

Colors!

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Week 5, Class 1

Data viz in Wild

Chris

Vinita

Shijing and David on deck

Agenda

- Color basics
 - 3 basic ways color is used
- Color blindness
- Some common problems with color use
- Quick discussion of palettes

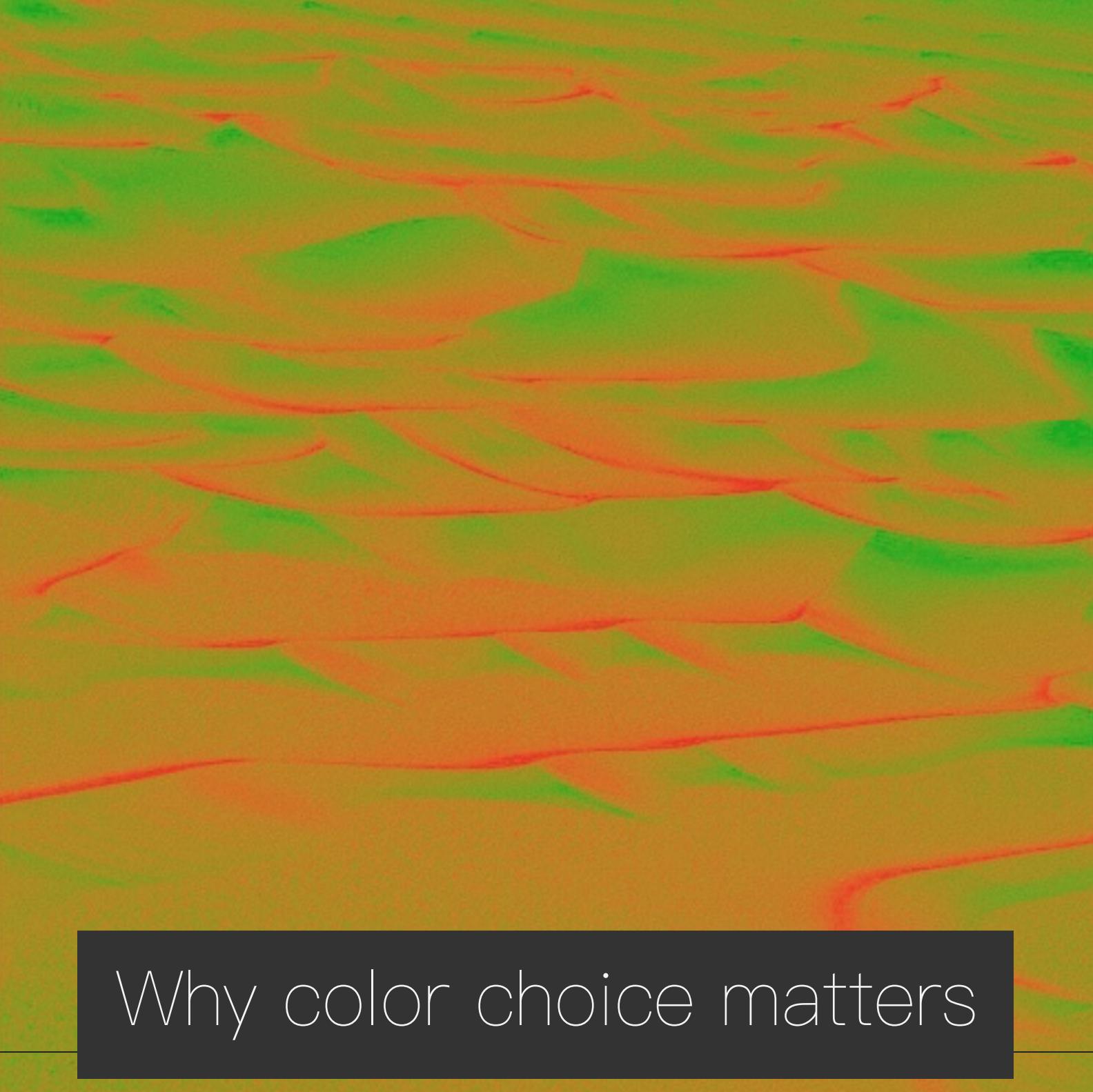
Learning Objectives

- 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

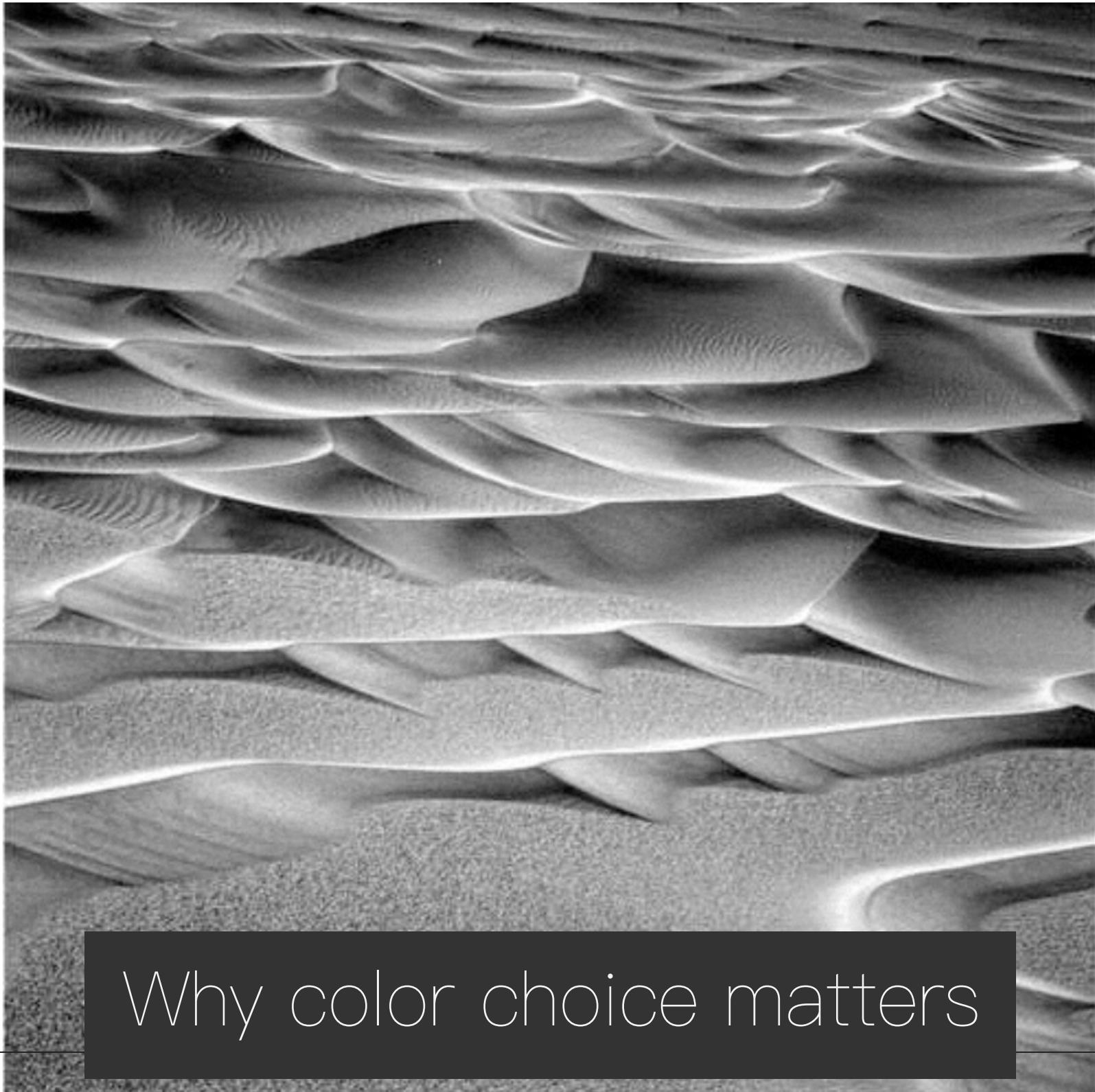
Before we get too deep

Some very practical advice

- 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?

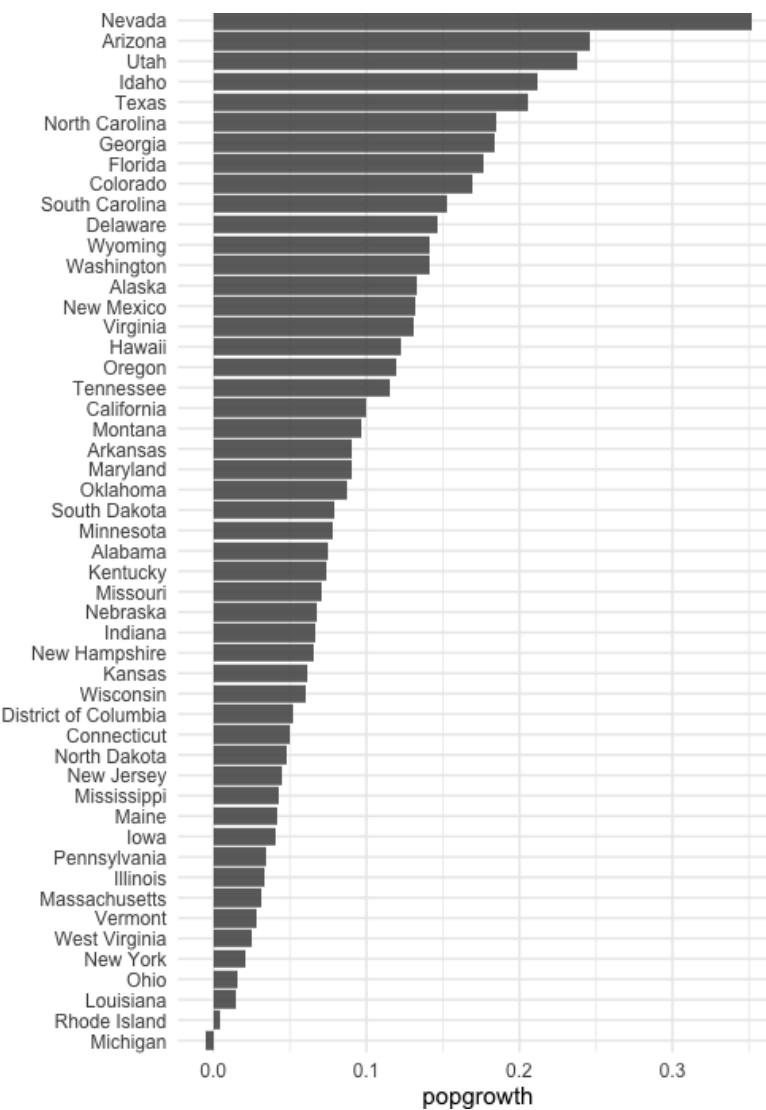
Imagine we have data like this

```
popgrowth_df
```

```
## # A tibble: 51 x 7
##   region      division       state    pop2000    pop2010  popgrowth
##   <fct>        <chr>        <fct>     <dbl>      <dbl>      <dbl>
## 1 Midwest     East North Central Michigan  9938444  9883640 -0.005514344
## 2 Northeast   New England      Rhode Island 1048319  1052567  0.004052202
## 3 South        West South Central Louisiana 4468976  4533372  0.01440956
## 4 Midwest     East North Central Ohio      11353140 11536504  0.01615095
## 5 Northeast   Middle Atlantic    New York   18976457 19378102  0.02116544
## 6 South        South Atlantic    West Virginia 1808344  1852994  0.02469110
## # ... with 45 more rows, and 1 more variable: area <dbl>
```

