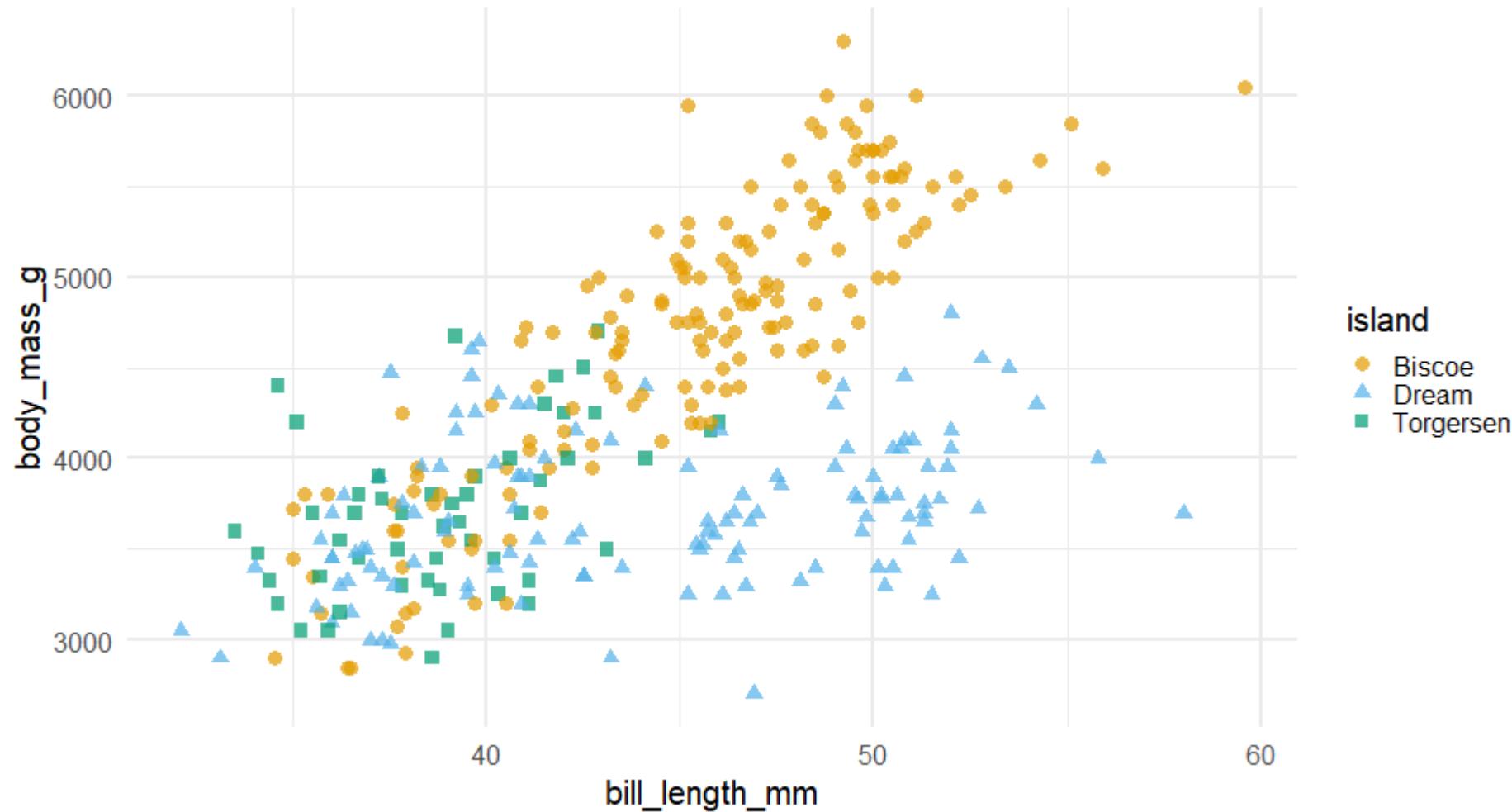
Tritanomaly



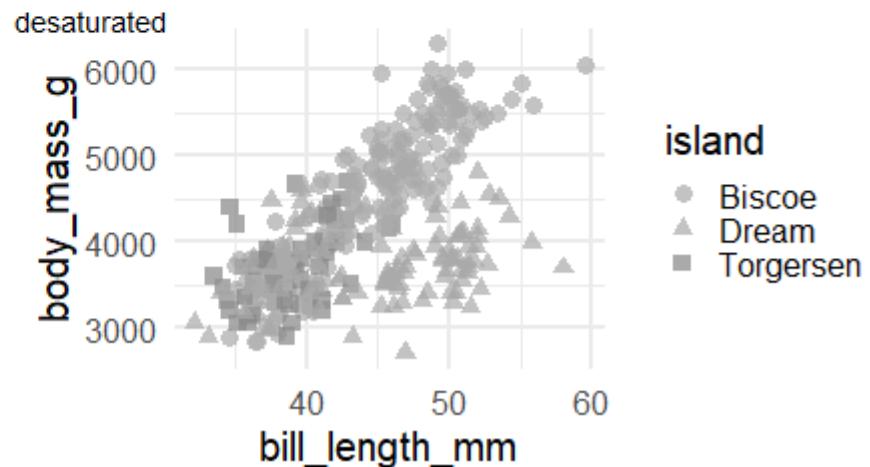
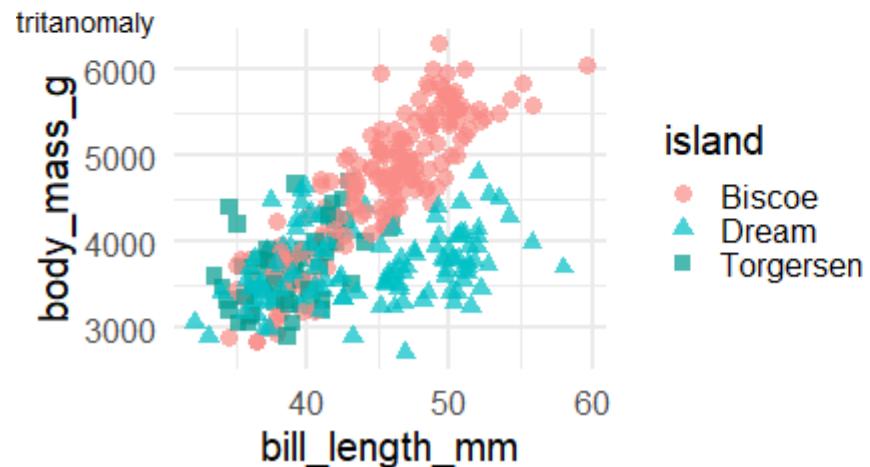