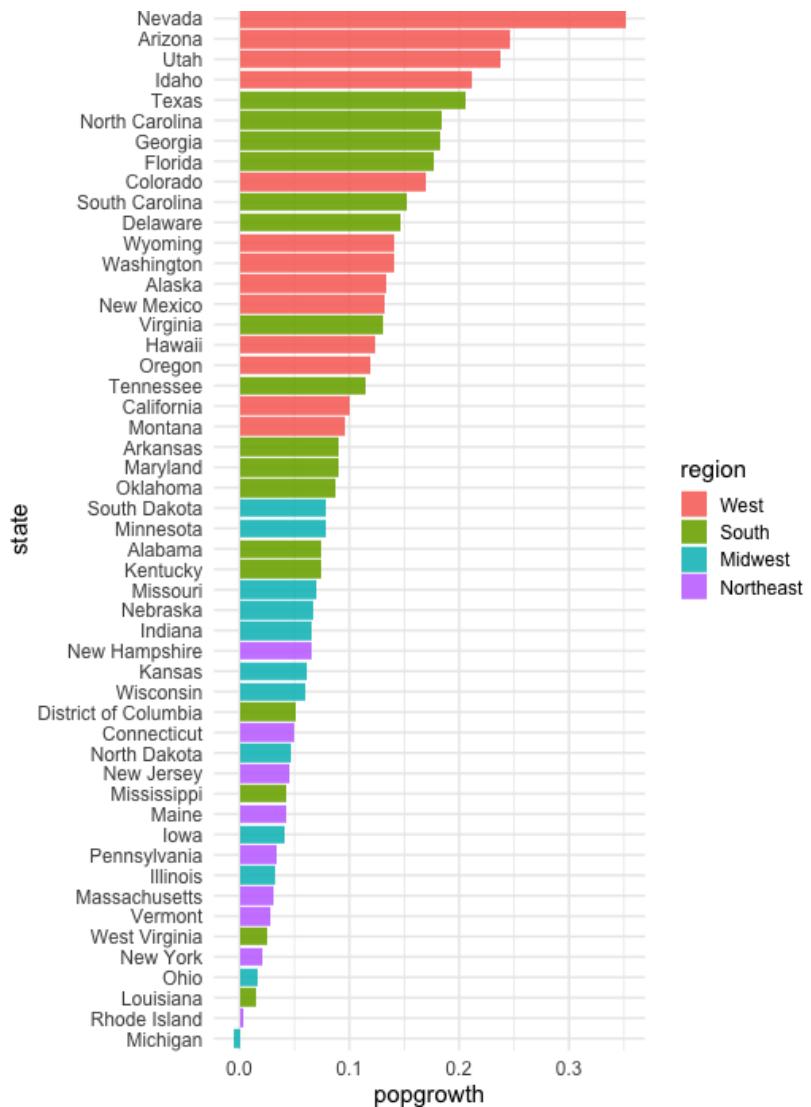
Maybe a plot like this

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

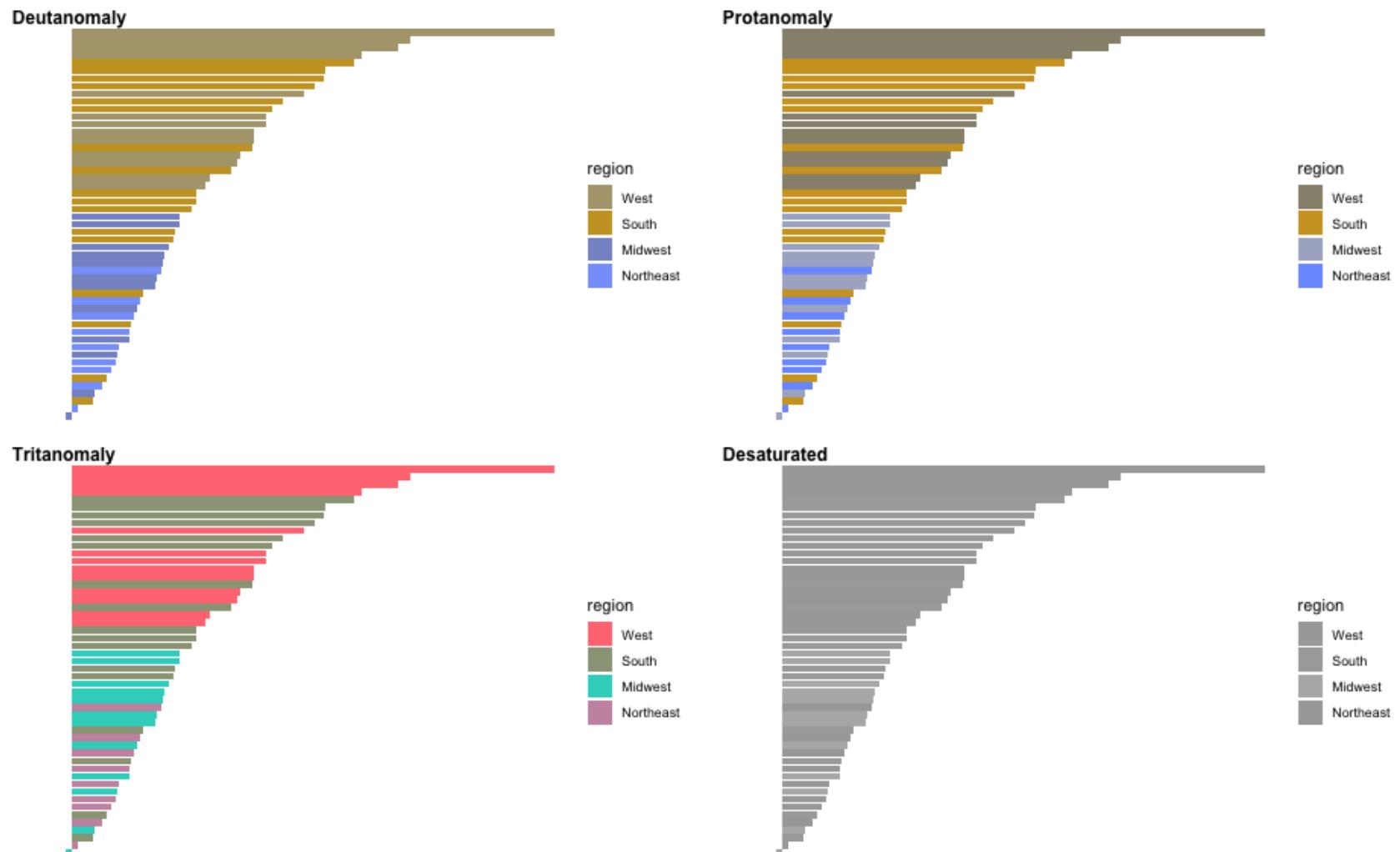


Alternatively, fill by region

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

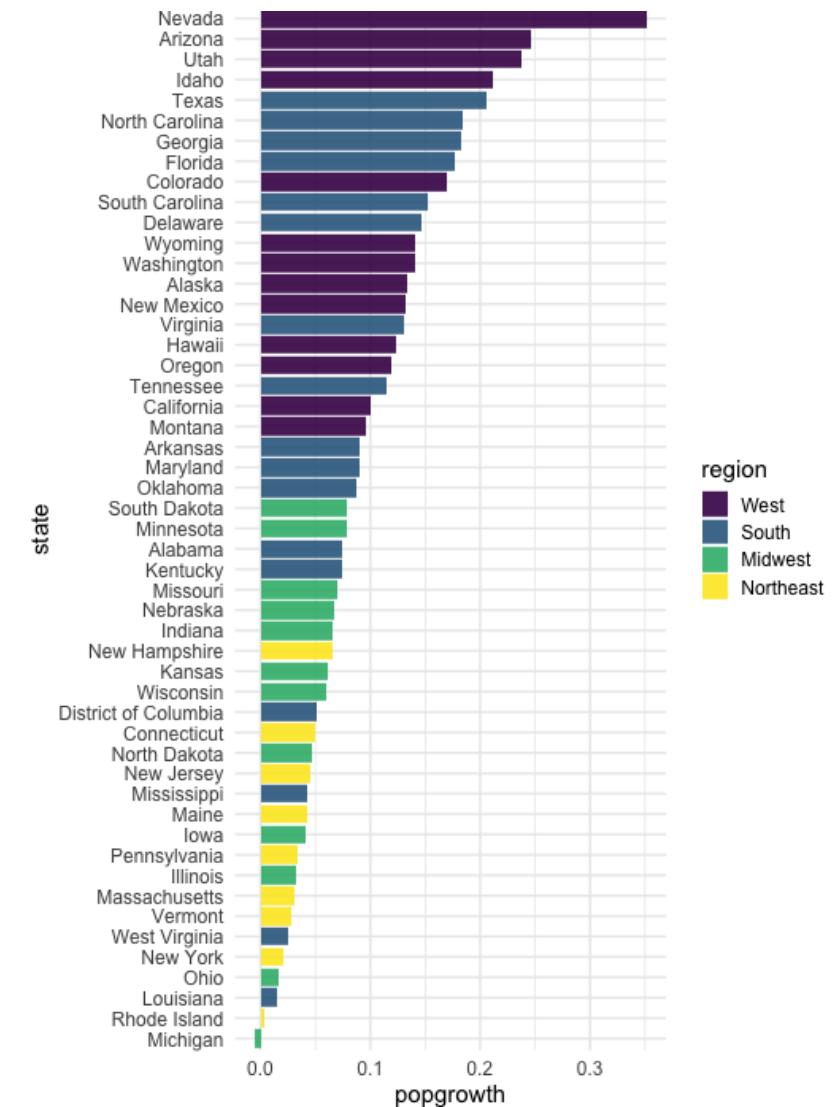


Problem with default palette

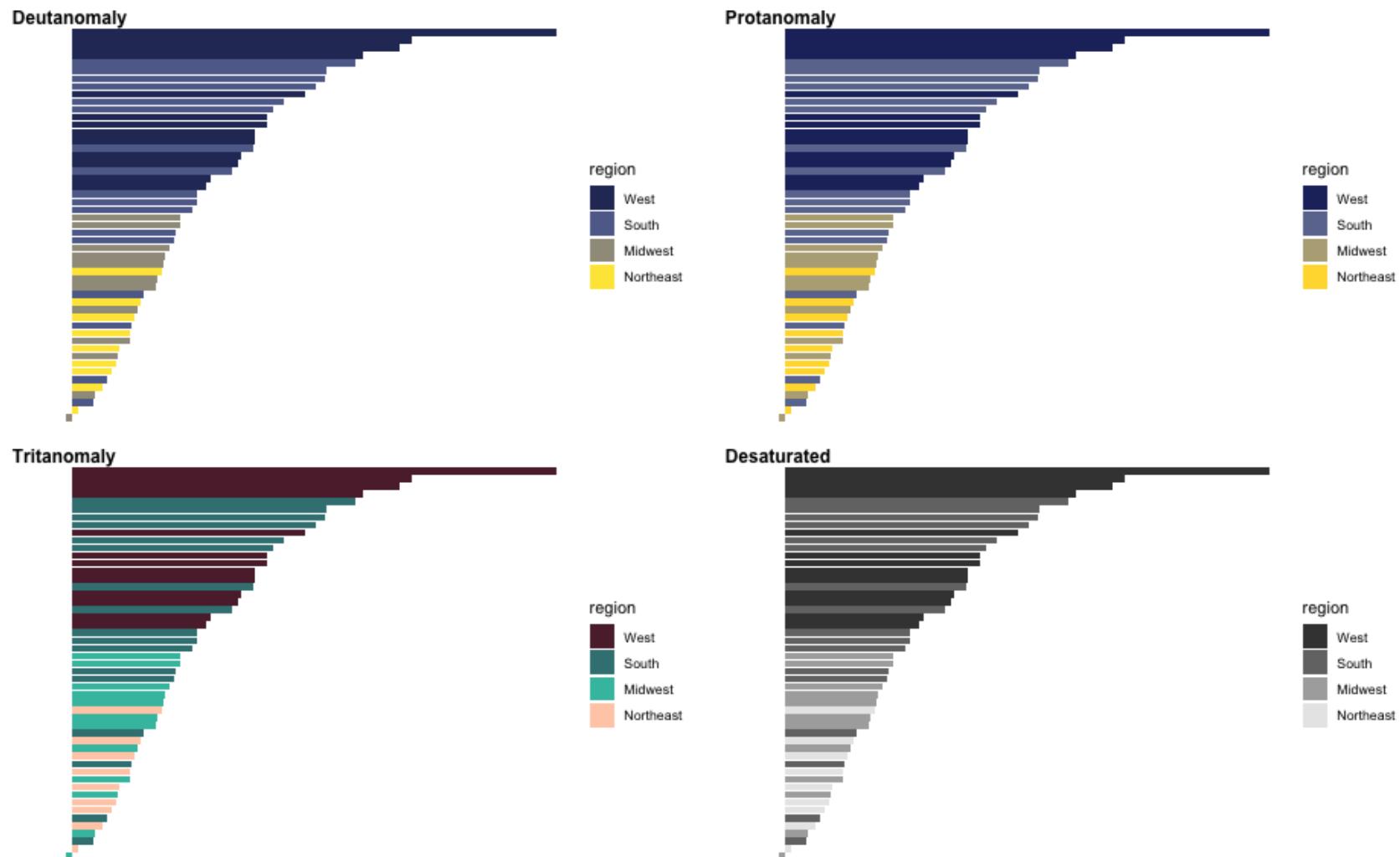


Alternative: viridis

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



Revised version

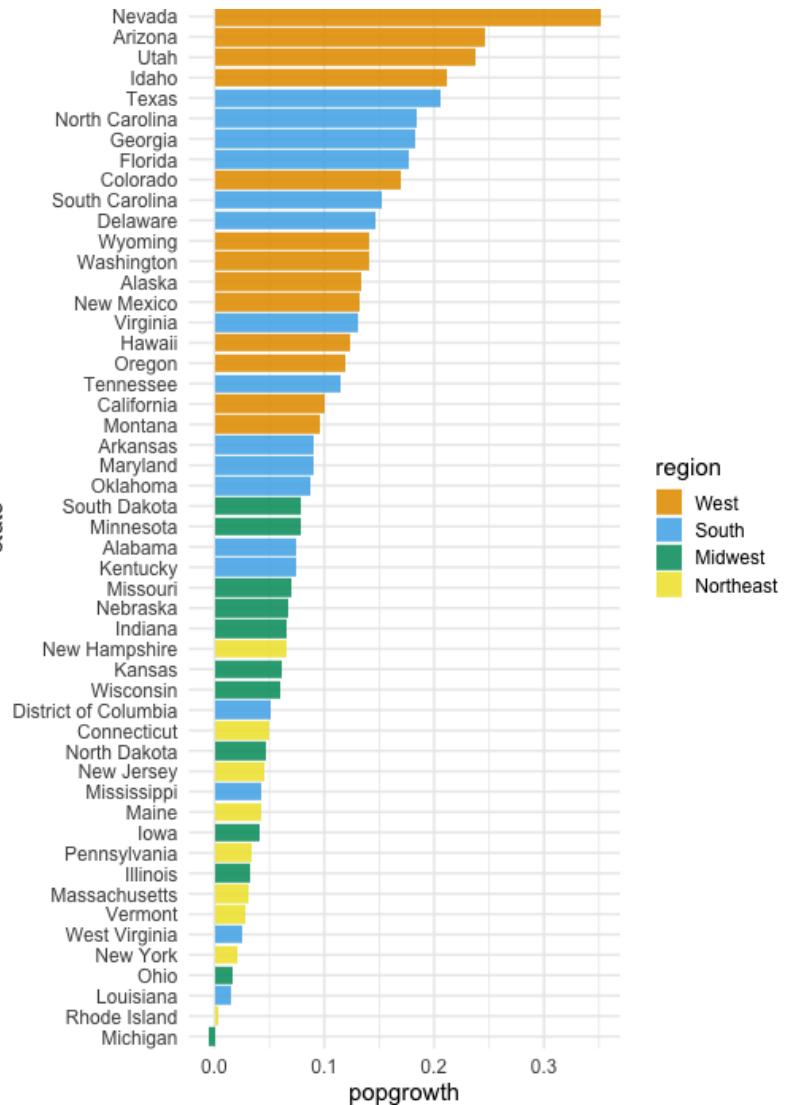


The Okabe Ito palette

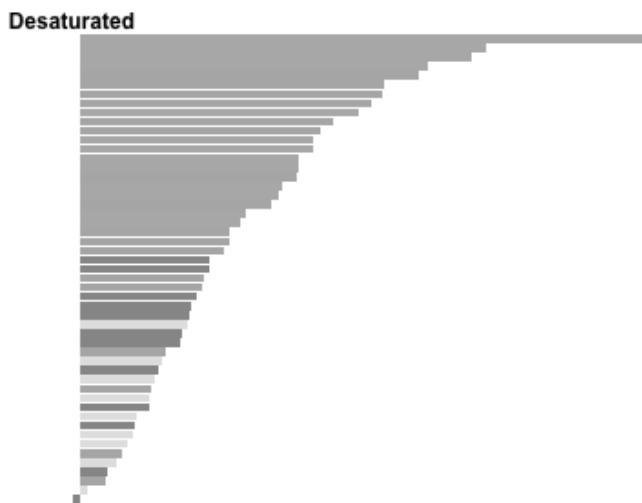
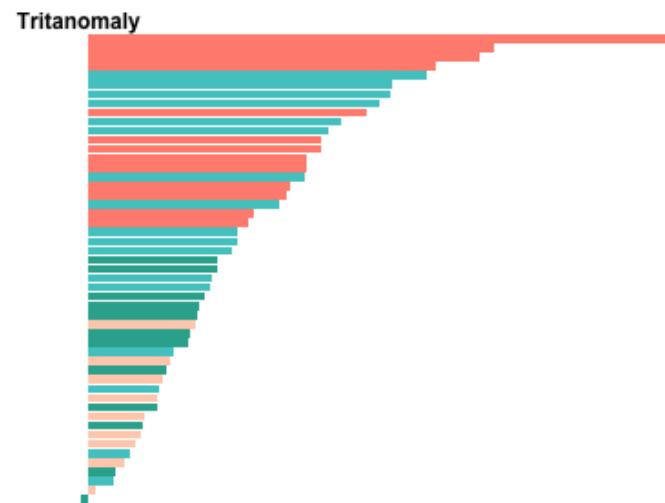
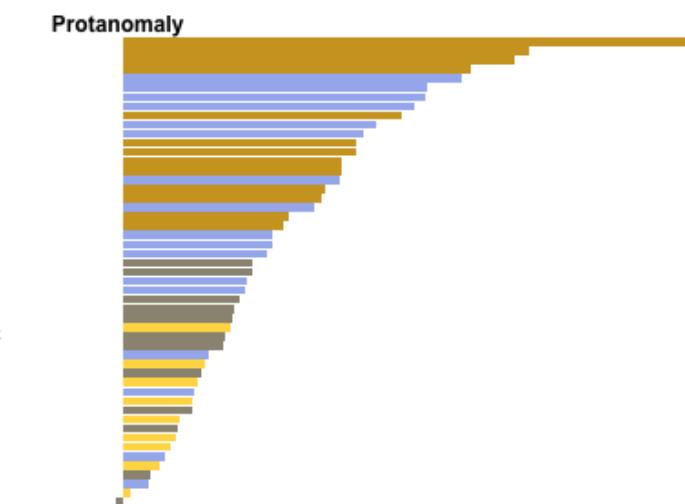
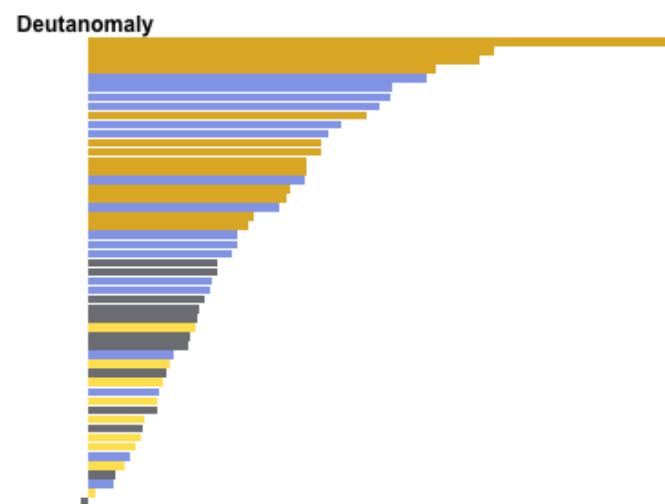
- From Color Universal Design

```
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?

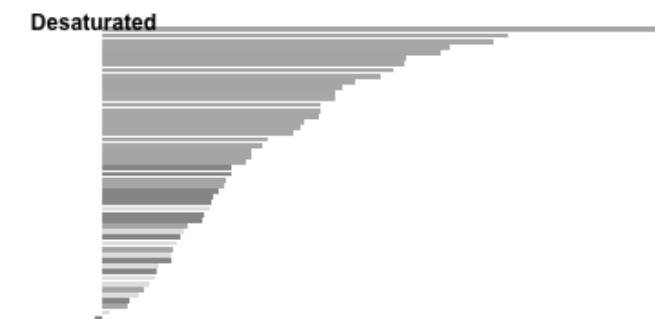
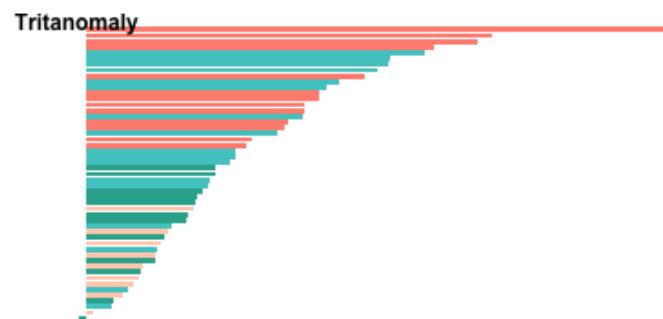
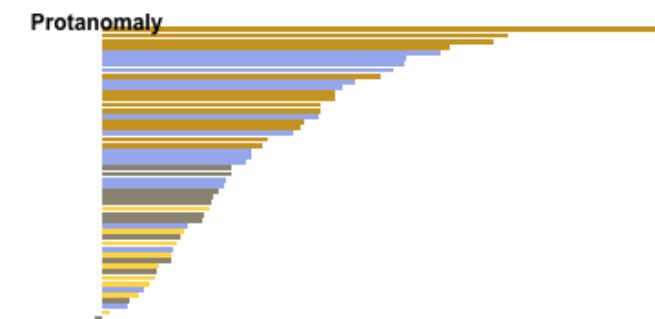
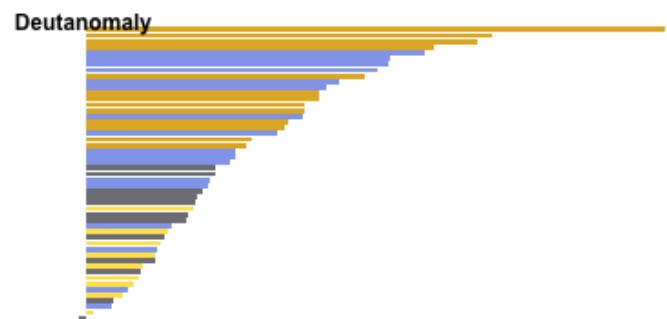
- Also part of the **{colorblindr}** package (here)
 - depends on the dev versions of **{colorspace}** and **{cowplot}**, which are useful packages in their own right

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

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