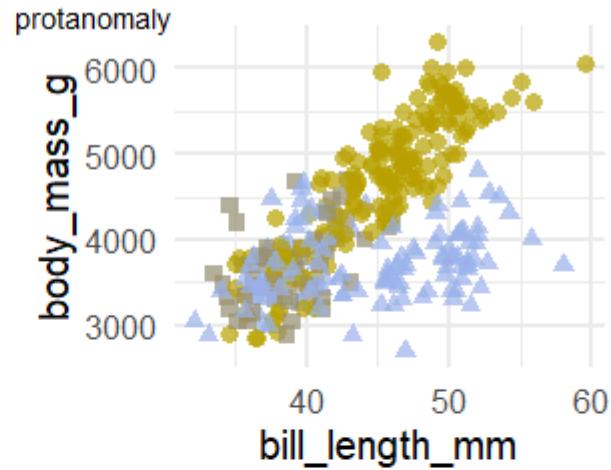
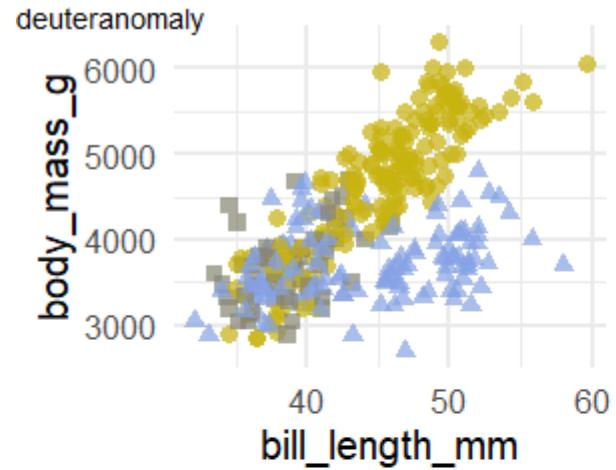
Desaturated



Better version



Color blindness check

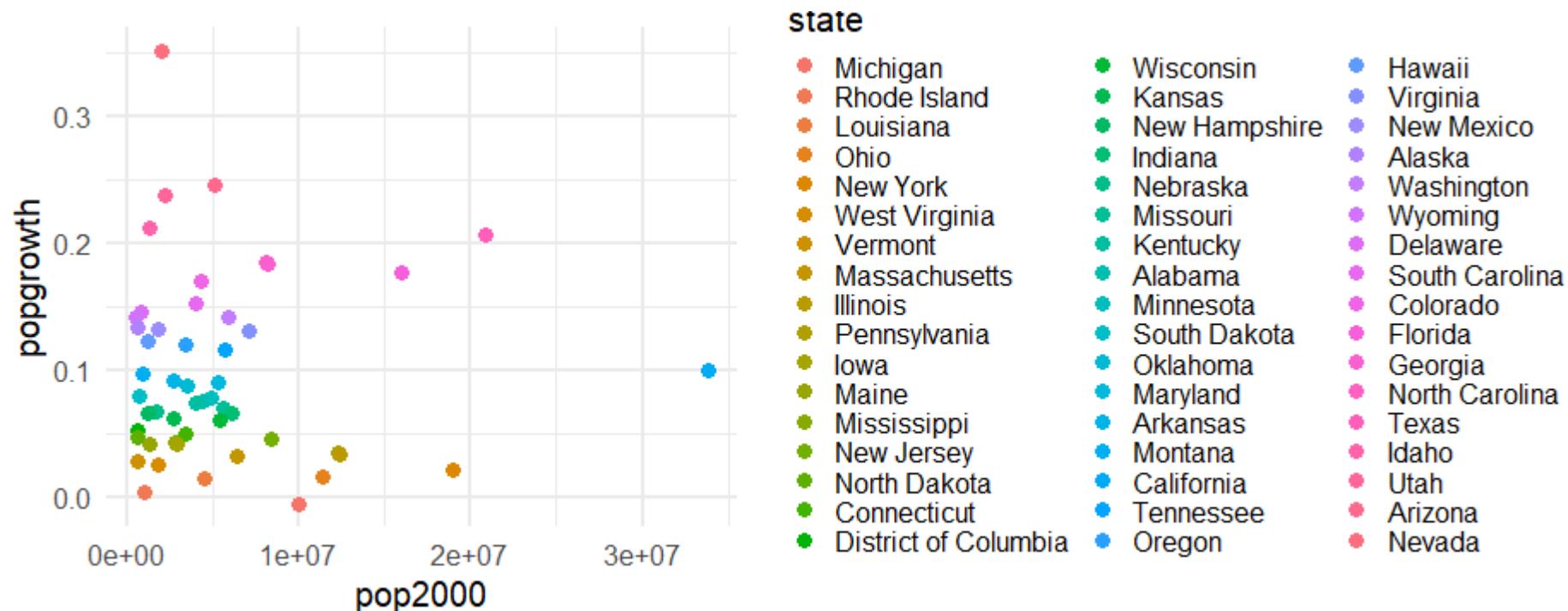


Common problems with color

Too many colors

More than 5-ish categories generally becomes too difficult to track

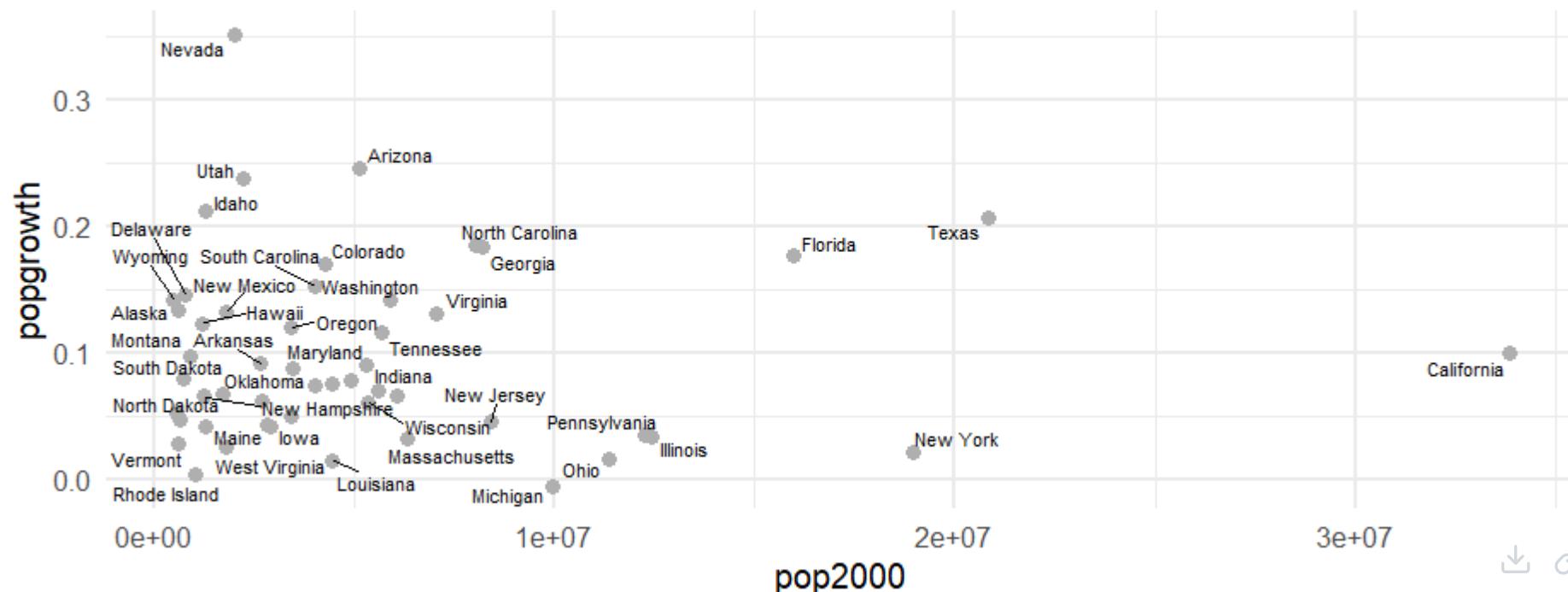
```
ggplot(popgrowth_df, aes(pop2000, popgrowth, color = state)) +  
  geom_point()
```



Use labels

More than 5-ish categories generally becomes too difficult to track

```
library(ggrepel)  
  
ggplot(popgrowth_df, aes(pop2000, popgrowth)) +  
  geom_point(color = "gray70") +  
  geom_text_repel(aes(label = state))
```