```
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)
```



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

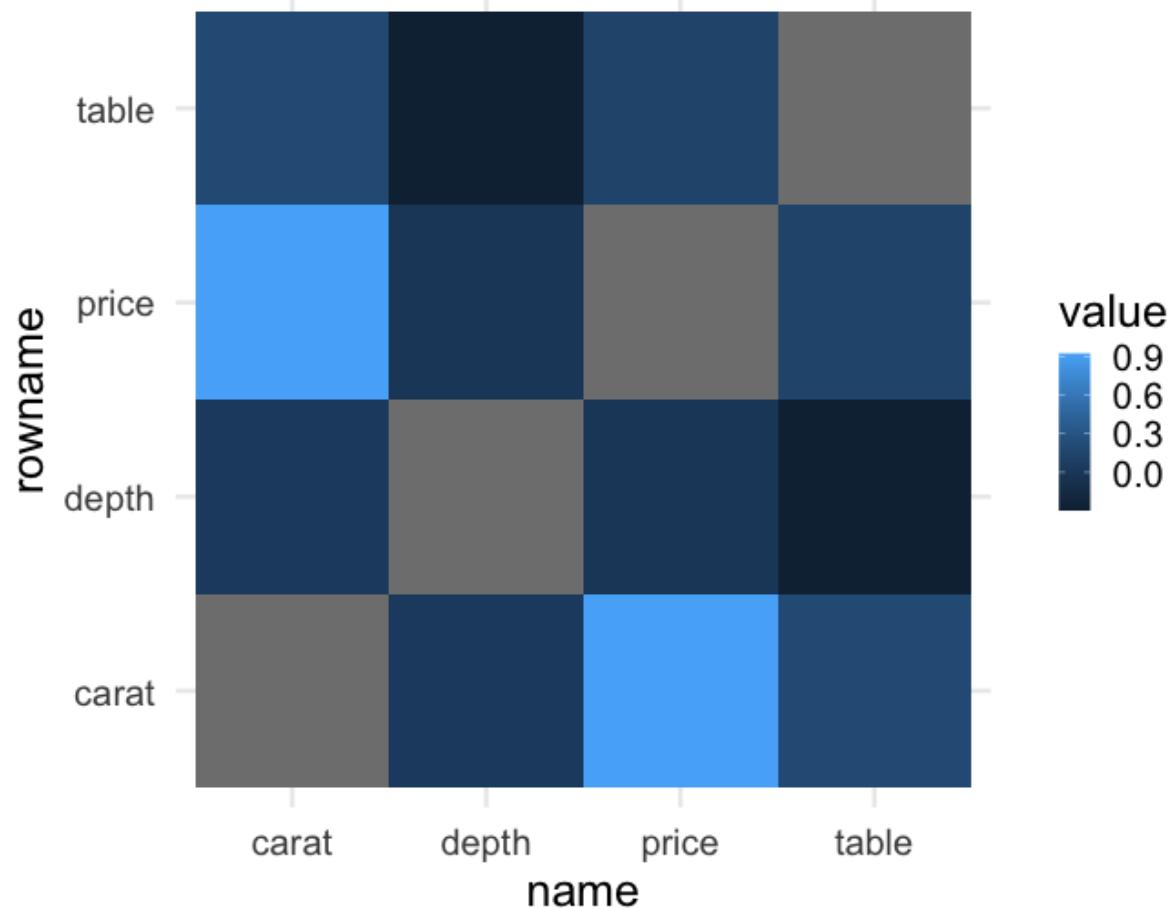
Heatmaps

First the data:

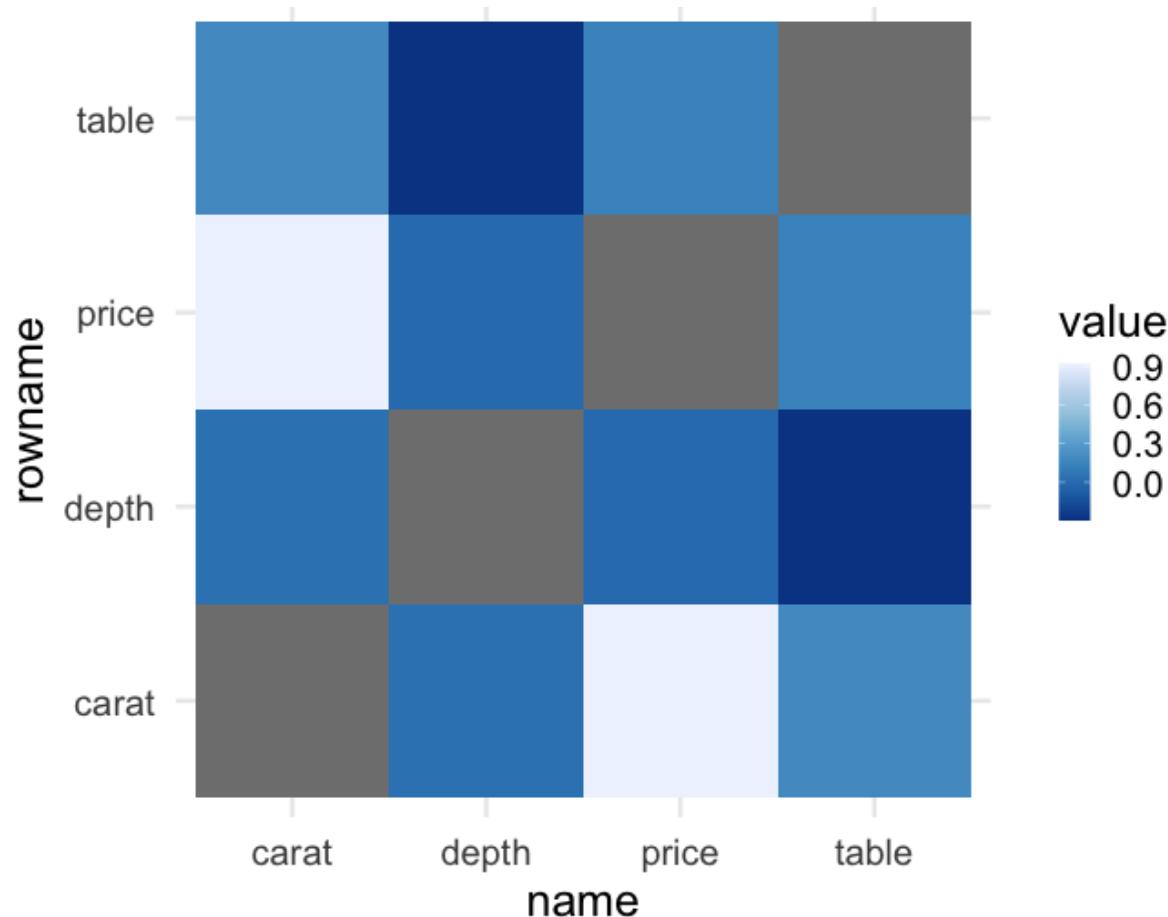
```
hm <- diamonds %>%
  select(table, price, depth, carat) %>%
  corrr::correlate() %>%
  pivot_longer(-rowname) %>%
  mutate(name = fct_reorder(name, value),
        rowname = fct_reorder(rowname, value))
hm
```

```
## # A tibble: 16 x 3
##   rowname name      value
##   <fct>   <fct>    <dbl>
## 1 table   table    NA
## 2 table   price    0.1271339
## 3 table   depth   -0.2957785
## 4 table   carat    0.1816175
## 5 price   table    0.1271339
## 6 price   price    NA
## # ... with 10 more rows
```

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



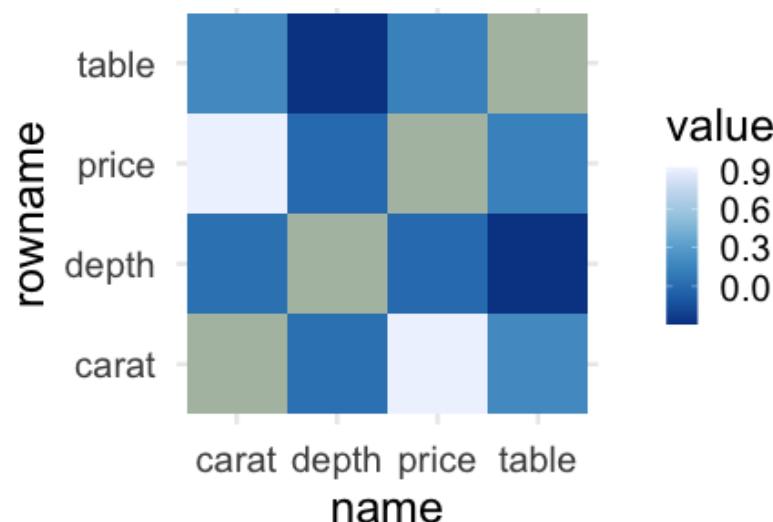
```
ggplot(hm, aes(name, rowname)) +  
  geom_tile(aes(fill = value)) +  
  coord_fixed() +  
  scale_fill_distiller(palette = "Blues")
```



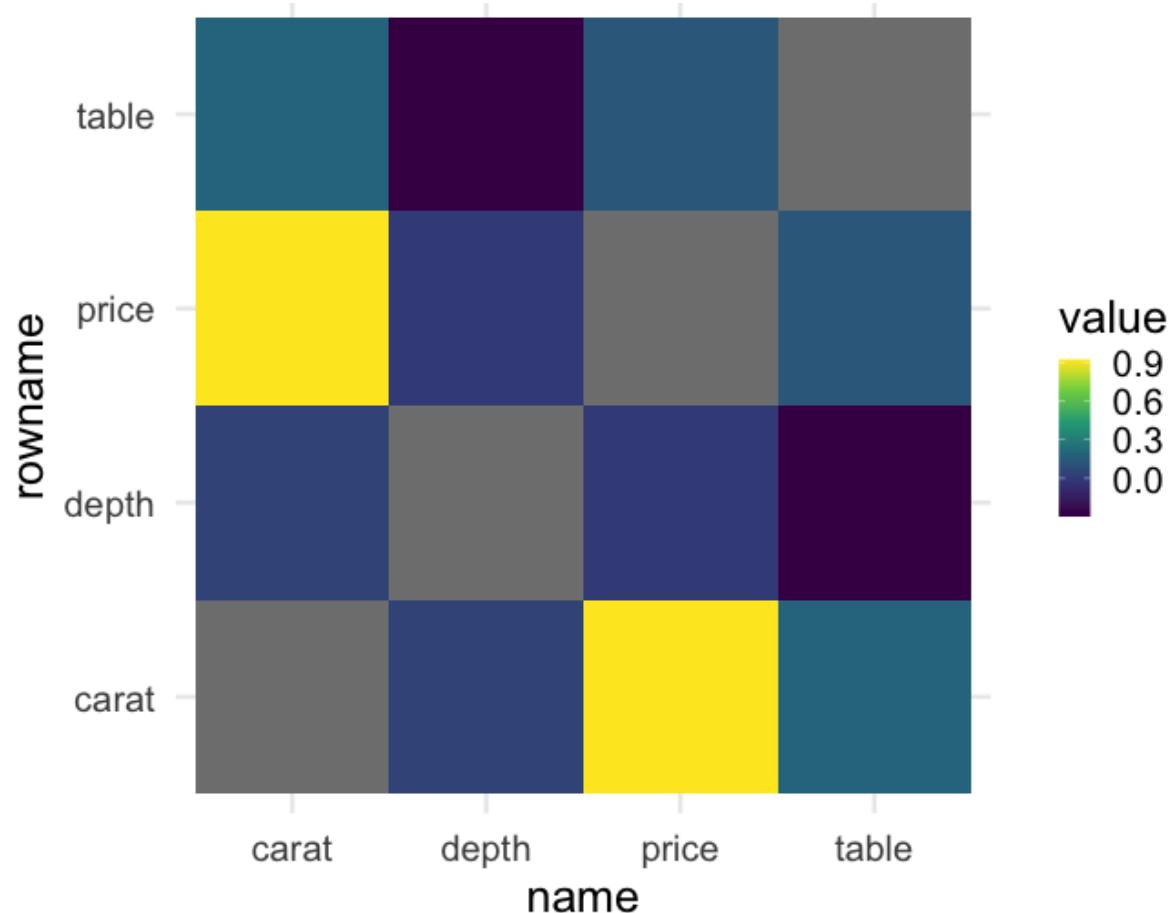
Change the NA value

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

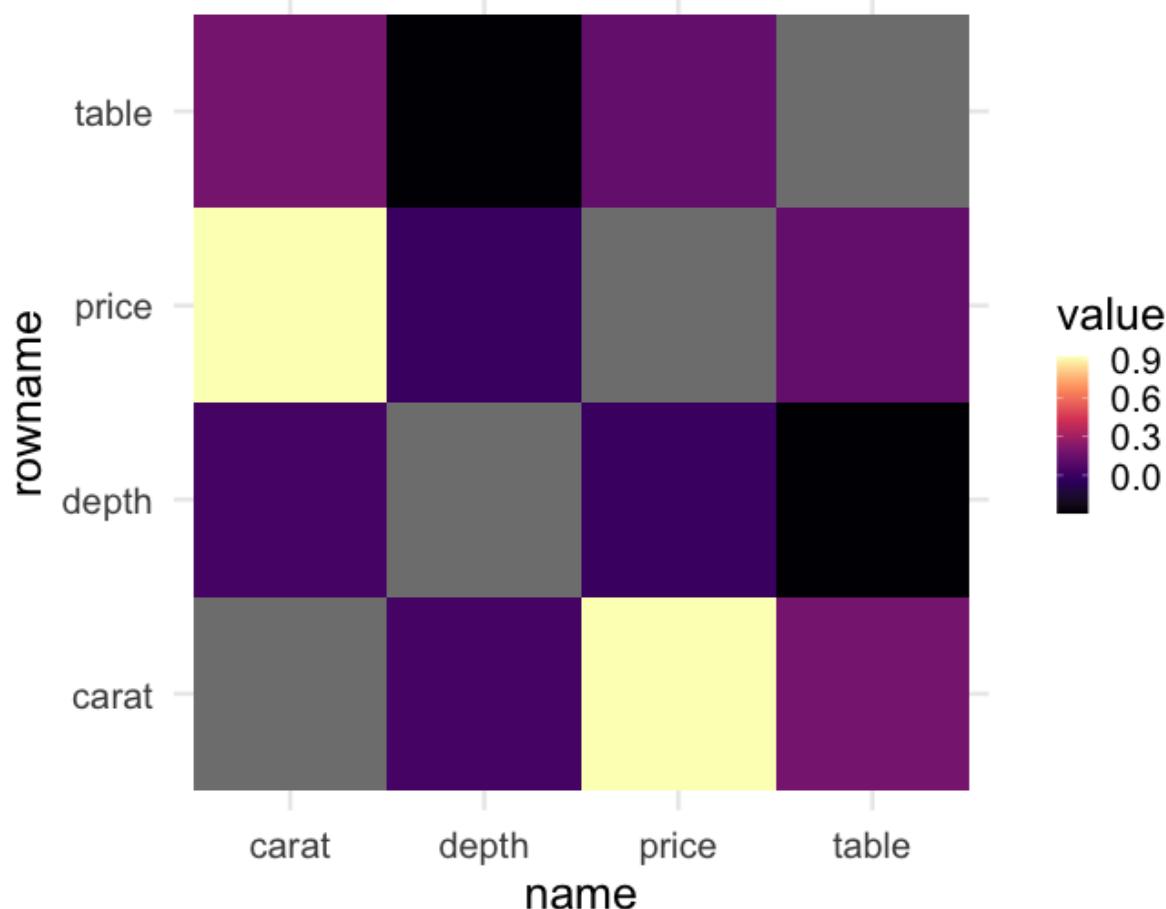
```
ggplot(hm, aes(name, rowname)) +  
  geom_tile(aes(fill = value)) +  
  coord_fixed() +  
  scale_fill_distiller(palette = "Blues",  
                      na.value = "#b0bfb0")
```



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



```
ggplot(hm, aes(name, rowname)) +  
  geom_tile(aes(fill = value)) +  
  coord_fixed() +  
  scale_fill_viridis_c(option = "magma")
```

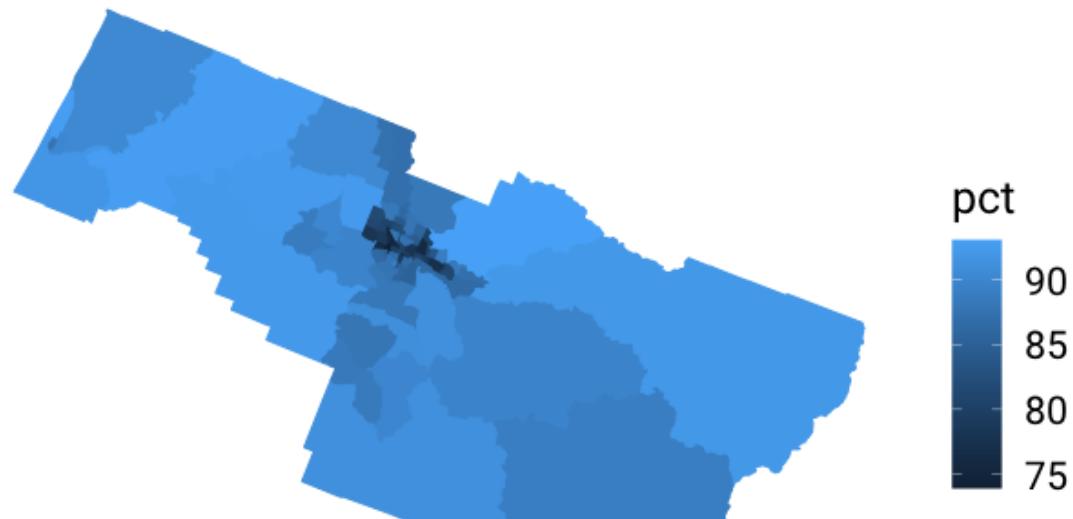