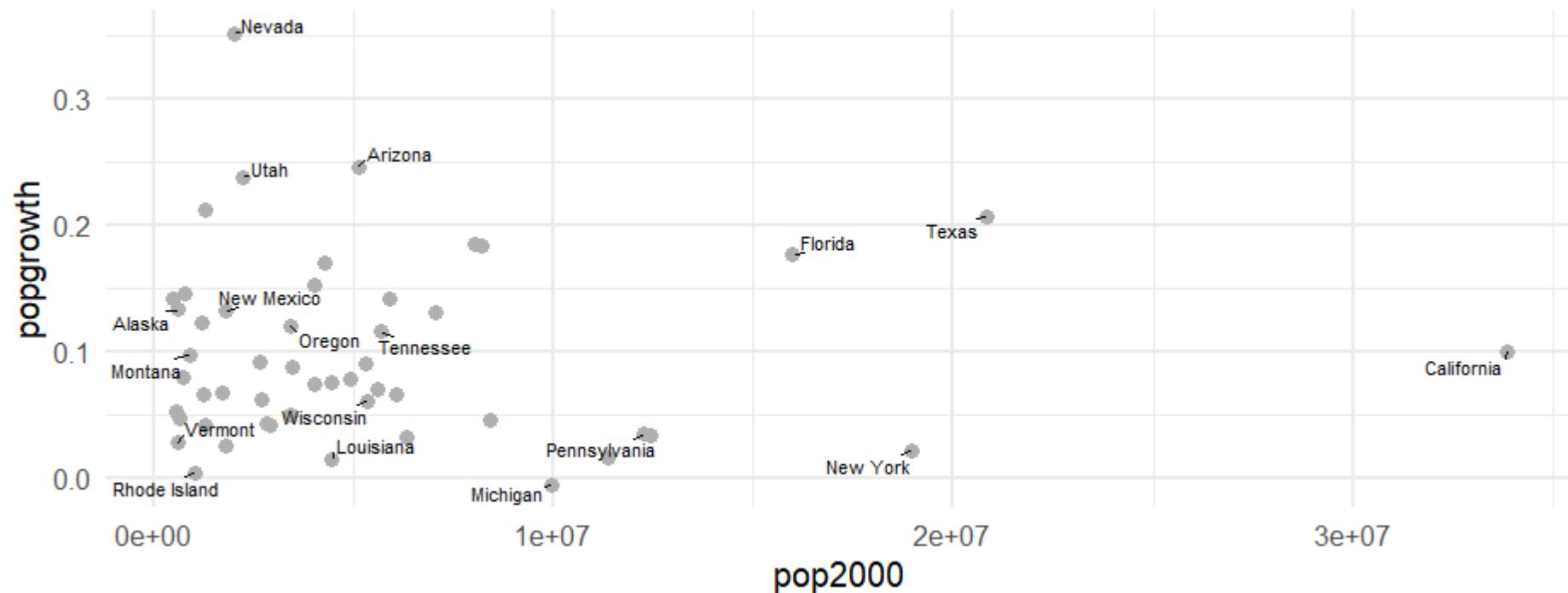
Better

Get a subset

```
to_label <- c("Alaska", "Arizona", "California", "Florida", "Wisconsin",
            "Louisiana", "Nevada", "Michigan", "Montana", "New Mexico",
            "Pennsylvania", "New York", "Oregon", "Rhode Island",
            "Tennessee", "Texas", "Utah", "Vermont")
subset_states <- popgrowth_df %>%
  filter(state %in% to_label)
```

```
library(ggrepel)
```

```
ggplot(popgrowth_df, aes(pop2000, popgrowth)) +  
  geom_point(color = "gray70") +  
  geom_text_repel(aes(label = state),  
                 data = subset_states,  
                 min.segment.length = 0)
```



(still lots more cleaning up we could do here...)

Rainbow palette

```
rainbow(3)
```

```
## [1] "#FF0000" "#00FF00" "#0000FF"
```

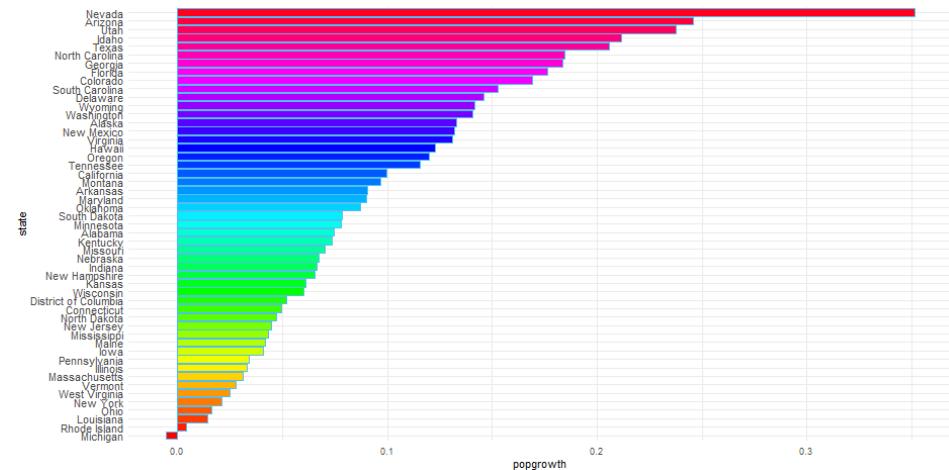
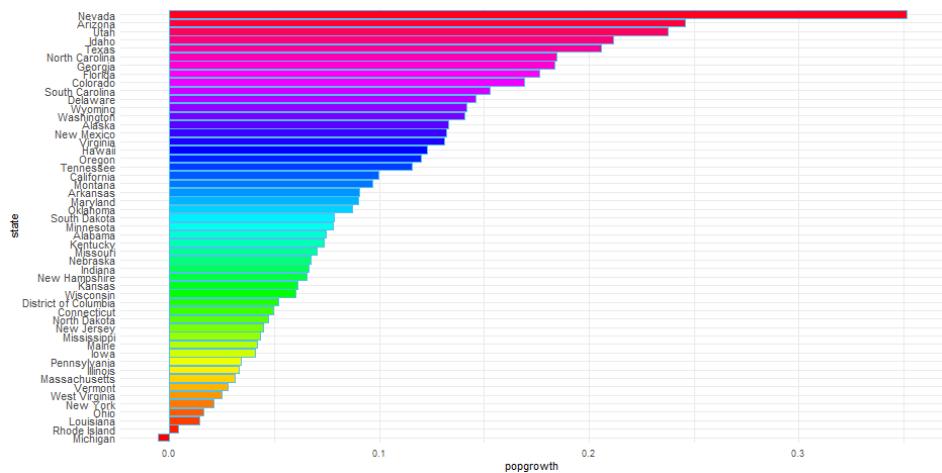
```
rainbow(7)
```

```
## [1] "#FF0000" "#FFDB00" "#49FF00" "#00FF92" "#0092FF" "#4900FF" "#FF00DB"
```

Pretty! Doesn't work well

See [here](#) for one (of many) articles on why this is the case

```
ggplot(popgrowth_df, aes(x = popgrowth, y = state)) +  
  geom_col(aes(fill = state)) +  
  scale_fill_manual(values = rainbow(51)) +  
  guides(fill = "none")
```