```
option = c("viridis", "magma", "inferno", "plasma")
```

Choropleths

Percentage of people identifying as White

Lane County

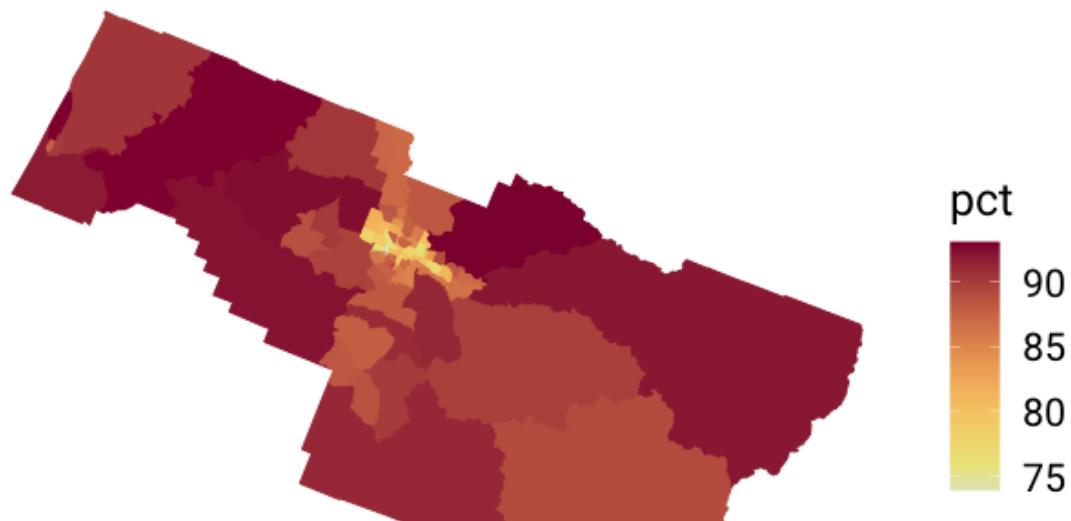


US Census Decennial Tract Data

Heat palette

Percentage of people identifying as White

Lane County



US Census Decennial Tract Data

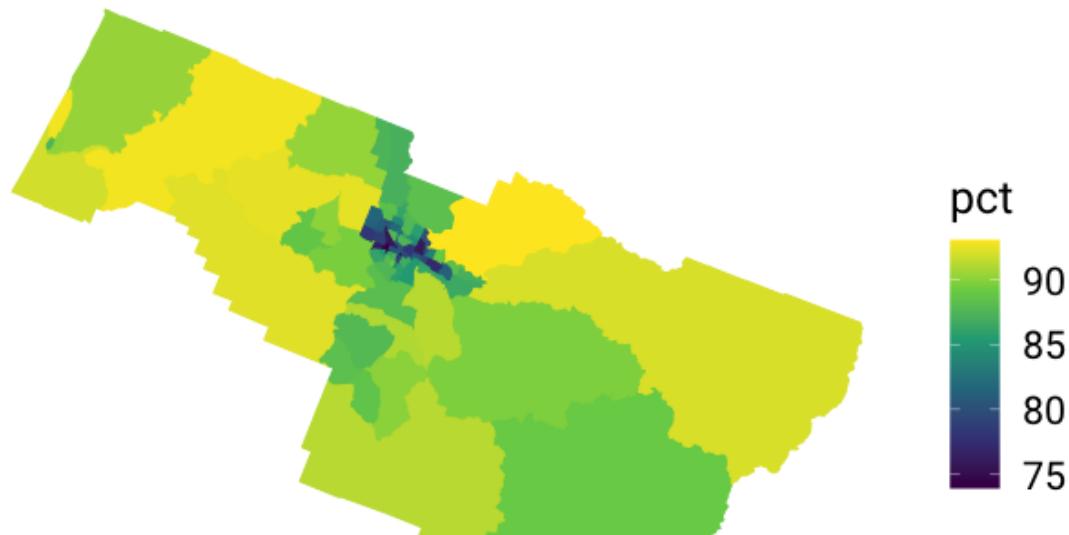
Options

- `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



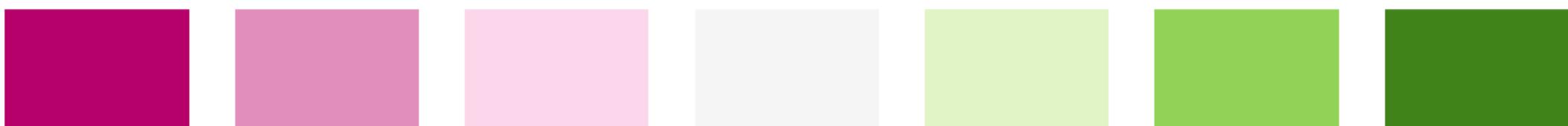
US Census Decennial Tract Data

Diverging palettes

CARTO Earth



ColorBrewer PiYG



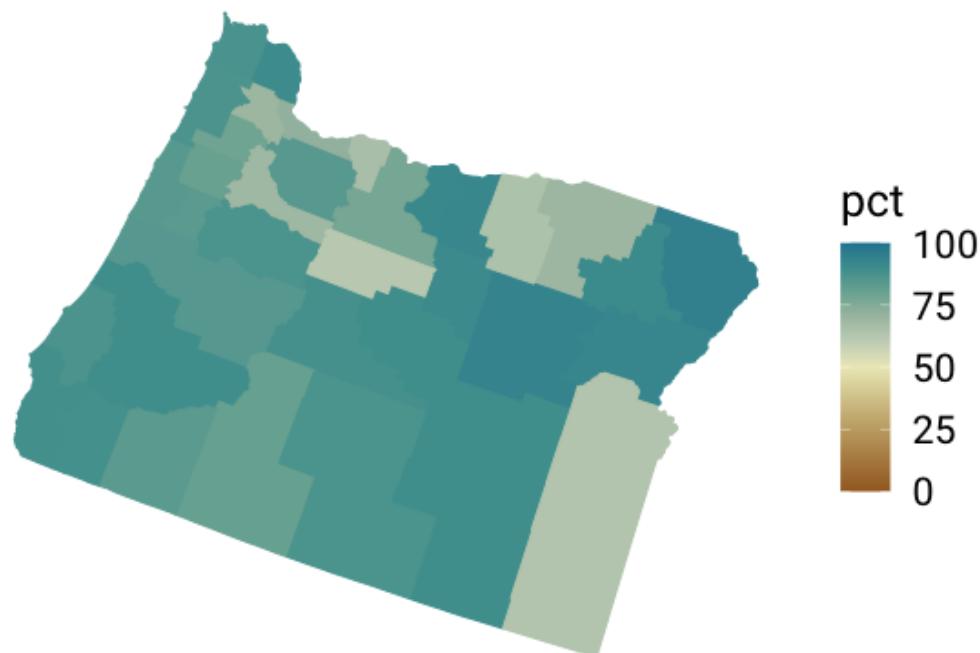
Blue-Red



Earth palette

Percentage of people identifying as White

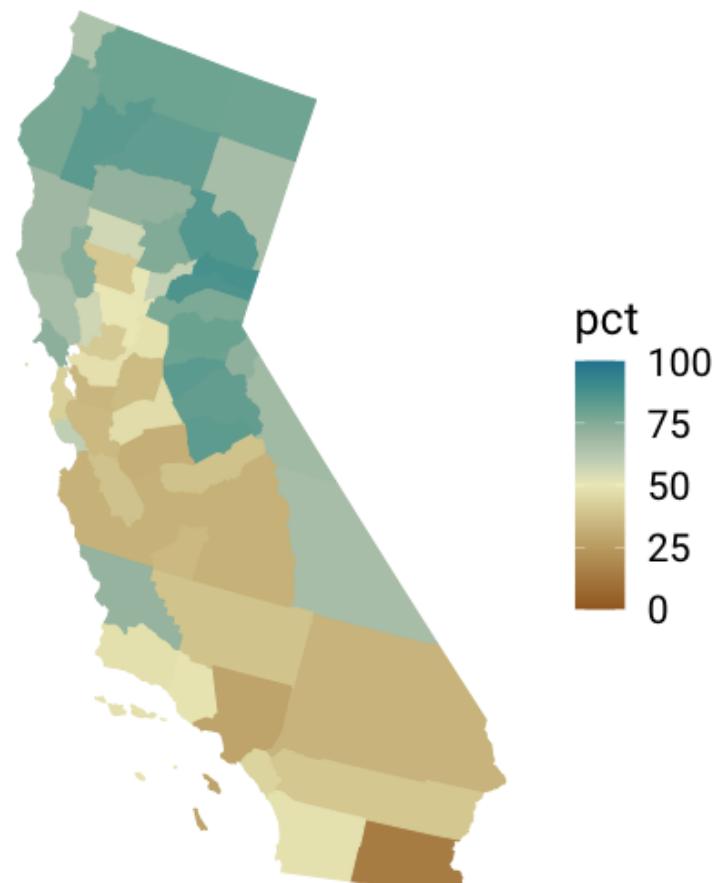
Oregon



US Census Decennial Tract Data

Percentage of people identifying as White

California



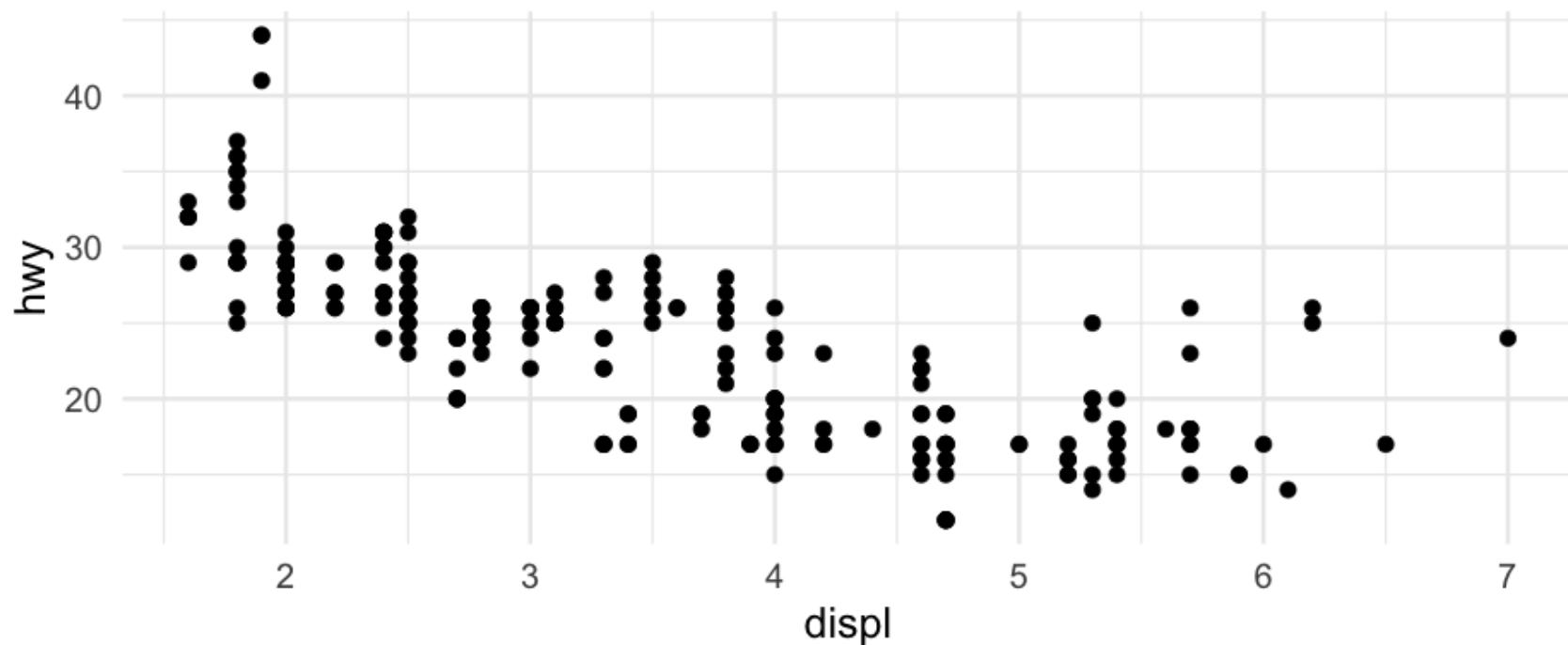
US Census Decennial County Data

Color as a tool
to highlight

MPG data

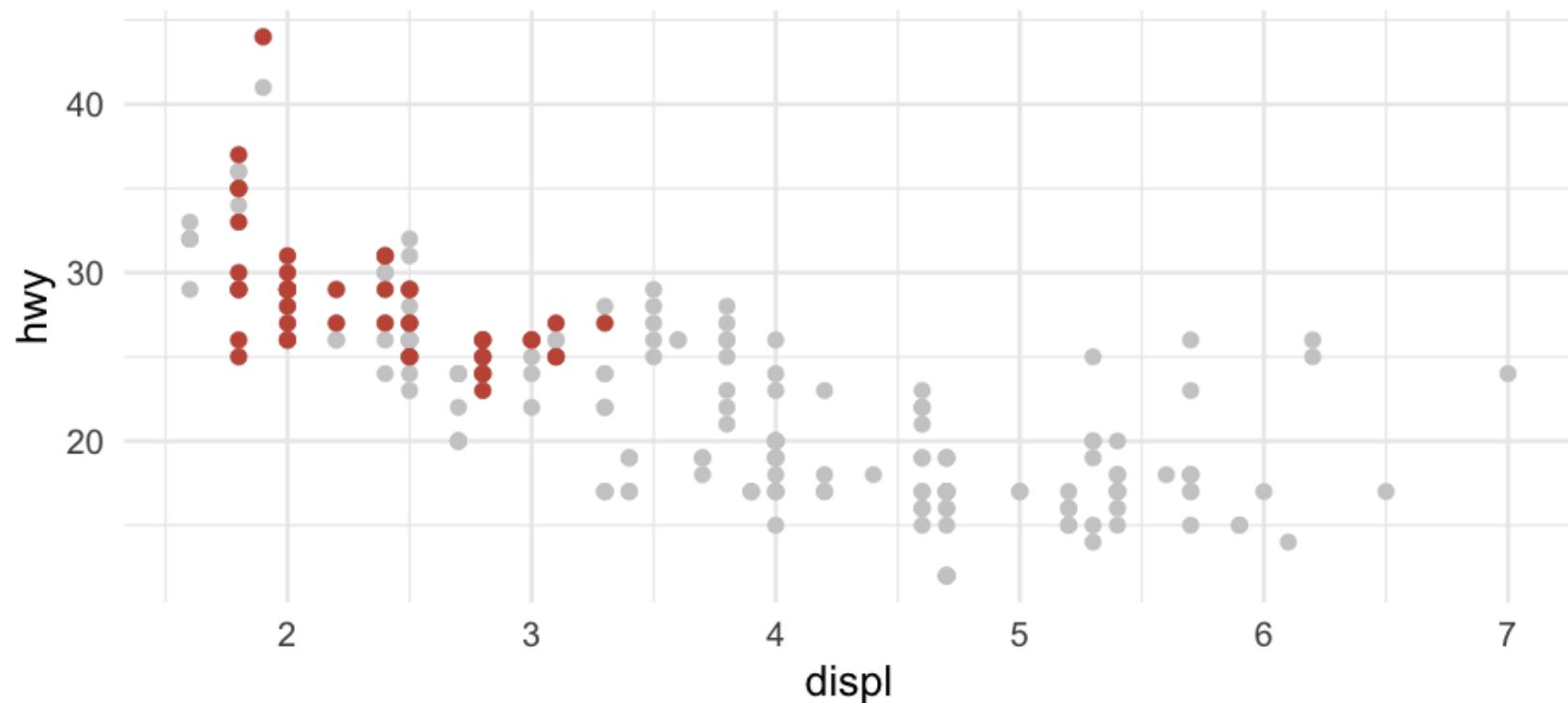
Basic scatterplot of weight to highway mpg

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



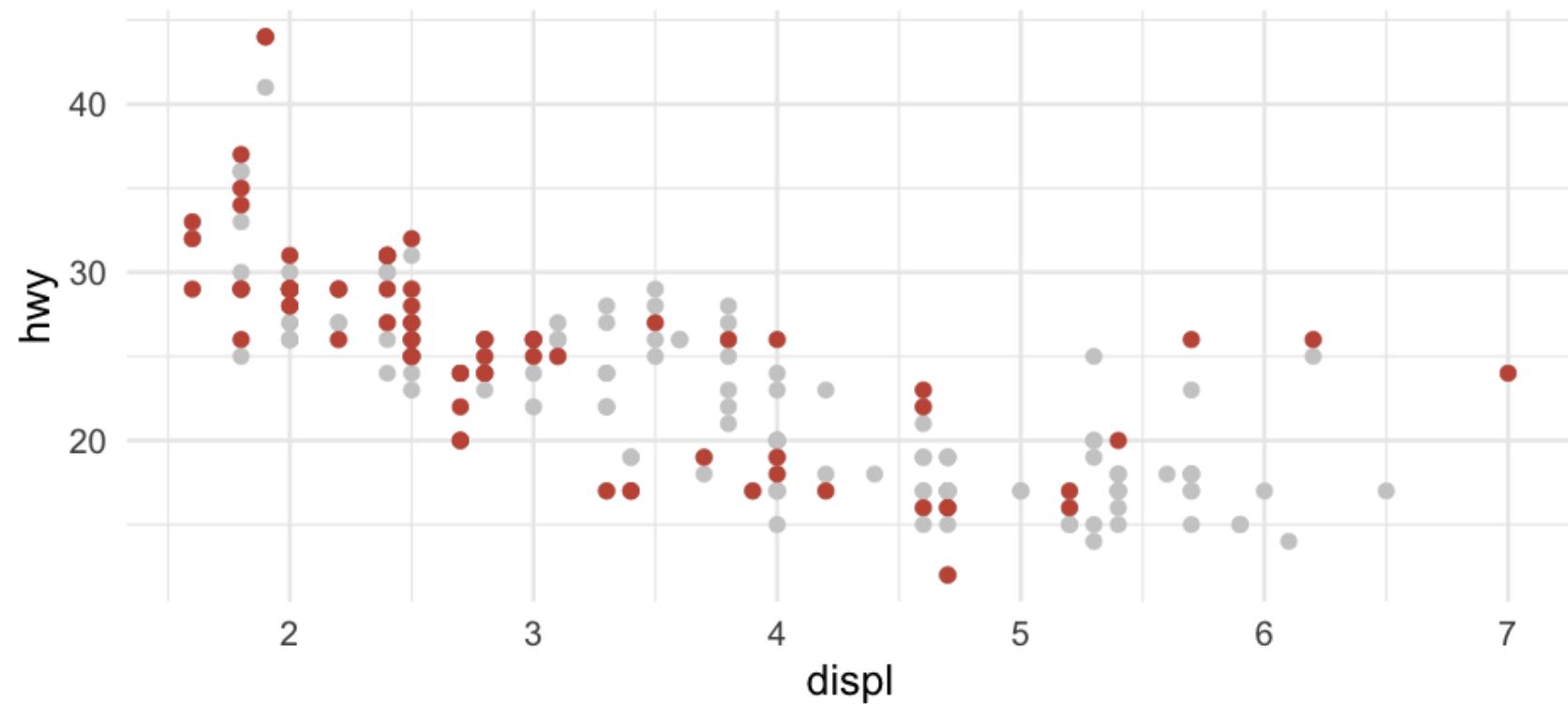
Highlight compact cars

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



Highlight manual cars

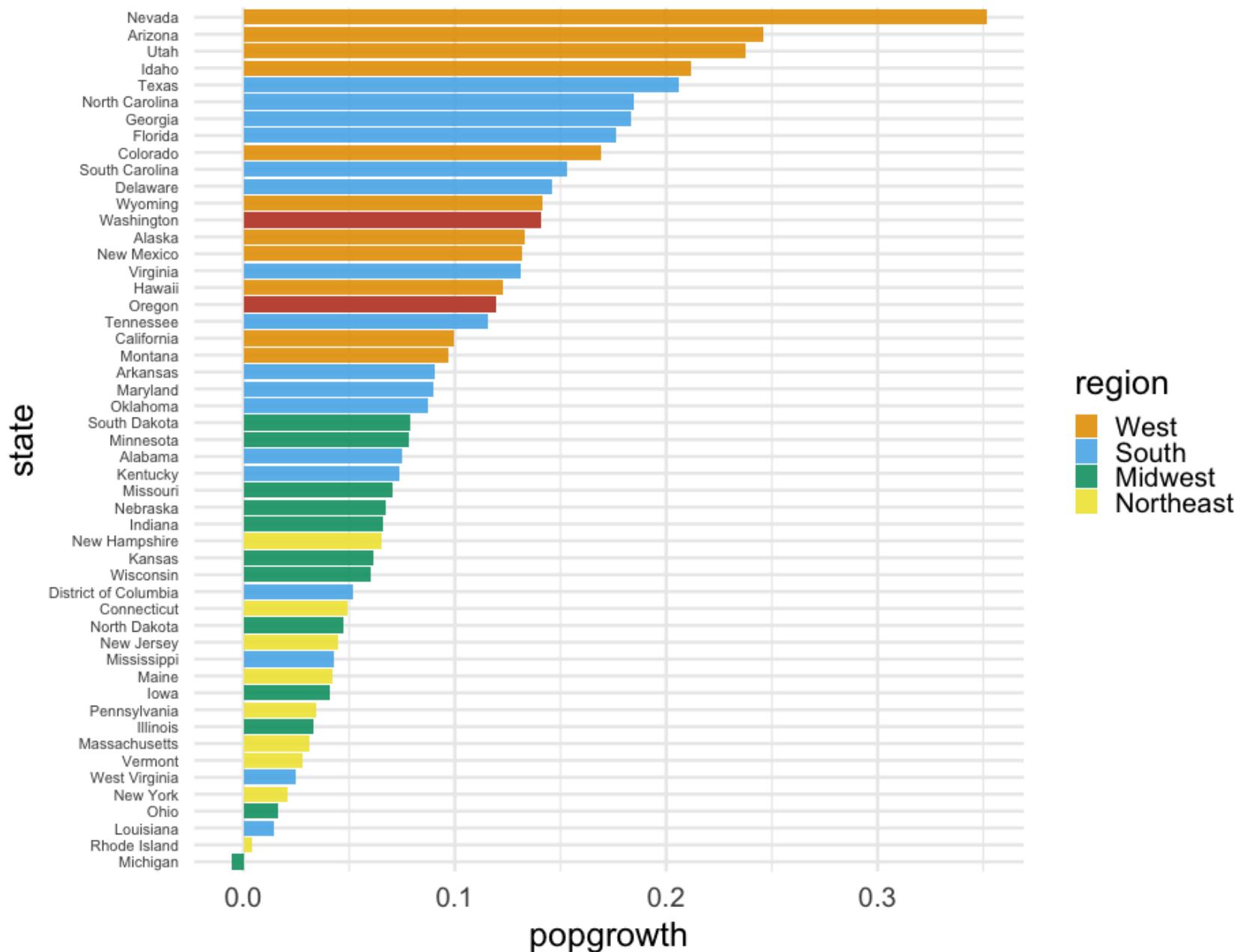
```
ggplot(mpg, aes(displ, hwy)) +  
  geom_point(color = "gray80") +  
  geom_point(data = filter(mpg, str_detect(trans, "manual")),  
             color = "#C55644")
```



Back to our states plot

Highlight Oregon and Washington

```
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 = "#C55644") +  
  scale_fill_OkabeIto()
```



Color labels

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

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

label_color
```

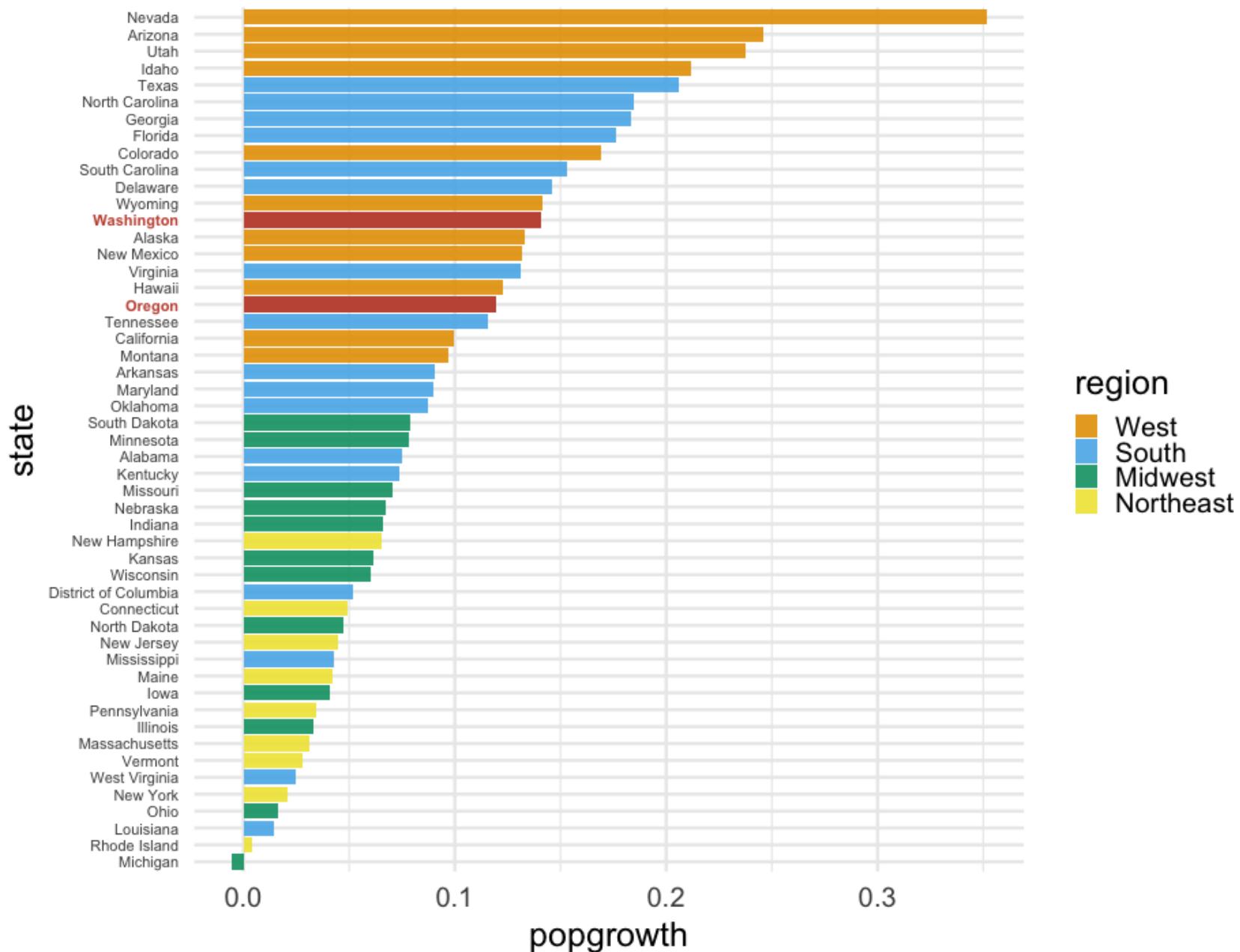
```
## [1] "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"
## [8] "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"
## [15] "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"
## [22] "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"
## [29] "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "#C55644" "gray30"
## [36] "gray30"  "gray30"  "gray30"  "#C55644" "gray30"  "gray30"  "gray30"  "gray30"
## [43] "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"  "gray30"
## [50] "gray30"  "gray30"
```

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

label_face
```

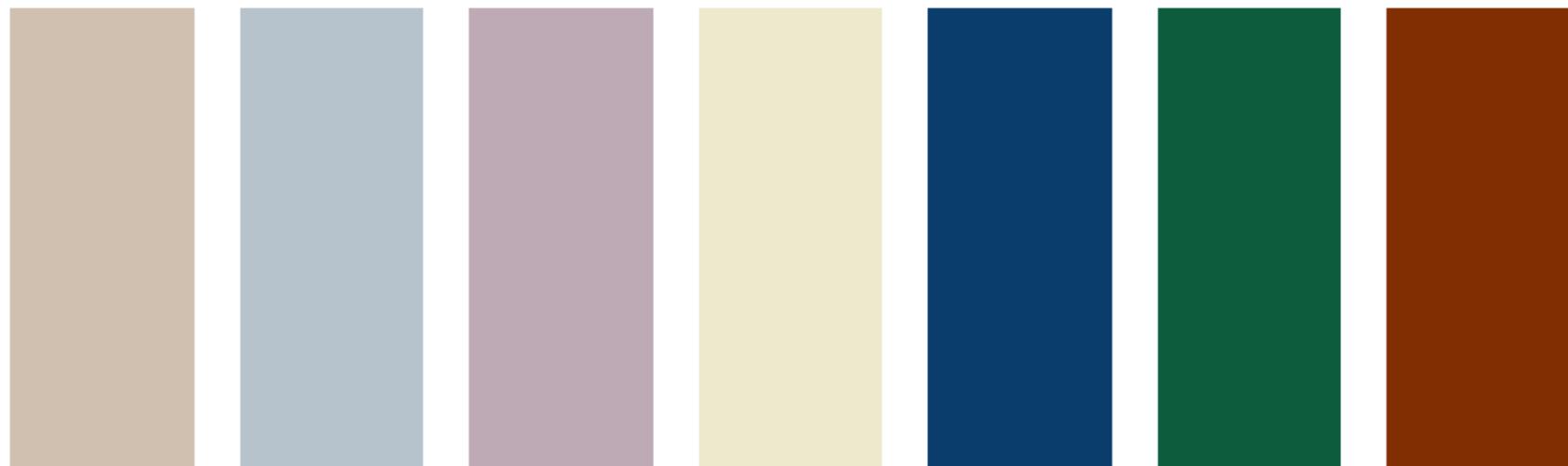
```
## [1] "plain"  "plain"  "plain"  "plain"  "plain"  "plain"  "plain"  "plain"  "plain"
## [10] "plain"  "plain"  "plain"  "plain"  "plain"  "plain"  "plain"  "plain"  "plain"
## [19] "plain"  "plain"  "plain"  "plain"  "plain"  "plain"  "plain"  "plain"  "plain"
## [28] "plain"  "plain"  "plain"  "plain"  "plain"  "plain"  "bold"   "plain"  "plain"
## [37] "plain"  "plain"  "bold"   "plain"  "plain"  "plain"  "plain"  "plain"  "plain"
## [46] "plain"  "plain"  "plain"  "plain"  "plain"  "plain"  "plain"  "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 = "#C55644") +  
  scale_fill_OkabeIto() +  
  theme(axis.text.y = element_text(color = label_color,  
                                    face = label_face))
```

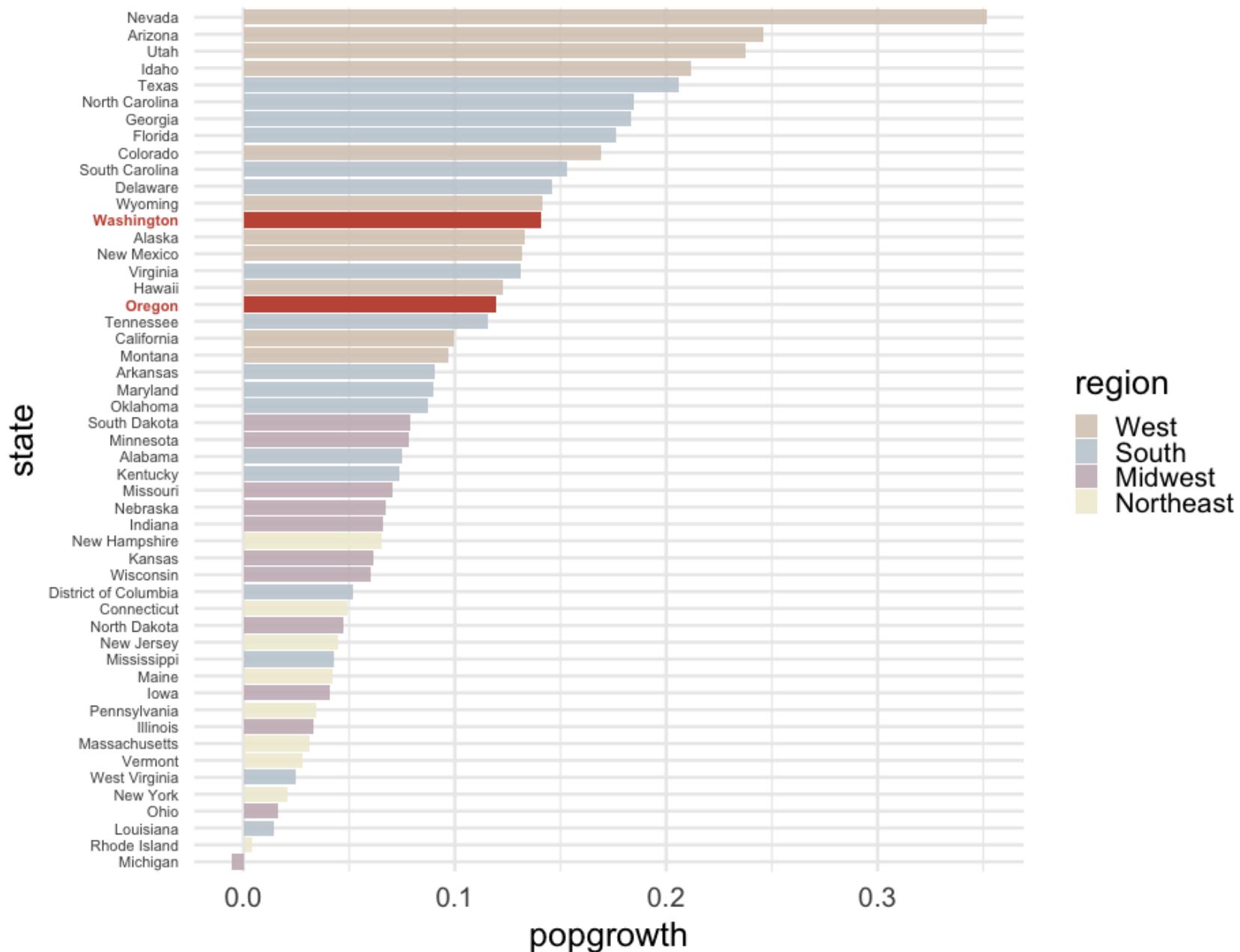


Even better