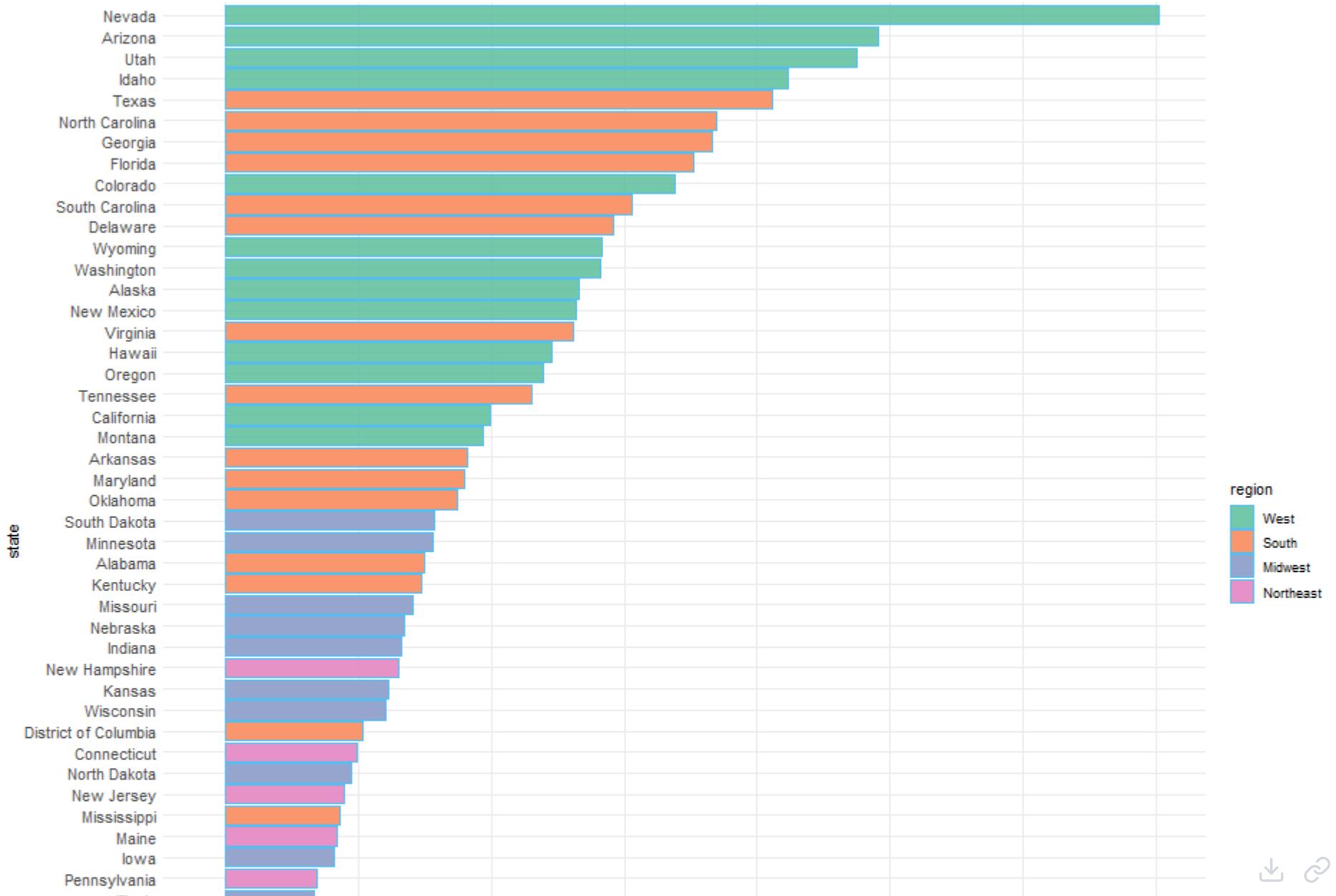


Last few notes on palettes

- Do some research, find what you like **and** what tends to work well
- Check for colorblindness
- Look into <http://colorbrewer2.org/>
 - `scale_color_brewer()` and `scale_fill_brewer()` ship with ggplot2

For example

```
ggplot(popgrowth_df, aes(x = popgrowth, y = state)) +  
  geom_col(aes(fill = region),  
           alpha = 0.9) +  
  scale_fill_brewer(palette = "Set2")
```



Paleteer package



The anatomy of a ggplot() theme

Theme system

ggplot2 Theme Elements

```
theme(element_name = element_function())
- element_text()
- element_line()
- element_rect()
- element_blank()
```

Axis elements:

axis.ticks
element_line()

axis.title
element_text()

axis.text
element_text()

axis.line
element_line()