```
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 = "#C55644") +  
  scale_fill_manual(values = accent_OkabeIto) +  
  theme(axis.text.y = element_text(color = label_color,  
                                    face = label_face))
```

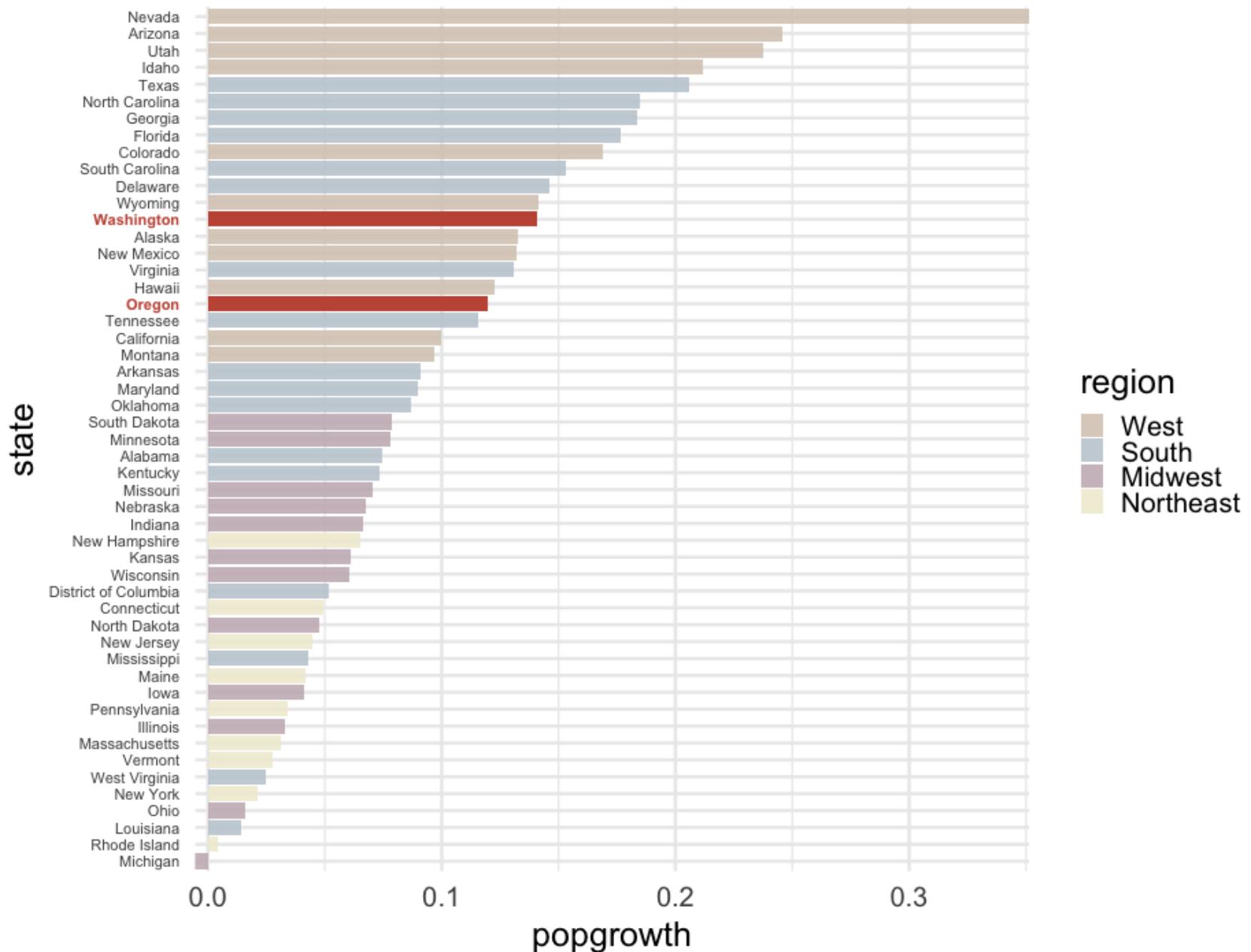


Or even better

```
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 = "#C55644") +
  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
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



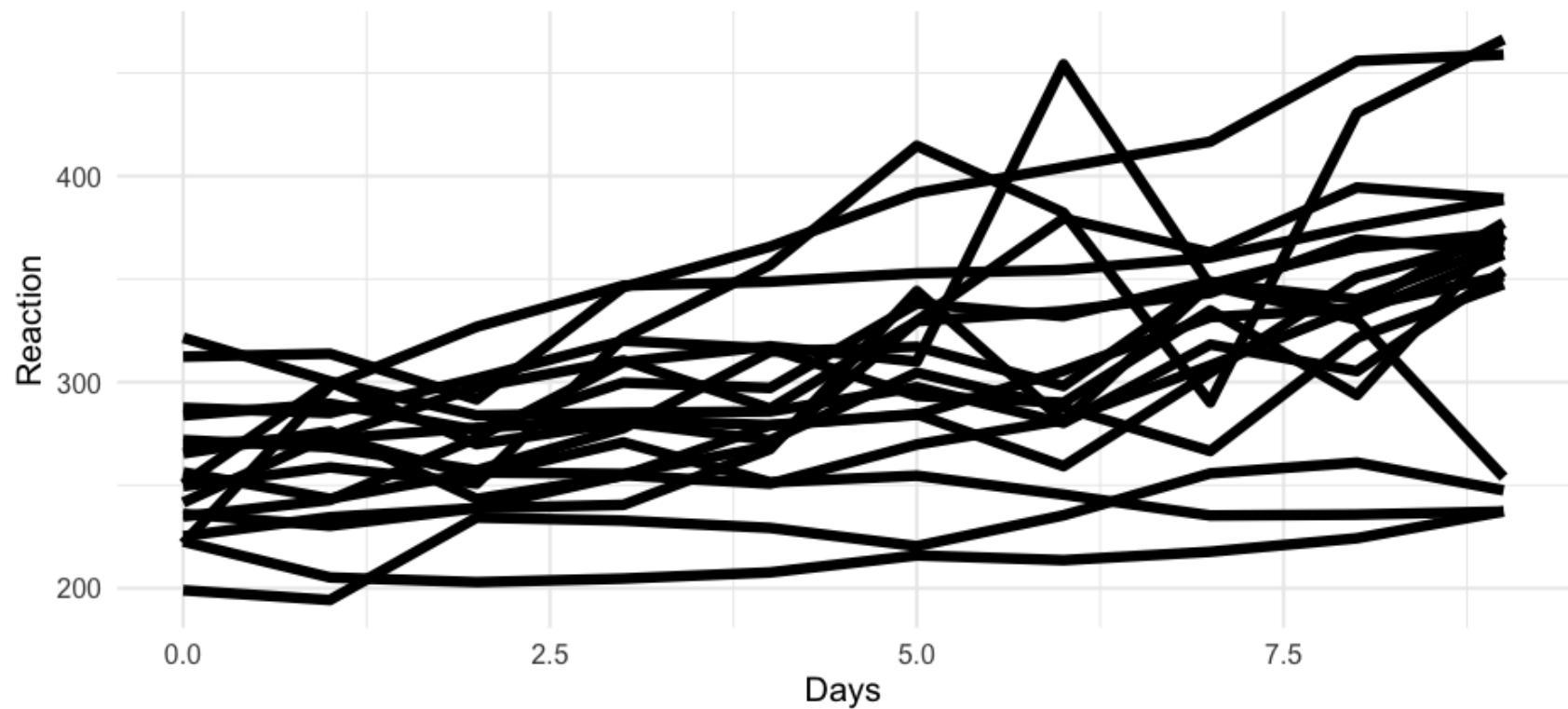
Last example

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

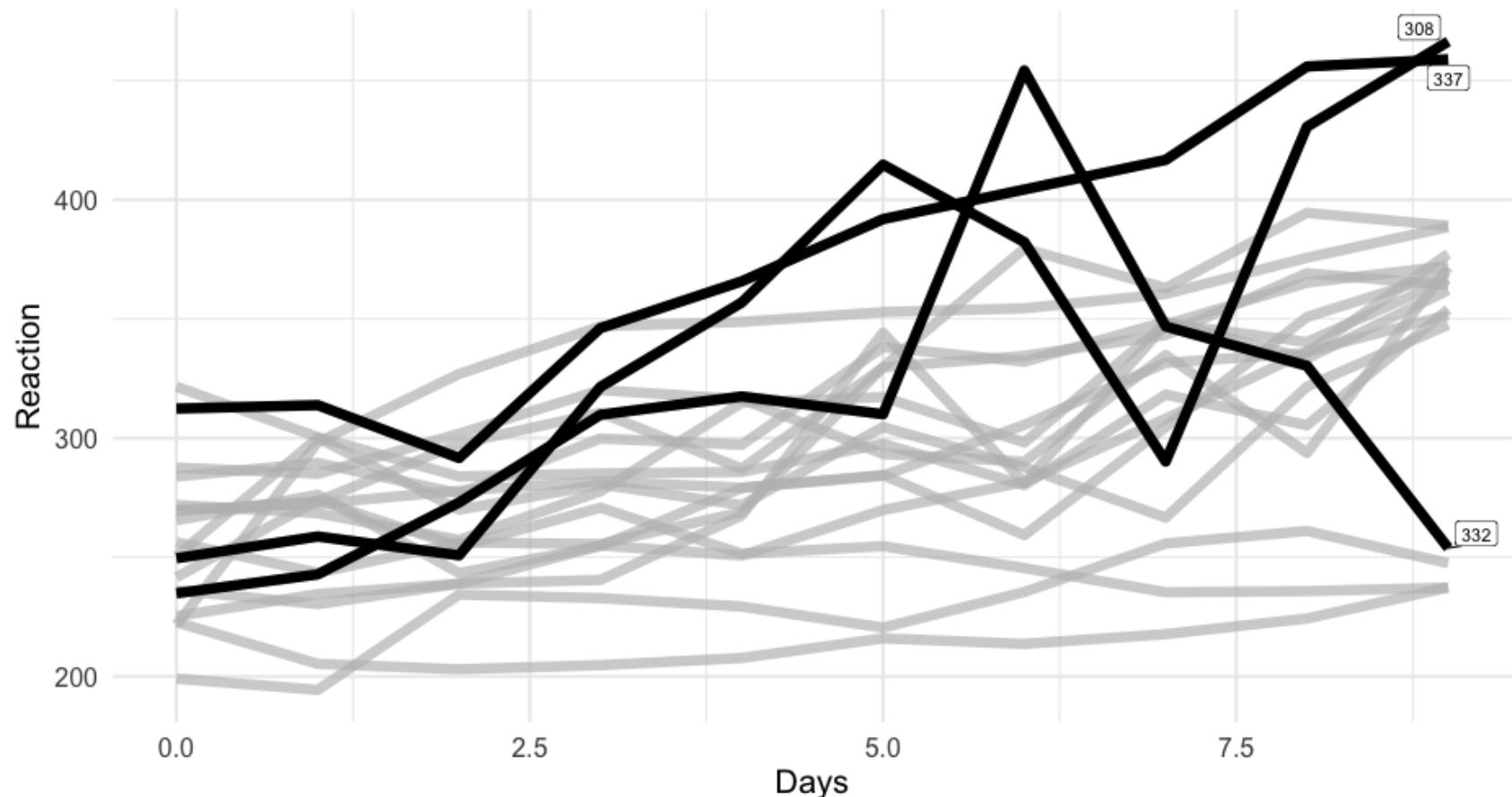
```
##   Reaction Days Subject
## 1      250     0    308
## 2      259     1    308
## 3      251     2    308
## 4      321     3    308
## 5      357     4    308
## 6      415     5    308
```

Plot by subject

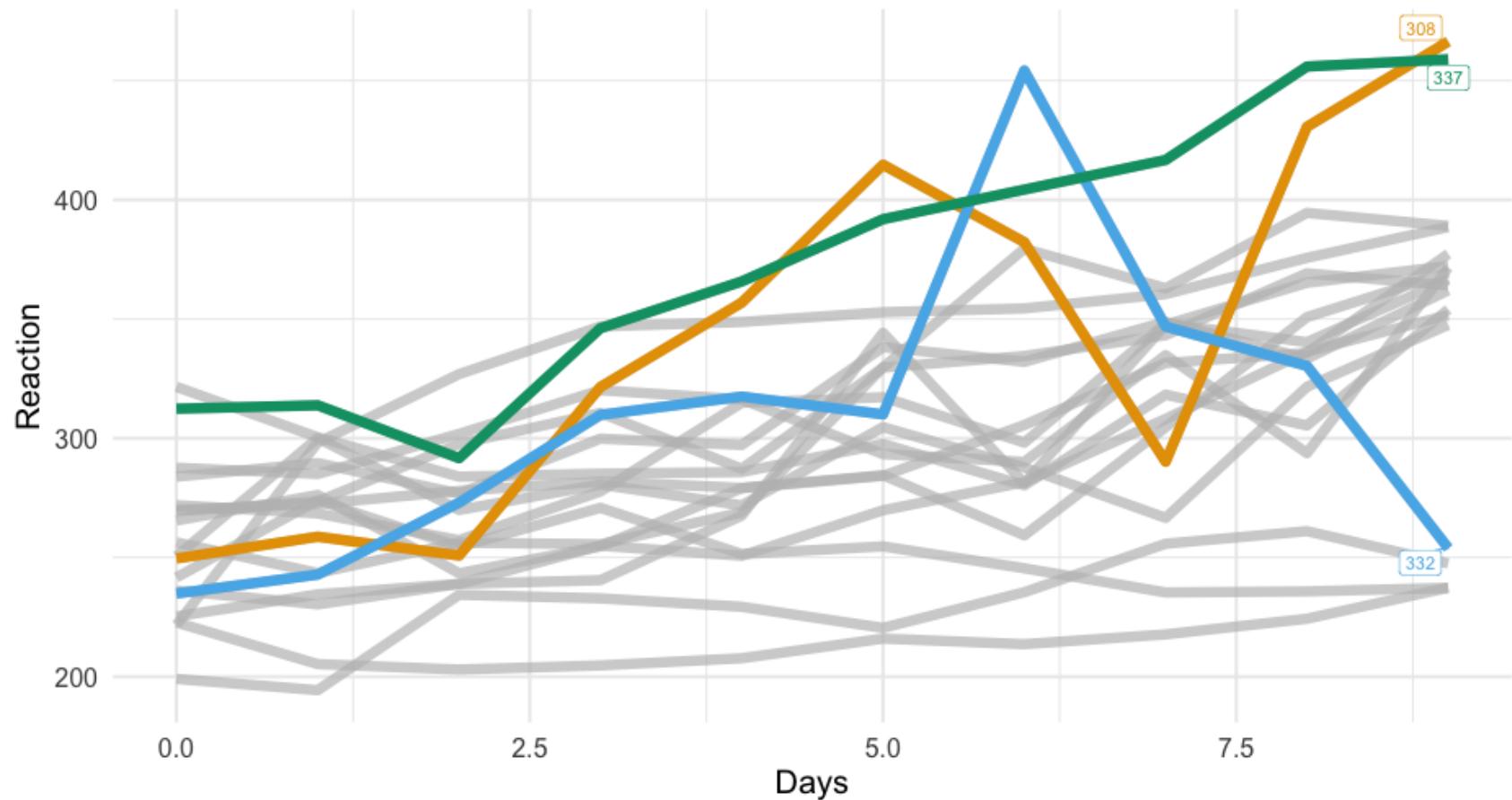
```
ggplot(sleepstudy, aes(Days, Reaction, group = Subject)) +  
  geom_line()
```