By Henry Wang

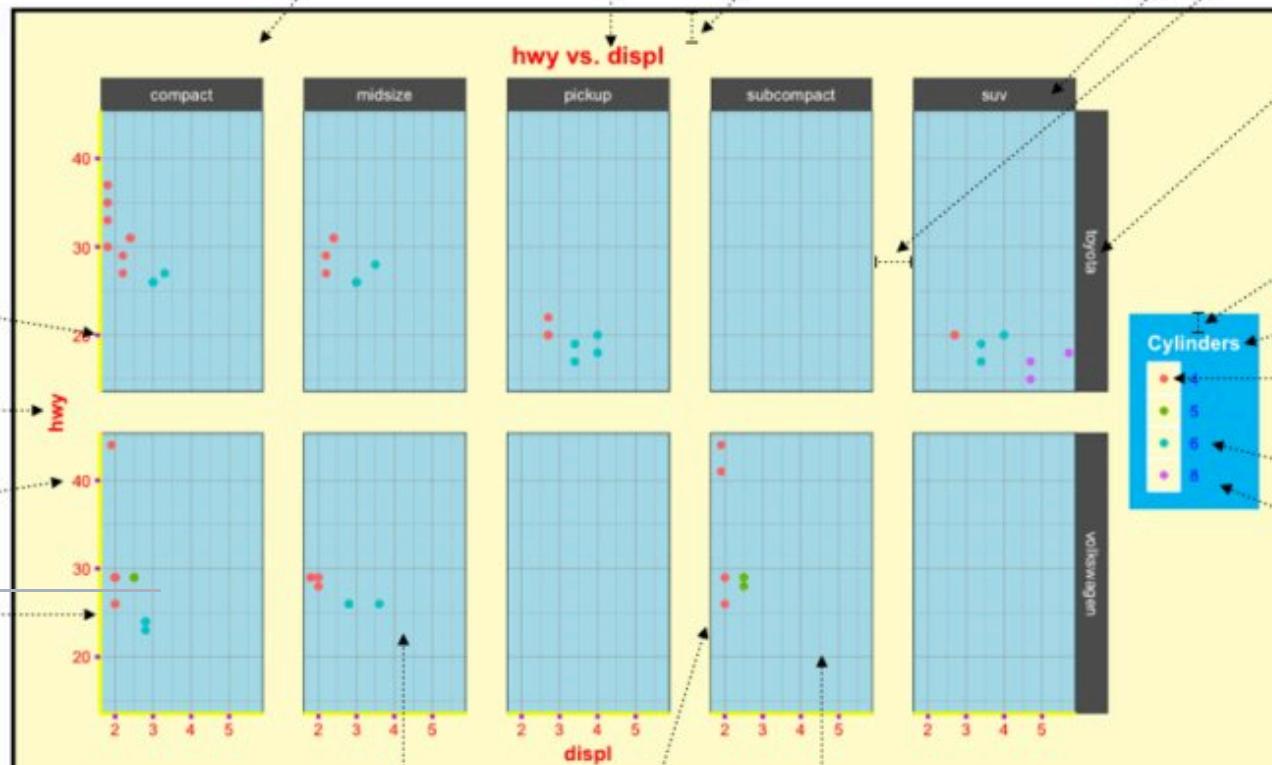
Plot elements:

plot.background
element_rect()

plot.title
element_text()

plot.margin
margin()

hwy vs. displ



Facetting elements:

strip.background
element_rect()

panel.spacing
unit()

strip.text
element_text()

Legend elements:

legend.margin
margin()

legend.title
element_text()

legend.key
element_rect()

legend.text
element_text()

legend.background
element_rect()

Theme elements

- Each element in the plot can be targeted
- Plot title = `plot.title`
- Grid lines = `panel.grid`
- Legend background = `legend.background`

Theme functions

- Use special functions to manipulate specific elements
- Text-based things = `element_text()`
- Rectangular things (backgrounds) = `element_rect()`
- Line-based things (axis lines, grid lines) = `element_line()`
- Disable element completely = `element_blank()`

How to learn `theme()`

- The `theme()` function has **94** possible arguments(!!!)
- You can get hyper-specific with things like `axis.ticks.length.x.bottom`
- The only way to learn how to use `theme()` is to use it and tinker with it

Review Lab 3

Lab 4

Q & A with Dr. Daniel Anderson

Next time

More colors, themes, refinement of plots