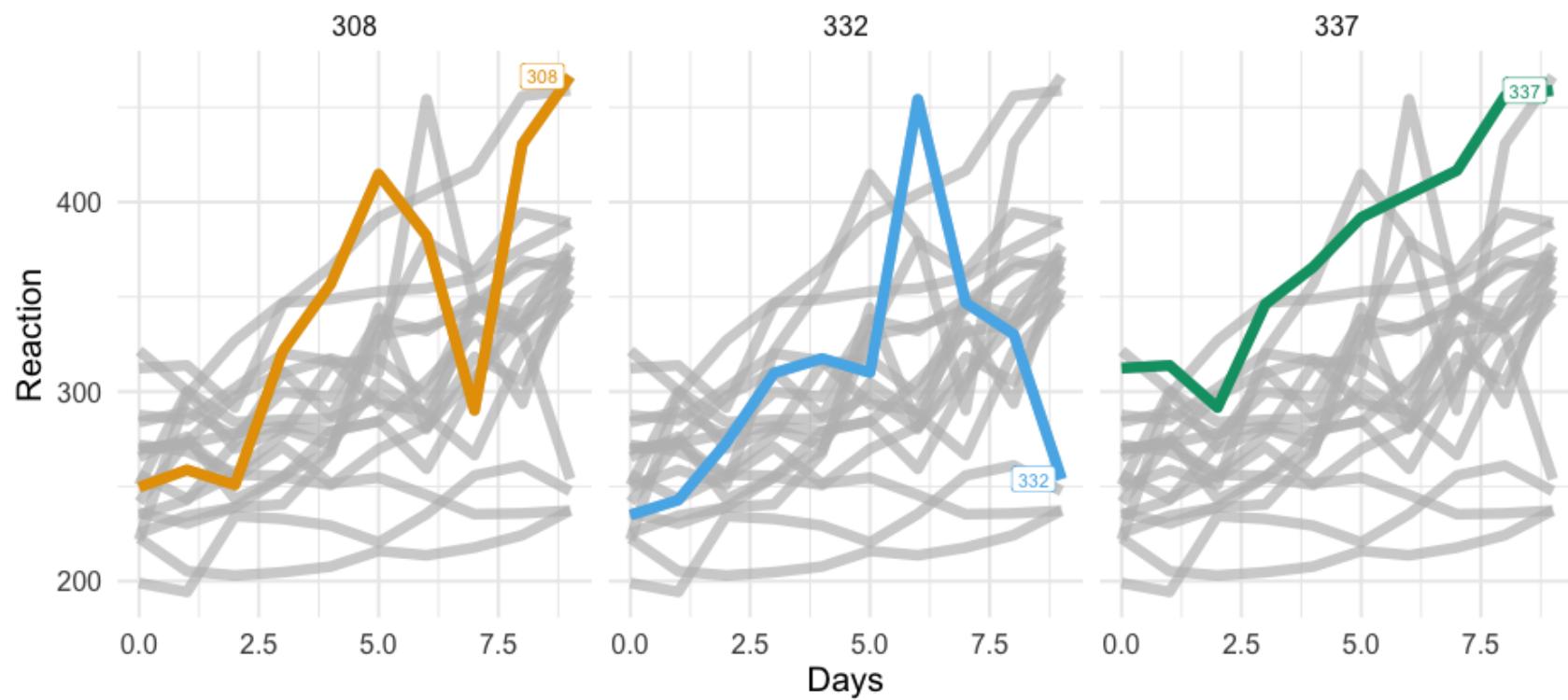
```
library(gghighlight)
ggplot(sleepstudy, aes(Days, Reaction, group = Subject)) +
  geom_line() +
  gghighlight(max(Reaction) > 400)
```



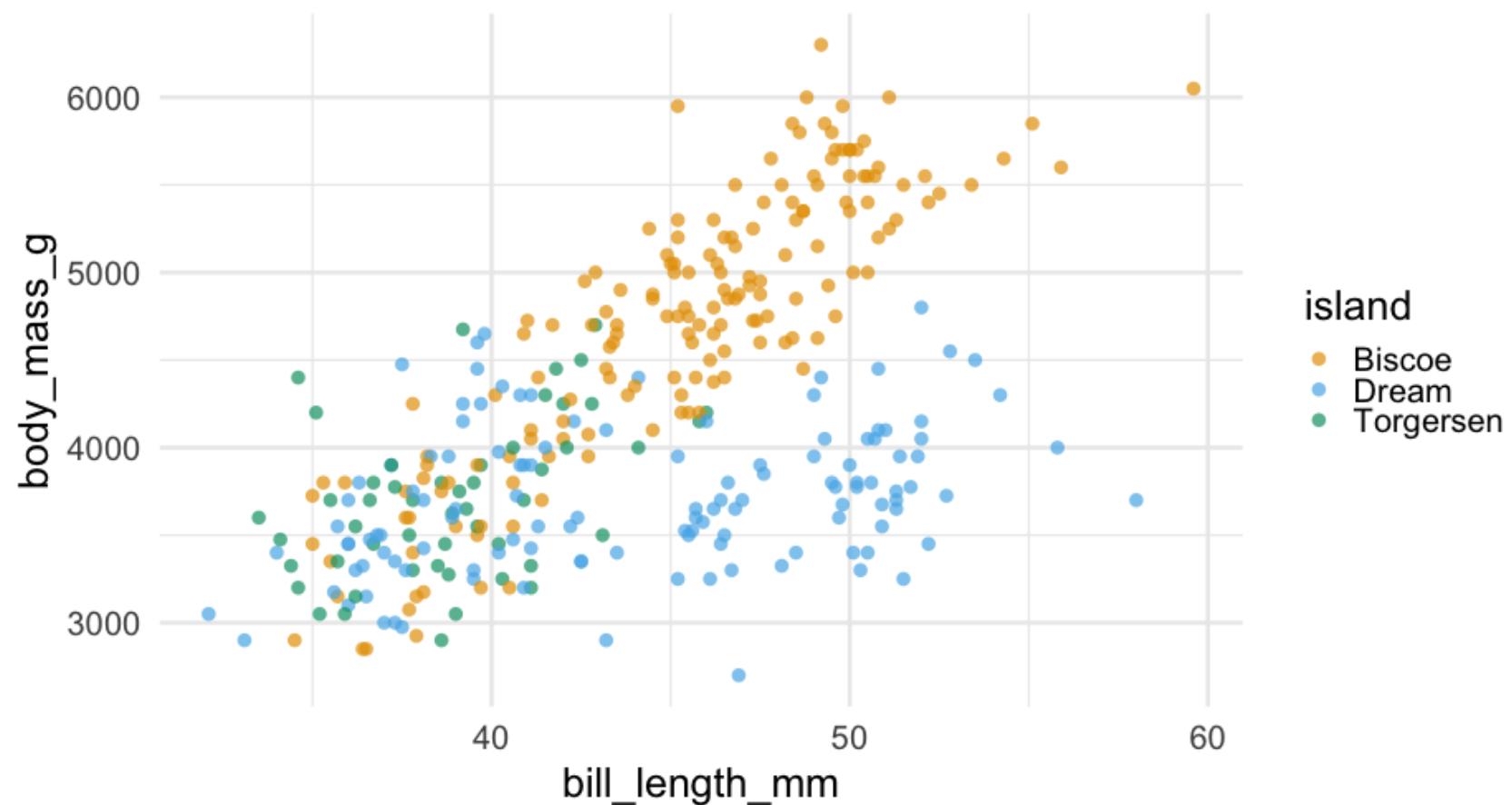
```
library(gghighlight)
ggplot(sleepstudy, aes(Days, Reaction, color = Subject)) +
  geom_line() +
  gghighlight(max(Reaction) > 400) +
  scale_color_OkabeIto()
```



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

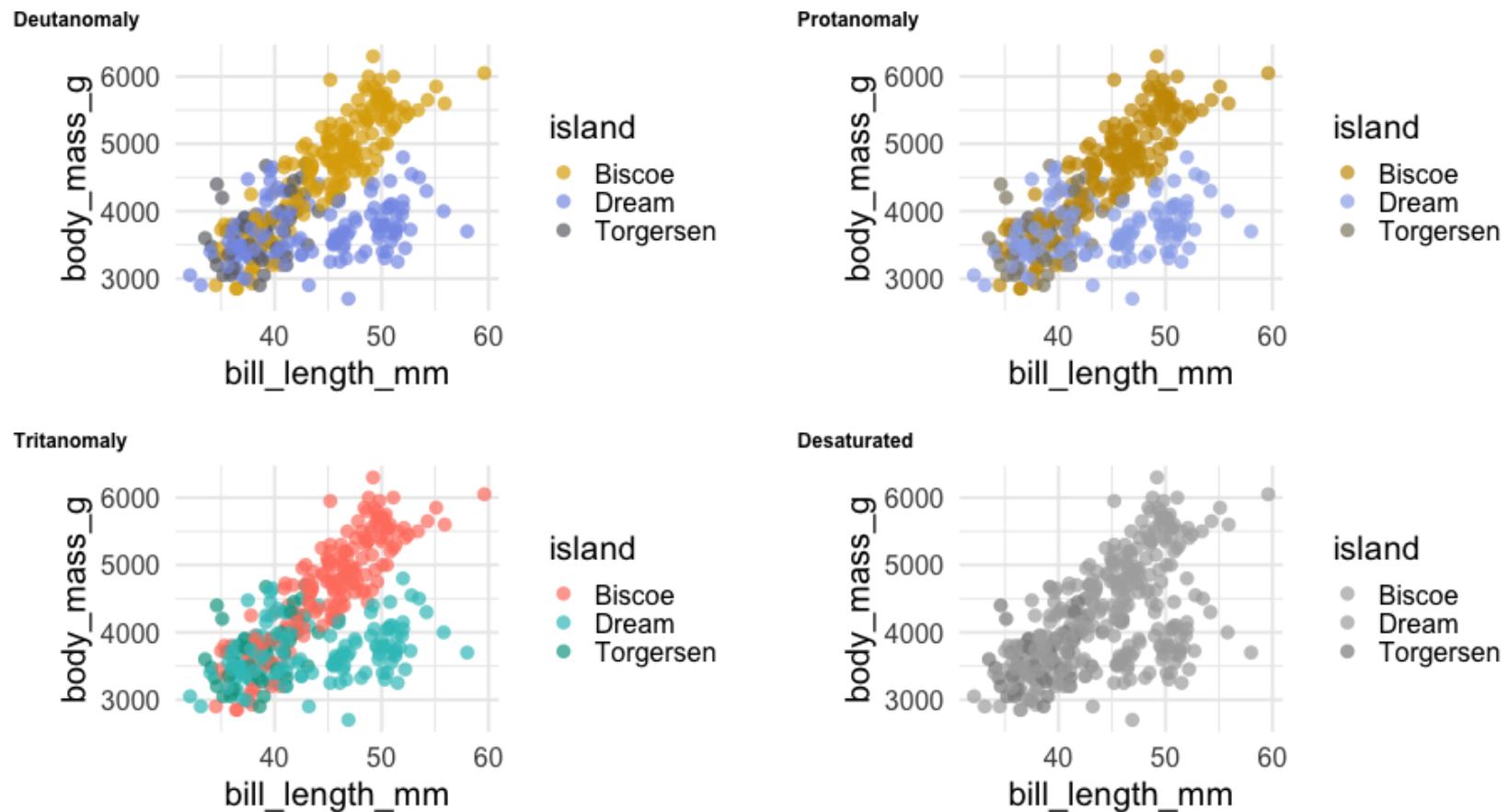


Double encodings

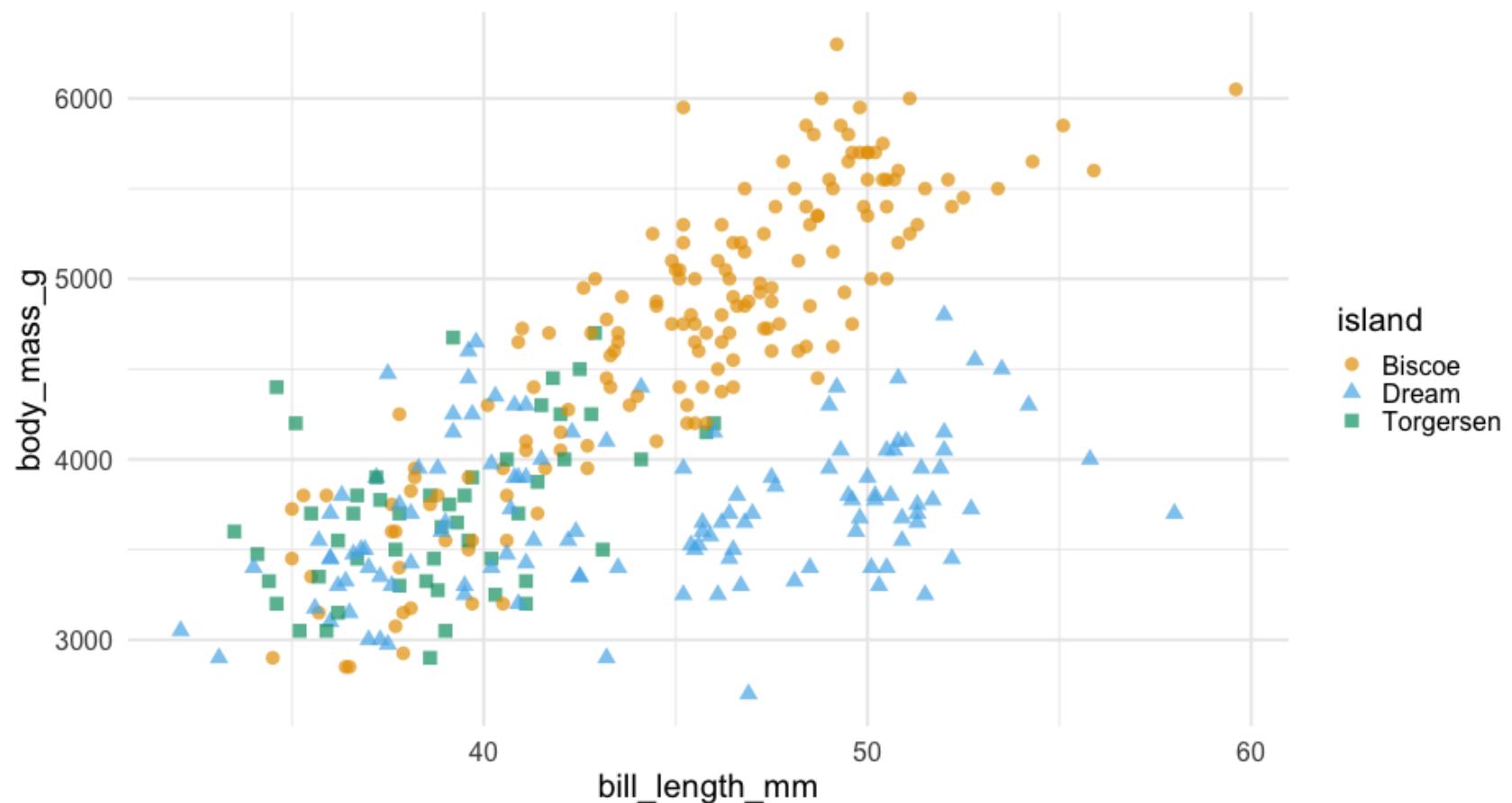


This plot is less than ideal. Why?

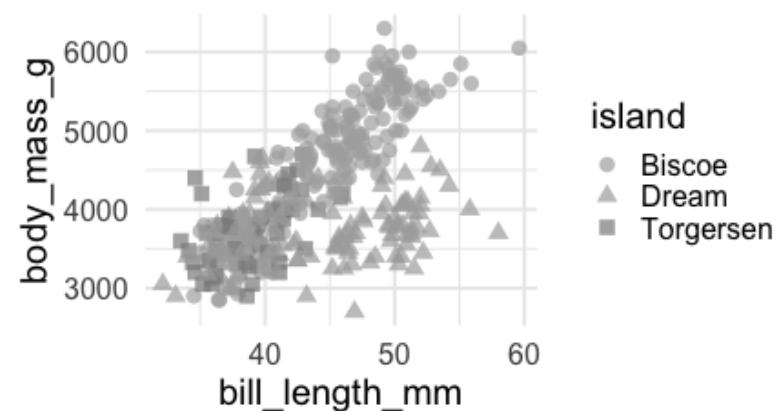
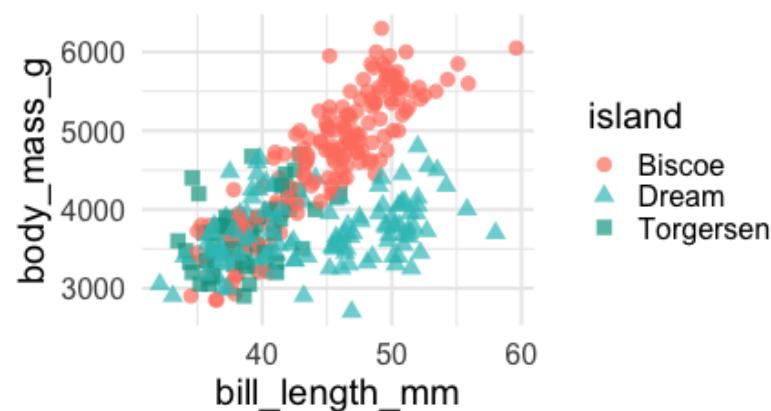
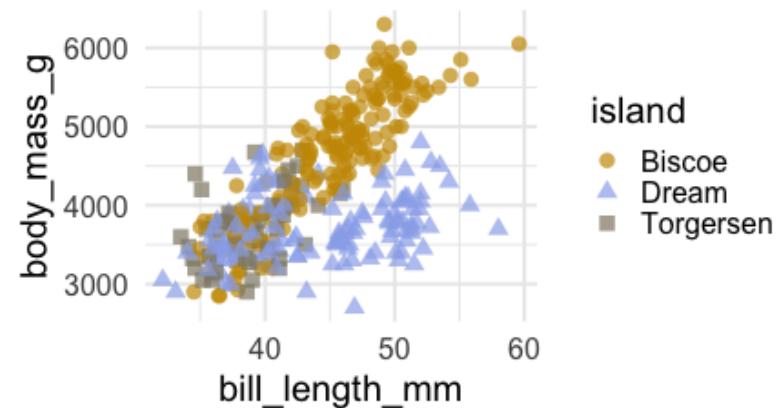
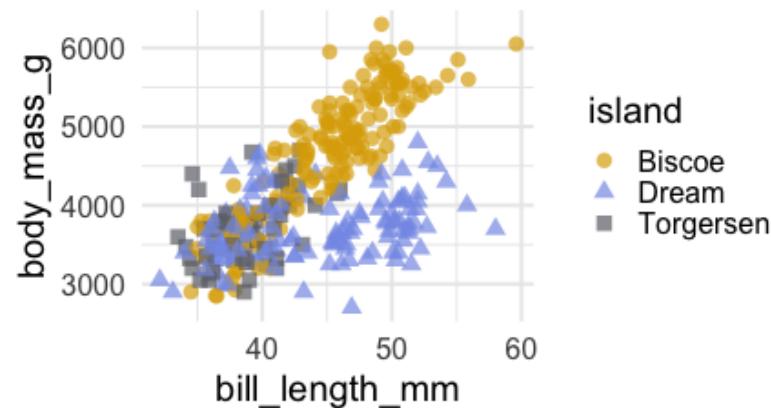
Color blindness



Better version



Color blindness check

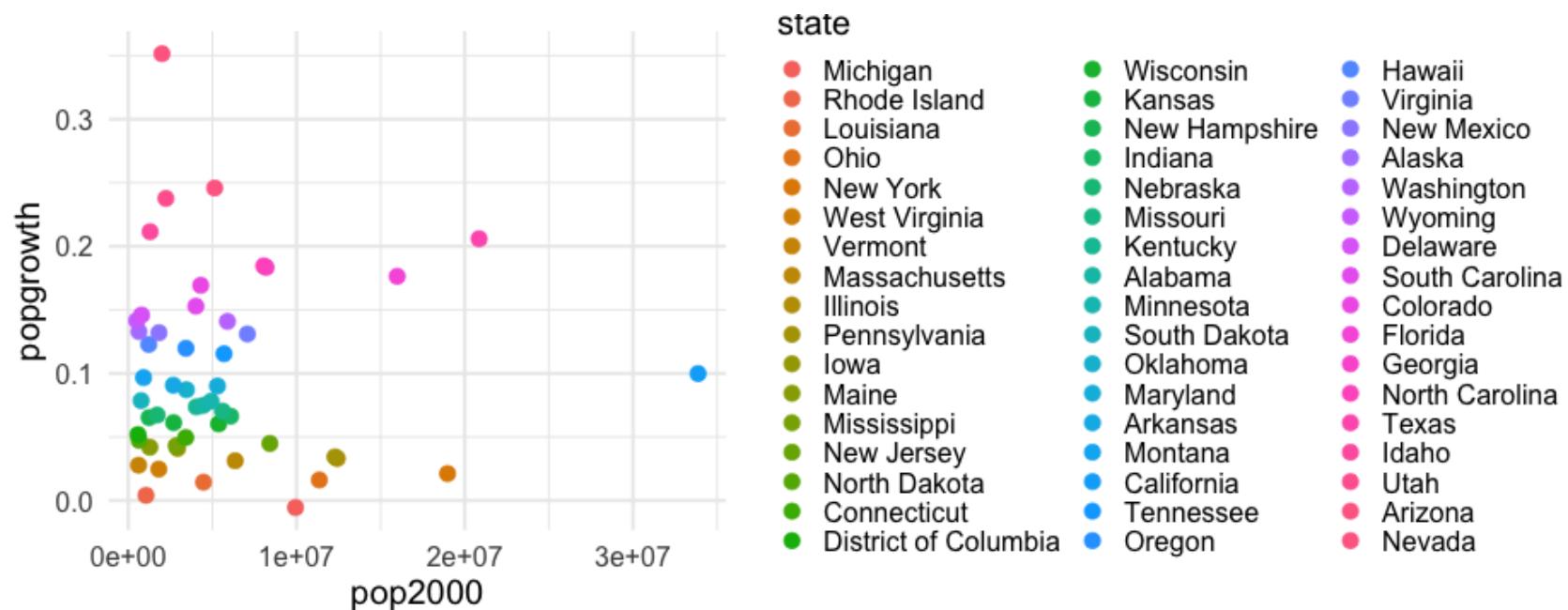


Common problems with color

Too many colors

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

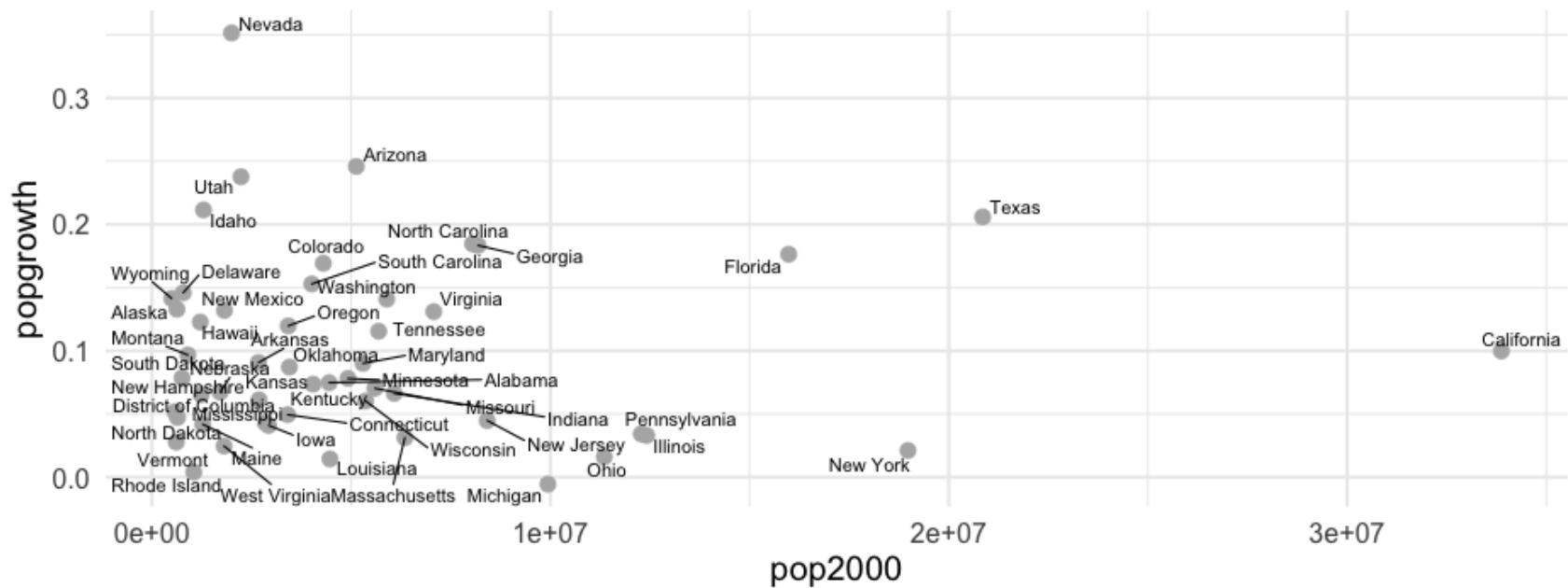
```
ggplot(popgrowth_df, aes(pop2000, popgrowth, color = state)) +  
  geom_point()
```



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))
```



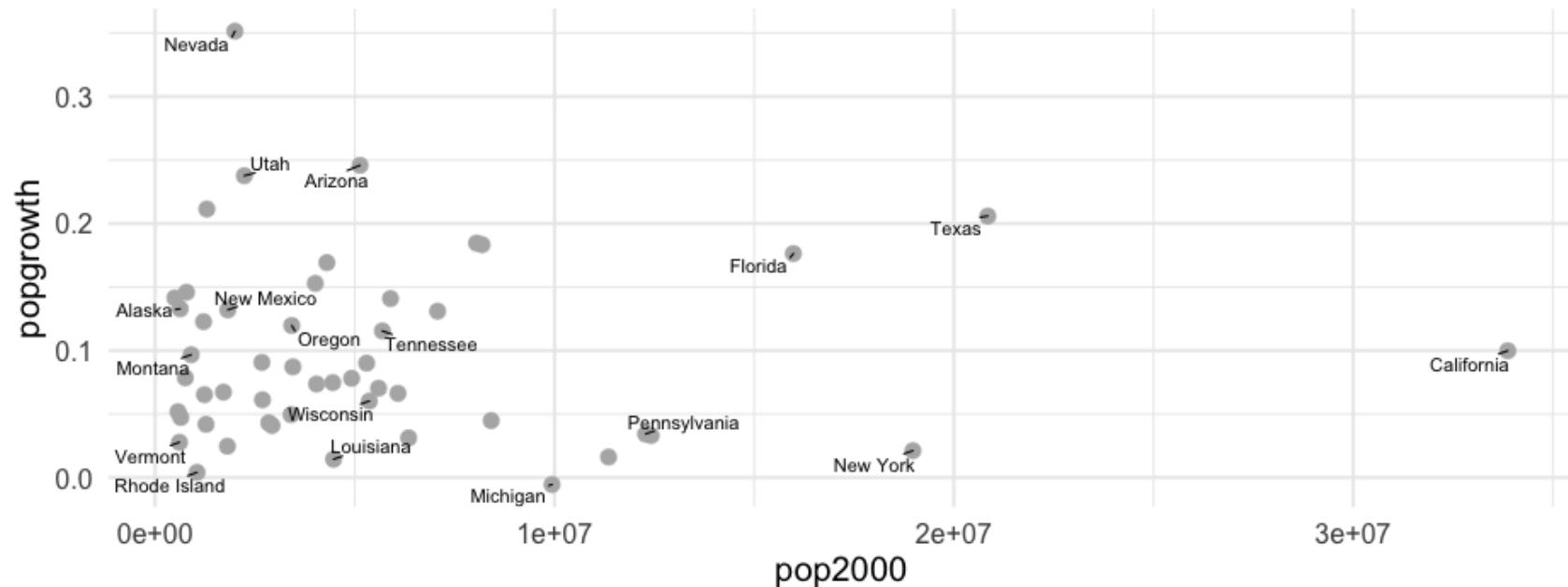
Better

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"
```

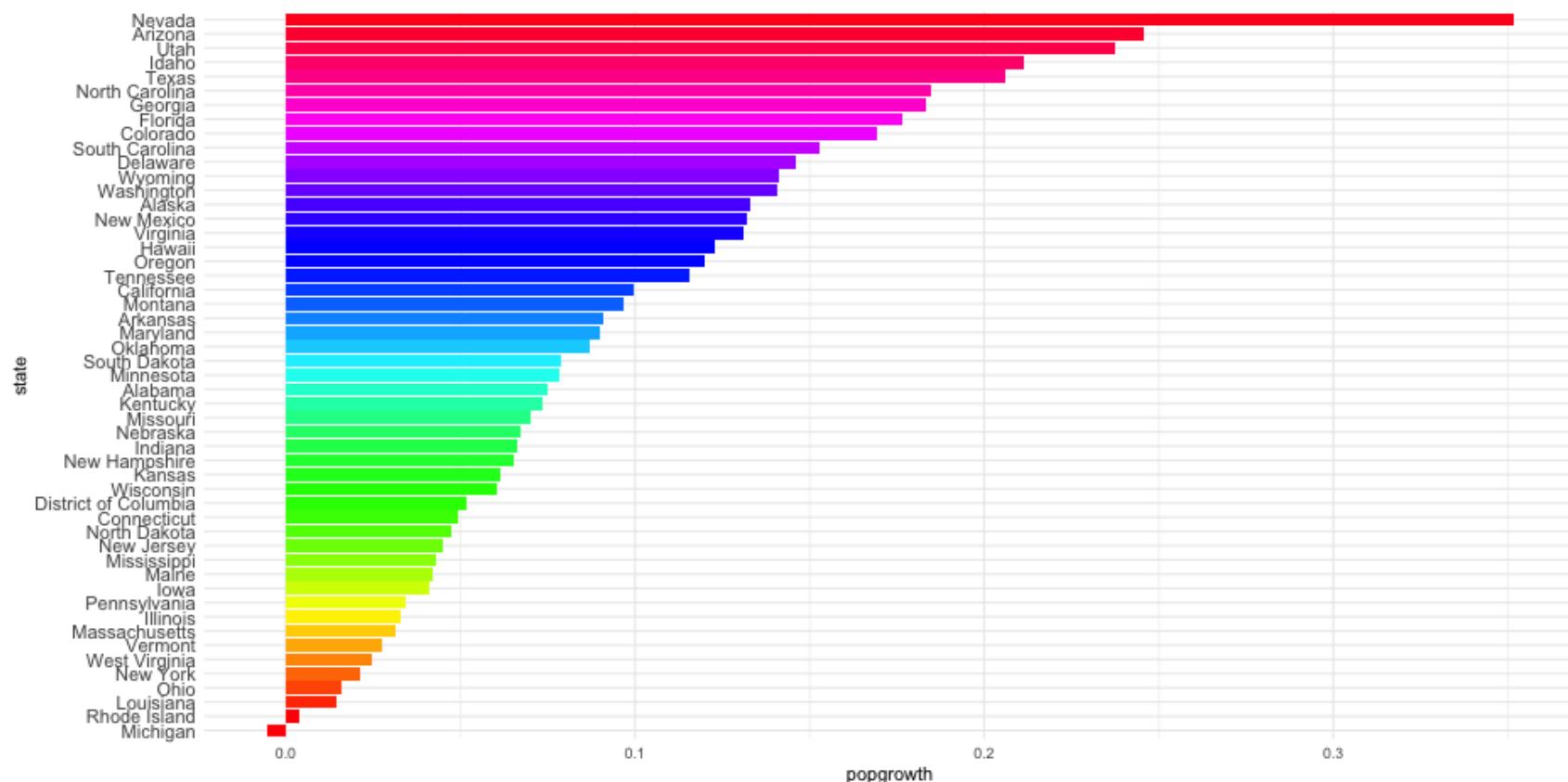
```
rainbow(7)
```

```
## [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")
```

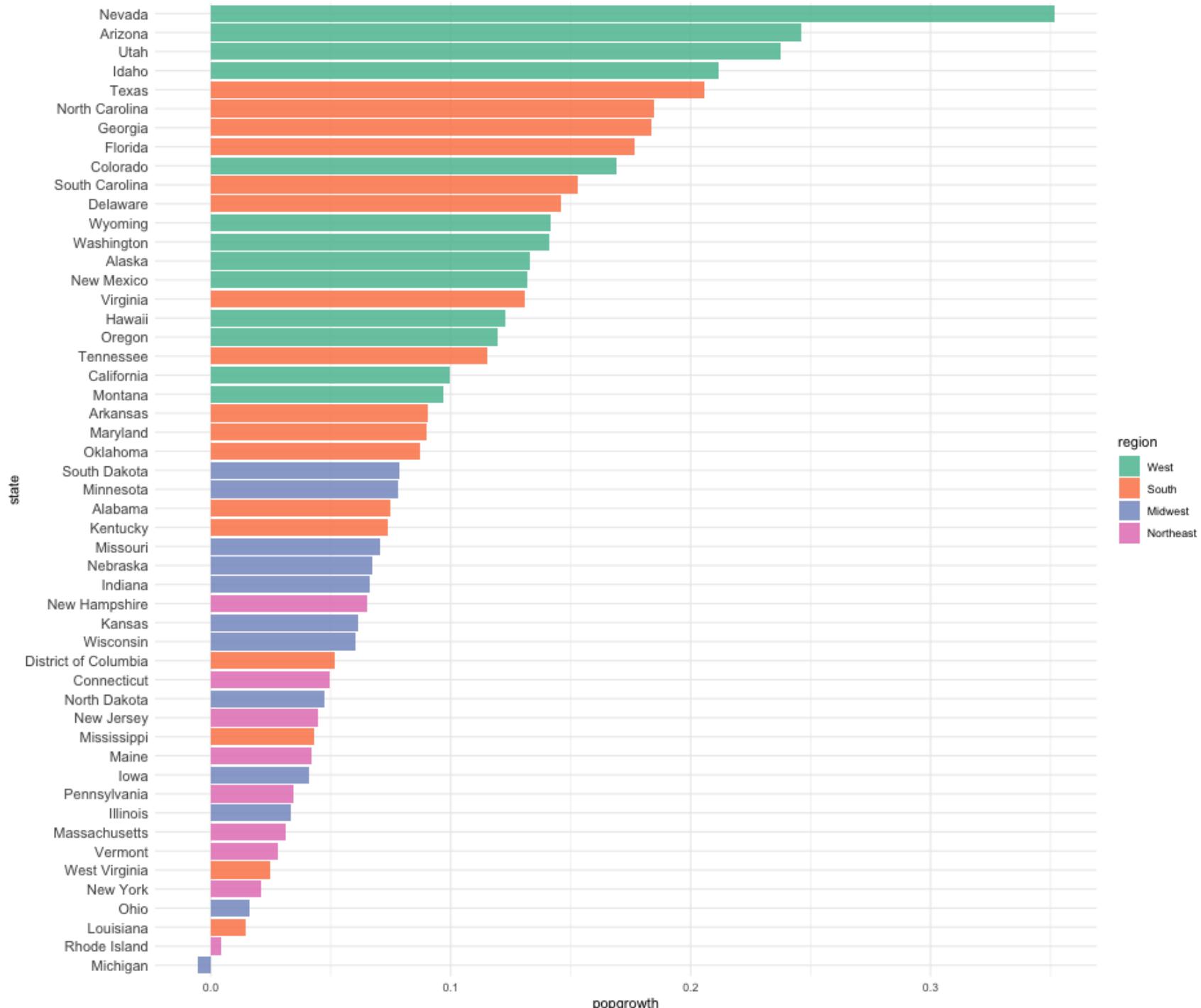


Last few note on palettes

- 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



Next time

Lab 3: Colors

Note – this will be our final lab to make sure you have sufficient time for your final projects